HIGH RESOLUTION NUMERICAL METHODS FOR COUPLED NON-LINEAR MULTIPHYSICS SIMULATIONS WITH APPLICATIONS IN REACTOR ANALYSIS

A Dissertation

by

VIJAY SUBRAMANIAM MAHADEVAN

Submitted to the Office of Graduate Studies of Texas A&M University in partial fulfillment of the requirements for the degree of

DOCTOR OF PHILOSOPHY

August 2010

Major Subject: Nuclear Engineering
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Approved by:

Chair of Committee, Jean C. Ragusa Committee Members, Marvin L. Adams, Jean-Luc Guermond, Jim E. Morel, Vincent A. Mousseau

Head of Department, Raymond J. Juzaitis

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ABSTRACT

High Resolution Numerical Methods for Coupled Non-linear Multi-physics Simulations with Applications in Reactor Analysis. (August 2010 )
Vijay Subramaniam Mahadevan, B.Tech., Bharathidasan University, India;
M.S., Texas A&M University
Chair of Advisory Committee: Dr. Jean C. Ragusa

The modeling of nuclear reactors involves the solution of a multi-physics problem with widely varying time and length scales. This translates mathematically to solving a system of coupled, non-linear, and stiff partial differential equations (PDEs). Multi-physics applications possess the added complexity that most of the solution fields participate in various physics components, potentially yielding spatial and/or temporal coupling errors. This dissertation deals with the verification aspects associated with such a multi-physics code, i.e., the substantiation that the mathematical description of the multi-physics equations are solved correctly (both in time and space).

Conventional paradigms used in reactor analysis problems employed to couple various physics components are often non-iterative and can be inconsistent in their treatment of the non-linear terms. This leads to the usage of smaller time steps to maintain stability and accuracy requirements, thereby increasing the overall computational time for simulation. The inconsistencies of these weakly coupled solution methods can be overcome using tighter coupling strategies and yield a better approximation to the coupled non-linear operator, by resolving the dominant spatial and temporal scales involved in the multi-physics simulation.

A multi-physics framework, KARMA (K(c)ode for Analysis of Reactor and other Multi-physics Applications), is presented. KARMA uses tight coupling strategies for
various physical models based on a Matrix-free Nonlinear-Krylov (MFNK) framework in order to attain high-order spatio-temporal accuracy for all solution fields in amenable wall clock times, for various test problems. The framework also utilizes traditional loosely coupled methods as lower-order solvers, which serve as efficient preconditioners for the tightly coupled solution. Since the software platform employs both lower and higher-order coupling strategies, it can easily be used to test and evaluate different coupling strategies and numerical methods and to compare their efficiency for problems of interest.

Multi-physics code verification efforts pertaining to reactor applications are described and associated numerical results obtained using the developed multi-physics framework are provided. The versatility of numerical methods used here for coupled problems and feasibility of general non-linear solvers with appropriate physics-based preconditioners in the KARMA framework offer significantly efficient techniques to solve multi-physics problems in reactor analysis.
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I also gratefully acknowledge the funding sources from Idaho National Laboratory for supporting my doctoral research since 2007.

I offer my sincerest tribute to my family for their patience and understanding over the past years. Without their continual support, completion of the research work and this dissertation would not have been even remotely possible.

And to all my friends in Bangalore and Chennai, and to my class-mates in College Station (David Ames, Yaqi Wang, Ayodeji Alajo and many more) who have been with me through thick and thin, I thank you all for your encouragement and support. I wouldn’t have made it without you!
To my father for helping me find my KARMA.

‘Blessed is he who has found his work; let him ask for no other blessedness.’
– Thomas Carlyle
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1. INTRODUCTION

‘All models are wrong, but some are useful.’
– George Box

High fidelity computer simulations of coupled multi-physics problems require solving large systems of non-linear, stiff, coupled equations. Many examples of non-linearly coupled multi-physics phenomena exist in various scientific fields, raising a need to develop stable and accurate numerical solution procedures. Some examples are:

1. Radiation diffusion where the radiation energy is strongly coupled to the material temperature field [1], [2].

2. Nuclear reactor analysis where the thermal power generated due to fission reactions in the fuel pin is strongly coupled with the thermal-hydraulics fields [3], [4].

3. Fluid-Structure-Interaction (FSI): the fluid and structural vibrations are coupled to each other. Applications in Automotive Systems, Nuclear Power Plants (NPP), Biomedical Applications, etc. [5], [6], [7].

4. Thermo-mechanical coupling: the temperature distribution affects the structural deformation and vice versa [8] [9].

Solution methods for non-linearly coupled multi-physics phenomena occurring have often relied on operator-split coupling strategies that introduce several types of errors in the solution fields. The new paradigm shift for multi-physics simulations is to quantify and reduce the sources of the errors due to the discretizations in space

---

This dissertation follows the style of Journal of Computational Physics.
and time and the resolution technique used to solve the non-linear coupling between the physics models. This requires stable and accurate numerical schemes that can tackle non-linearly coupled, stiff multi-physics problems arising from the discretization of the various physics Partial Differential Equations (PDEs) with widely varying characteristic time and length scales. The use of verified physical models for problems of interest and the accurate resolution of these characteristic physical scales are not trivial. This Dissertation is aimed at combining consistent numerical methods with principles in software engineering to create a coupled physics framework that is verifiable and can help better quantify these simulation errors.

Numerical simulation using computers is considered as the third pillar of science, besides theory and experiments. This dependence on computers as a virtual laboratory has been recognized in recent years, due to rapid growth in computer speed and affordable memory. Hence, cost effective development and design of scientific applications can be considerably accelerated by the use of simulations on powerful computing systems. In order to make use of solutions from computer simulations for multi-physics problems, it is important to predict the behavior of these non-linearly coupled systems. Predictive science, defined as the development and application of verified and validated (V&V) computational simulations to predict the properties and dynamic response of complex systems particularly in cases where routine, separable, experimental tests, while important, are difficult. This definition, borrowed from the Predictive Science Academic Alliance Program (PSAAP) of the US DOE-NNSA [10], is applicable to a wide variety of scientific and engineering applications. This emphasis on predictive capabilities has resurged the need to create and utilize robust numerical techniques for solving coupled problems with high resolution.

The current work focuses primarily on the development and usage of existing analysis and numerical methods for creating a unified and verified tool with predictive capability in the field of multi-physics nuclear reactor computation. Typically the current practice of multi-physics simulations in reactor applications, combines
models or algorithms from a diverse set of disciplines. The path towards the predictability of such computations requires the effective integration of both software and numerical methods. Generally speaking, the three pillars of predictive science include code verification, model validation, and uncertainty quantification in the computed solution.

The Dissertation is concerned with the development of a multi-physics software platform and the verification of numerical methods for multi-physics applications and an application of uncertainty propagation. Verification is typically an exercise in mathematics, where one assures that the equations are solved correctly, i.e., the software has been coded precisely and implemented according to the physics specifications and requirements. This is an integral part of any software development cycle for simulating physical phenomena.

The need to quantitatively predict the behavior of physical phenomena requires that the sensitivity of the solution fields to uncertainties in the parameters involved in the simulated physical models need to be ascertained. If not, the value of the simulations in comparison to real world experimental results is limited. Noting these as the basic requirements for a complex multi-physics code, we shall systematically develop relevant physical models and use efficient, high resolution numerical techniques in the current work.

In the next subsection, a short background on the current state of coupling methods is provided along with an introduction to the coupling methods that are implemented in the current work.

1.1 Background

Let the non-linear vector-valued function representing a coupled PDE system be written in a general form as

$$\mathbf{F}(\mathbf{y}) = \mathbf{N}(\mathbf{y})\mathbf{y} - \mathbf{b} = 0,$$  \hspace{1cm} (1.1)
where $\mathbf{y}$ is the solution vector that is dependent on both space and time respectively and $\mathbf{F} : \mathbb{R}^n \rightarrow \mathbb{R}^n$, $\mathbf{F}$ is the non-linear operator representing the coupled system and $n$ is the total number of unknowns. For ease of comprehension, we can write $\mathbf{F}$ as in the second equality of Eq. (1.1), where $\mathbf{N}$ is also a non-linear operator and $\mathbf{b}$ is the load vector. It helps to represent $\mathbf{y}$ as a vector comprised of the solution vector for each of the $M$ physics components involved, i.e., $[\mathbf{y}_1, \mathbf{y}_2, \ldots, \mathbf{y}_M]^T$. A similar definition holds for $\mathbf{F}(\mathbf{y})$ and its $m$-th component is the non-linear residual stemming from the $m$-th physics component and may depend effectively on all other fields, e.g., $\mathbf{F}_m(\mathbf{y}) = \mathbf{F}_m(\mathbf{y}_1, \mathbf{y}_2, \ldots, \mathbf{y}_M)$.

In the next subsections, the application of different coupling strategies to resolve the non-linear problem in Eq. (1.1) is presented.

1.1.1 Loose Coupling Strategies

In the past few decades, high fidelity modeling of non-linear multi-physics problems has been subdivided into several distinct domains of physics and solved individually as mono disciplinary blocks with specialized codes, without rigorous coupling between the different physics. Although naive, this coupling strategy, mathematically described as Operator-Split (OS) technique, is widely used. With the advent of Parallel Virtual Machines (PVM) and Message Passing Interfaces (MPI) in the 1990’s, the OS coupling of several existing specialized single physics codes has become the main multi-physics paradigm in reactor analysis. This kind of modeling is based on coupling several existing specialized mono-disciplinary codes using a 'black-box' strategy, where the input of one code is the output of other, thereby producing solutions that are weakly coupled. The schematics of such models is shown on Fig. 1.1, where the system of PDEs arising from the spatial and temporal discretization of physical models is decomposed into simpler sub-problems. Each physics component is solved by an independent, specialized single-physics code and the data between codes is exchanged through message passing paradigms. Often, this strategy
is non-iterative and the non-linearities due to the coupling in between the physics components are not resolved over a time step, reducing the overall accuracy in the time stepping procedure to first-order $O(\Delta t)$, even though high-order time integration might have been used for the individual physics components; see [11, 12]. Note that this explicit linearization of the problem in the OS strategy does not resolve the non-linearities between the different physics. Yet, these isolated physical models in reality describe physical phenomena that are tightly intertwined and rely heavily on the solution field of one another.

For illustration, consider the non-linear coupled system shown in Eq. (1.1). In OS loose coupling strategy, the non-linear operator is linearized as follows through an explicit treatment:

$$F(y^{\ell+1}) = N(y^\ell)y^{\ell+1} - b, \quad (1.2)$$

Hence the new update to the solution is obtained by solving the system

$$N(y^\ell)y^{\ell+1} = b. \quad (1.3)$$

Although OS allows parts of the problem to be treated implicitly and others explicitly, the lack of iterations in the conventional strategy degrades the solution accuracy in time to first order and the explicit linearization imposes a conditional stability limits for the time-step selection. The direct implication of using smaller time steps to achieve a reasonable accuracy is that the computations need greater CPU time and resources. Despite these drawbacks, this is still one of the major coupling paradigms used today for solving non-linear multi-physics systems.
The attractive feature of such a coupling strategy is that the legacy of many man-years of mono-disciplinary code development and V&V (validation and verification) is preserved. It is of prime importance to analyze the coupling strategies that can produce highly accurate solutions even in the complex scenarios usually encountered in multi-physics applications. As mentioned before, nuclear reactor analysis is a good example of highly non-linear, coupled multi-physics problem and the non-linearities at the heart of reactor design, analysis and safety calculations provide a good state-space to test high-fidelity numerical methods for multi-physics problems.
Physical phenomena such as the ones found in reactor accidents, involve rapidly varying transients that are represented by a stiff system of differential equations. Stiff problems are characterized by solutions having fast varying modes together with slower varying modes, requiring time integrators that can handle such disparate time scales. Stiff problems necessitate the use of implicit time discretization for stability reasons, indicating that OS coupling could prove disadvantageous in terms of efficacy (cost for obtaining a certain accuracy in the solution).

Current examples of OS coupling in the field of nuclear reactor analysis involve the following pairs of neutronics/thermal-hydraulics codes: CRONOS/FLICA [13, 14], PARCS/TRACE [15] and NESTLE/RELAP [16]. Even though more advanced OS strategies exist and can be up to second-order accurate in time, they are complicated to use in coupled legacy codes and hence are not currently employed. For more details regarding these higher order OS schemes, we refer the reader to [17, 18, 19, 20].

1.1.2 Tight Coupling Strategies

An alternative to loosely coupled OS strategies is to converge the non-linearities between the physics at every time level to obtain a tightly coupled solution that is consistent with the non-linear system of PDEs. This preserves higher order temporal accuracy of specialized schemes that can be used for resolving the disparate temporal scales in the different physics. Even though the cost/time step can be larger than that of an OS time step, it is essential to stress that the stability of the higher order discretization scheme can be maintained using this procedure, unlike the explicit linearization method where the solution is only conditionally stable.

To devise such a tightly coupled solution procedure, a non-linear iterative scheme needs to be applied to solve the coupled physics and converge the non-linearities to within user’s specified tolerances. Two techniques for non-linear system of equations are mentioned next: the Fixed-point or Picard iteration technique and the well known Newton’s method.
1.1.2.1 Picard Iteration

Picard iteration technique is a simple non-linear iterative method can be used to converge the non-linearities over the different physics when an OS coupling technique is employed to couple multiple physics codes. Picard iterations can restore the convergence order of a higher order scheme and eliminate the loss of accuracy due to the crude explicit linearization in loosely coupled strategy. The schematic for such a method is shown in Fig. 1.2. This essentially involves iterating over the solution obtained by successively solving Eq. (1.3).

![Fig. 1.2. High-order, Converged OS Coupling Strategy](image)

The advantage of such a coupling scheme is that it is non-intrusive and can easily use existing framework of codes to obtain a tightly coupled solution. But the primary disadvantage of using such a strategy to restore the accuracy is the increase in computational cost and memory usage to converge the solution.

Since Picard iteration is only linearly convergent, some form of non-linear acceleration techniques are necessary to make this scheme efficient and feasible [11]. Previous research using Aitken’s iterated $\Delta^2$ technique suggests that usage of such acceleration schemes can be advantageous and efforts to apply Wynn-Epsilon [21] and other schemes should be pursued as future extensions.
1.1.2.2 Newton Iteration

Current OS strategies may offer flexibility in the way the different physics are solved but involve complexities in terms of resolving the non-linearities and finding an high-order accurate solution. Instead, the coupled non-linear problem can be tackled by recasting it as a root finding problem, in a form amenable for the application of Newton-type methods.

Applying a Newton’s method to system of equations in Eq. (1.1), we obtain the following recurrence equation:

\[
J(y^\ell)\delta y = -F(y^\ell) \quad (1.4)
\]
\[
y^{\ell+1} = y^\ell + \delta y. \quad (1.5)
\]

where \( \ell \) is the Newton iteration index, \( \delta y \) is the solution update, and \( J(y^\ell) \) is the Jacobian matrix evaluated at \( y^\ell \). The Jacobian matrix is defined as

\[
J(y) = \frac{\partial F(y)}{\partial y}. \quad (1.6)
\]

Note that in Eq. (1.6), the Newton linearization accurately accounts for the true Jacobian of the non-linear system while the OS linearization in Eq. (1.3) neglects a term in the Jacobian matrix expansion \( (J(y) = N(y) + \partial_y N \cdot y \approx N(y)) \). This additional term contributes to the stability and robust convergence properties of the Newton iteration as compared to the Picard iteration shown earlier.

At each Newton’s iteration, a linear system of equations involving the Jacobian matrix, Eq. (1.4), needs to be solved. As the number of physics components grows, so do the total number of unknowns, resulting in a large memory usage to store the Jacobian matrix. However, employing a Jacobian-free approximation avoids the need for the expensive Jacobian calculation and storage of the matrix since only the action of the Jacobian on a vector is needed to solve the linear system.
Noticing the similarities between the tightly coupled methods with Picard iteration by solving recursively Eq. (1.3) and with Newton iteration by solving Eq. (1.4), we introduce a unified framework that is referred henceforth as the Matrix-free Nonlinear Krylov (MFNK) method. Note that this is based on the Jacobian-free Newton-Krylov (JFNK) method proposed by Brown and Saad in the early 1990’s [22], that has enjoyed much success in recent years in several multi-physics applications [23]. When Newton’s iteration is used as the non-linear solver, MFNK reduces to the original JFNK technique.

Several researchers have analyzed (a) the applicability of this tightly coupled method to obtain high-order accurate solutions and (b) the feasibility of the method in terms of total computational cost [24,25,26]. These prior results indicate that this scheme can tackle the widely varying time scales occurring in multi-physics problems efficiently, as compared to an OS coupling strategy. Note that the application of these tight coupling methods based on Picard or Newton iteration is not only limited to PDEs written in the conservative form alone as in Eq. (1.1).

With the aforementioned background ideas, the motivations for the current research work is laid out next.

1.2 Research Motivation

To overcome the issues stated in section 1.1, a fully implicit treatment of the coupling terms needs to be used to preserve accuracy and obtain unconditional stability. The difficulties in implementing such a scheme is that the spatial and temporal discretizations of all the physics need to be non-linearly consistent. With such discretizations, the coupling terms in the physics are also treated implicitly and hence higher order accuracy is ensured by resolving the non-linearities accurately. In the current work, a new code system is created based on the MFNK framework with higher order spatio-temporal schemes for all the physics in addition to the ability to simultaneously test OS coupling schemes side-by-side. Also, most existing mono-
disciplinary codes for reactor physics simulation were written one to three decades ago to run on computers that existed during that period. Due to the current advances in computing, it would be rather imprudent to develop a new multi-physics code that does not take advantage of the state-of-the-art multi-core, multi-processor parallel architectures that are available now and with expandability to more advanced technologies in the future.

Predictability of the solution is a driving factor in this research and hence it is imperative to obtain a completely verifiable code where the numerical convergence order from the spatial and temporal treatment of the coupled PDEs can be measured against the theoretical orders seamlessly. With computational efficiency in mind, the matrix-free approach through MFNK for the non-linear solve eliminates large storage requirements of the discretized systems and competent numerical and physics based preconditioning techniques \cite{27} can be used to considerably reduce the cost of the linear Krylov iterations. Previous work using JFNK for non-linear diffusion-reaction and advection-reaction problems \cite{28} have shown promising results and serve as the basis for the new coupling strategy being implemented here. It is expected that such a scheme will enable achieving the higher orders of spatio-temporal accuracy for all coupled solutions.

The prime motivation behind the new code is not to employ high fidelity physics models coupled to each other with high resolution but rather to create consistent coupling methodologies that can test the feasibility of using physics-based preconditioned MFNK schemes for real-world problems in reactor design and safety analysis.

1.3 Dissertation Organization

The layout for this dissertation is as follows: in Section. \(2\) we discuss the equations for the physics models used to describe nuclear reactor cores and the governing relations that couple the different physics. In Section. \(3\) we provide a detailed overview of the different spatio-temporal discretizations, the numerical techniques
based on JFNK scheme, and the preconditioning methods employed to reduce the number of Krylov iterations. In Section. (4), a new code system that implements the physics models of Section. (2) and the numerical methods of Section. (3) is introduced and details regarding the software architecture are provided. Next, the code system is put to test using problems created with Method of Manufactured Solutions (MMS) and benchmarks to verify higher order treatment of all the physics models in Section. (5). Finally, in Section. (6) we discuss the details of using the MFNK framework to solve eigenvalue problems occurring commonly in nuclear reactor analysis and compare it to state-of-art schemes like Arnoldi and Jacobi-Davidson iterations. Also, the MFNK technique is applied to a stiff non-linear multi-physics problem based on a radiation diffusion physics model to emphasize the flexibility of applying the implemented code for problems not related to nuclear reactor simulations. Then, we draw conclusions and point out avenues for future research in Section. (7).
2. PHYSICS MODELS

‘The more you see how strangely Nature behaves, the harder it is to make
a model that explains how even the simplest phenomena actually work.’
– Richard Feynman

In this section, details regarding the physics models used in this Dissertation are
provided. All models have been deliberately chosen to be of “coarse” fidelity as the
purpose of this research work is not to validate the physical models themselves but
to present a multi-physics verification study that will help develop better intuition
regarding efficient coupling strategies. At a later stage, when the details about the
implementation code are given, a description will be provided for employing higher
fidelity physical models interchangeably within the MFNK framework, when they
are deemed necessary for the physics being solved.

In the realm of reactor analysis, there are three primary domains of physics that
play a pivotal role in determining core operation and safety. These are:

1. Neutronics - Describes the neutron population distribution and the interaction
   of neutrons with the material in the reactor core. The primary solution fields
calculated is the scalar flux as a function of position, time and neutron energy.

2. Thermal conduction - Describes the distribution of temperature in the fuel
   pin due to the sensible heat generated from the energetic neutron fission re-
   actions. The solution fields of interest are usually the temperature profile in
   the fuel element from which the peak fuel temperature and the maximum clad
temperature at the surface of the pin are obtained.

3. Coolant channel flow - Describes the flow of coolant fluid through the core that
   removes the thermal energy from the fuel pins. The models used can be for
   single or multi-phase fluid to calculate the density, momentum in all directions,
total energy, temperature and the pressure drop across the core.
Other physics components include structural mechanics that describes the behavior of thermal expansion in the fuel pins and structures comprising the core, kinetics of chemical reactions occurring due to flow of borated water in PWRs. In the current research, only the three basic physics models listed above have been used to create a multi-physics model to analyze nuclear reactor transients.

2.1 Neutronics

Neutronics is the branch of physics that deals with the calculation of neutron flux and neutron reaction rates in the different materials inside the reactor core. These reaction rates need to be calculated accurately in order to determine the power produced in a nuclear reactor and to calculate the temperature solution fields, which are strongly coupled to thermal energy generated in the fuel.

High-fidelity description of neutronics is usually provided by a neutron balance equation or the ‘neutron continuity equation’ for discrete energy groups that describes the neutron population in the phase-space domain. But finding a numerical solution to the neutron scalar flux \( \phi \) from the neutron continuity equation is an arduous task in itself, without coupling to other physics, especially when the reactor domain is large and heterogeneous and when many neutron energy groups \( G \) and delayed precursor groups \( K \) are employed. We base our neutronics model on the time-dependent Multi Group Neutron Diffusion (MGND) equation to solve for the neutron scalar flux.

\[
\frac{1}{v^g} \frac{\partial \phi^g(\vec{r}, t)}{\partial t} = - \vec{\nabla} \cdot D^g(\vec{r}, t) \vec{\nabla} \phi^g(\vec{r}, t) + \Sigma_{t,g}(\vec{r}, t) \phi^g(r, t) = \sum_{g'=1}^{G} \Sigma_{s}^{g' \rightarrow g}(\vec{r}, t) \phi^{g'}(\vec{r}, t) + \chi_p^g \sum_{g'=1}^{G} \nu^{g'} \Sigma_{f}^{g'}(\vec{r}, t) \phi^{g'}(\vec{r}, t) + \sum_{j=1}^{J} \chi_{d,j}^g \lambda_j C_j(\vec{r}, t) \quad \forall g \in [1, G], \forall \vec{r} \in D
\]  

(2.1)
The notations used here are standard [29]. The system of equations is closed with appropriate boundary and initial conditions. We can see that the neutron flux $\phi$ is dependent on the position in the core, the energy of neutrons and on time.

A nuclear core is typically composed of hundreds of different materials and isotopes, each with different crosssections. The crosssection of a material is greatly affected by the temperature and density of the material and depends on the energy of the incident neutron. In this coarse grain neutron diffusion model, the heterogeneity of the materials have been averaged to create fuel assembly homogenized material crosssections (piece-wise constant crosssection values per assembly) that preserve the total reaction rates in the core. The crosssections are usually tabulated, or provided in a closed form approximation, as a function of fuel and coolant temperatures (extension to additional parameters, such a boron concentration, void history, control rod history, . . . , is straightforward). The tabulated crosssection values are obtained using table look-up and $R_p$ interpolation, where $p$ is the total number of parameters used.

The Ordinary Differential Equations (ODEs) for the evolution of delayed neutron precursor concentrations are given by

$$
\frac{dC_j(\vec{r}, t)}{dt} + \lambda_j C_j(\vec{r}, t) = \beta_j \sum_{g'=1}^{G} \nu \Sigma_{f}^{g'}(\vec{r}, t) \phi^{g'}(\vec{r}, t) \quad \forall j \in [1, J].
$$

(2.2)

The precursor concentration balance is obtained based on the rate of production from fission reactions and losses due to radioactive decay given by the half-life $\lambda$.

The energy production due to the fission or radiative capture events is given by

$$
Q(r, t) = \sum_{g=1}^{G} \left[ \kappa_{f}^{g} \Sigma_{f}^{g} + \kappa_{c}^{g} \Sigma_{c}^{g} \right] (\vec{r}, t) \phi^{g}(\vec{r}, t),
$$

(2.3)
where the $\kappa$ coefficients represent the amount of energy released per reaction event, the subscript $f$ represents fission reactions and subscript $c$ represents radiative capture reactions.

The design of reactors is often carried out in Steady-State (SS) where the distribution of the neutron flux is considered to be in equilibrium. In this static state with fissile material present and no external source, the MGND equation reduces to an eigenvalue problem. The dominant eigenvalue of the system, called the effective multiplication factor $k_{\text{eff}}$, is defined as

$$k_{\text{eff}} = \frac{\text{Number of neutrons in one generation}}{\text{Number of neutrons in the preceding generation}} \quad (2.4)$$

The determination of this parameter is done by solving the following modified form of the MGND equation Eq. (2.1),

$$- \nabla \cdot D_g(\vec{r}, t) \nabla \phi_g(\vec{r}, t) + \Sigma_{t,g}(\vec{r}, t) \phi_g(\vec{r}, t) - \sum_{g' = 1}^{G} \Sigma_{s}^{g'} g(\vec{r}, t) \phi_{g'}(\vec{r}, t) = \frac{X_p}{k_{\text{eff}}} \sum_{g' = 1}^{G} \nu \Sigma_{f}^{g'}(\vec{r}, t) \phi_{g'}(\vec{r}, t) \quad \forall g \in [1, G]. \quad (2.5)$$

with appropriate boundary conditions.

This generalized eigenvalue problem relates the fundamental eigenvalue (dominant) representing the $k_{\text{eff}}$ and its corresponding eigenmode representing the scalar flux $\phi^g(r, t)$ for SS conditions. Since the flux is obtained as a solution of the eigenproblem, only the shape of the flux can be ascertained and the magnitude is determined based on the total power load chosen during operation. Criticality provides information for the design of a reactor and also serves as a tunable parameter to determine the conditions for continuous power output.

In the current work, the statics are governed by Eq. (2.5) and the dynamics of solution field evolution is described by the MGND equation Eq. (2.1) and precursor
equations Eq. (2.2). These closed set of equations, along with boundary and initial conditions, form the neutronics model of this work.

2.2 Thermal Conduction Model

The energy production due to the fission reaction in the fuel elements generate sensible heat energy which is deposited locally in the fuel. This energy is conducted outward towards the surface of the fuel pellet, the gap and the outer cladding so that it can be transferred to the coolant. The conservation equation to model this physics in Cartesian coordinates ($\vec{r} = x, y, z$) can be written simply as

$$\rho(T)C_p(T)\frac{\partial T(\vec{r},t)}{\partial t} - \vec{\nabla} \cdot k \vec{\nabla} T(\vec{r},t) = q(\vec{r},t), \quad (2.6)$$

with appropriate boundary conditions on the outer trace of domain $\mathcal{D}$.

Here, the density ($\rho$) in $kg/m^3$, specific heat ($C_p$) in $J/kg\cdot ^\circ C$, and conductivity ($k$) in $W/m\cdot ^\circ C$ can depend on the temperature $T(\vec{r},t)$ and hence Eq. (2.6) is a non-linear equation by itself.

The boundary term coupling the conducting solid to the fluid is given by

$$k(T)\partial_n T|_w = h_c(T_w, T_f)(T_w - T_f), \quad (2.7)$$

where $T_w (^\circ C)$ is the (solid) wall temperature, $T_f (^\circ C)$ is the coolant temperature at the interface, and $h_c(T_w, T_f)$ ($W/m^2/\circ C$) is the convective heat transfer coefficient obtained by means of a closure relation.

The non-linear heat conduction model employed here represents the core as a porous medium where the fuel, the fluid flowing in the channel and the supporting structures are homogenized together and properties found accordingly. This is strictly used to verify and to test the code implementation since the model used is described typically in the same space as neutronics and the equation is a simple scalar non-
linear parabolic equation with non-constant heat source and mixed or Robin BC at the solid-fluid interface.

As a refinement of the porous model described above, a two dimensional diffusion-reaction equation in cylindrical coordinates can be used to find the fuel profile in a pin with the given average power density distribution. This model is described by

$$\frac{\partial \rho C_p T(\vec{w}, t)}{\partial t} - \frac{1}{r} \frac{\partial}{\partial r} \left( r k_r \frac{\partial T(\vec{w}, t)}{\partial r} \right) - \frac{\partial}{\partial z} \left( k_z \frac{\partial T(\vec{w}, t)}{\partial z} \right) = q_{avg}(\vec{w}, t), \quad (2.8)$$

where $q_{avg}(\vec{w}, t)$ is the average power density in a fuel pin.

Using this model, the average temperature profile and behavior of a region (traditionally a full assembly or part of it in a lattice) can be ascertained and used to find parameters to estimate the peak clad temperature (based on oxidation limits) and other safety parameters such as the maximum fuel temperature in order to eliminate the possibility of fuel melting. A sample schematic 1-d subchannel model that is traditionally used in reactor analysis codes is shown in Fig. 2.1.

![Subchannel Model](image)

Fig. 2.1. Subchannel Model

In such a subchannel model, the average power density corresponding to an assembly location, for a given axial region can be calculated to provide the necessary source terms for determining the temperature profile in the fuel pin. This is representative of the average fuel behavior in that region. The fuel surface temperature is coupled also to the coolant flow in the channel and is accounted using appropriate boundary conditions Eq. (2.7).
2.3 Coolant Fluid Flow Model

The coolant flowing in a channel outside the clad of the fuel element gains enthalpy by convection and removes heat generated and conducted in the fuel elements. The thermal hydraulics physics and heat conduction are coupled due to the heat transferred from the fuel pin surface to the coolant by means of convection. The temperature of the coolant is directly dependent on the temperature of the outer clad surface, which, in turn, is a direct function of the fission reaction rate, thereby making all physics coupled to one another. In addition, a volumetric heat source can also be present in the bulk of the coolant to model radiative capture energy release and direct gamma heating.

Typically, in nuclear reactors, the flow of the coolant/moderator fluid occurs in channels of vertical columns. Higher fidelity descriptions may use three-dimensional Navier-Stokes equations in either the conservative or non-conservative variable sets with appropriate turbulence models. In the current work, a simplified approach is taken and the coolant is modeled using a single-phase fluid flowing vertically in one-dimensional channels. The model allows for one or multiple 1-d average channels (the maximum number of channels being the number of right prisms describing the fuel assemblies in the neutronics model; a simple user-defined mapping is employed to assign channels to fuel assemblies). The fluid convects the heat generated either in the bulk of the fluid or at the fuel pin clad interface.

The governing equations for the fluid flow are solved in terms of the conservative variables and are given as:

\[
\begin{align*}
\partial_t \rho + \partial_z (\rho u) &= 0 \quad (2.9) \\
\partial_t (\rho u) + \partial_z (\rho u^2) + \partial_z P &= f_{w\rho} |u| u + \rho b_f \quad (2.10) \\
\partial_t (\rho E) + \partial_z (u(\rho E + P)) &= \partial_z q + S, \quad (2.11)
\end{align*}
\]
where $\rho$ is the fluid density, $\rho u$ its momentum, $\rho E$ its total energy, $P$ the pressure, $f_w$ is the ratio of dimensionless wall-friction factor and the hydraulic diameter $D_h$, $q$ the conduction of thermal energy in the fluid, $b_f$ is the net body force acting in the direction of velocity $v$ (for instance, acceleration due to gravity in the downward direction) and $S$ external source terms (energy from the fuel pin) through convective transfer. An equation of state closes the system of fluid equations:

$$P = f^\text{EoS}(\rho, \rho e),$$

where the internal energy is $\rho e = \rho E - \frac{1}{2} \rho u^2$.

An example of a closure relation for the equation of state Eq. (2.12) is given by the ideal gas law:

$$P = \rho e (\gamma - 1),$$

where $\gamma = \frac{C_p}{C_v}$, the ratio of specific heat at constant pressure to constant volume. Alternately, a more generic closure relation can be written by means of a linearized relation that is dependent on density and temperature:

$$P = P_0 + \alpha (\rho - \rho_0) + \beta (T - T_0),$$

where $\alpha, \beta$ are the constants that are valid about the linearization point $(P_0, \rho_0, T_0)$. Note that $\alpha$ is related to the speed of sound in the fluid and provides a simple way to alter the Mach number $(Ma)$ in calculations employing manufactured solutions. This is useful in verifying the numerical scheme used to treat this system of equations since they are stiff in the flow regimes of concern in nuclear reactors where low Mach flows dominate.

$$\alpha = \left[ \frac{\partial P}{\partial \rho} \right]_0 \propto \frac{1}{Ma^2},$$

As the fluid velocity becomes small in magnitude compared to the speed of sound in the medium, it is very difficult to solve the low-speed flow equations with a con-
ventional compressible algorithm because of their slow convergence. The difficulty in solving the compressible equations for low Mach numbers [30] is associated with the large disparity between the acoustic wave speed and the fluid speed, which contributes to stiffness, resulting in an indefinite system of equations.

Efforts to derive schemes that can tackle all speed flows (from low Mach to supersonic) using physics-based semi-discrete formulations [31,32], linearized perturbation equations [30] and methods based on asymptotic expansions (pressure separation formulation) in terms of $Ma$ [33] have been investigated previously. In the current research we consider a variation of the method introduced by Harlow [31] to tackle the stiff and low Mach flow regimes that are encountered in nuclear reactor applications. It is also important to note that the semi-discrete and asymptotic expansion methods share similar traits in tackling low Mach flows and further investigations to derive an elegant relation between these family of solvers is necessary to fully understand their mathematical implications.

2.4 Closing Remarks

The aim of the research presented in this Dissertation is to focus primarily on using better coupling techniques for a given physical equation model. Hence the physics models chosen in the current work are coarse but descriptive enough to analyze transient problems occurring in nuclear reactors.
3. METHODS FOR MULTI-PHYSICS SIMULATIONS

‘Knowing thus the Algorithm of this calculus, which I call Differential
Calculus, all differential equations can be solved by a common method...
not only addition and subtraction, but also multiplication and division,
could be accomplished by a suitably arranged machine.’
– Gottfried Wilhelm von Leibniz

In this section, details regarding the numerical methods employed to tackle the
coupled physics problems are provided. In addition to these coupling techniques,
which include a discussion of solution methods for non-linear and linear systems
and preconditioners, we also describe space and time discretization techniques with
adequate references to supporting materials.

3.1 Spatial Discretization

Boundary Value problems (BVP) and Initial BVPs for PDEs are often used to
model physical phenomena and hence a consistent and accurate discretization of these
equations to resolve the length and time scales correctly is pertinent. Parabolic and
Hyperbolic systems of PDEs or mixed systems are typically encountered as governing
equations for multi-physics applications. Let the bounded solution domain $\Omega$ be in a
d-dimensional space $\mathbb{R}^d$ with boundary $\Gamma$. Appropriate boundary conditions should
also be prescribed in order to close the system and yield a well-posed problem.

There are several options available for treating the spatial terms in these PDEs;
Finite Difference (FD), Finite Volume (FV) and Finite Element (FE) methods. All
of these methods, in one form or other, rely on replacing the true solution for the
original differential equation with a discrete form of the solution using approximate
expansions in terms of piecewise (higher order) polynomials. This reduces the prob-
lem to a finite system of coupled equations.
The spatial discretization of the mathematical models in the current work is performed using FE methods. This method is based on the variational form of the boundary value problem. The primary reasons for this choice are the ease of use for arbitrary geometries and irregular domains, the ability to employ nonuniform meshes to reflect sharp solution gradients, and the ability to obtain more easily high-order approximations. Also, rigorous a-priori error estimates of the discretization error based on the order of the polynomial basis functions are available, at reasonable costs, and can be used to adapt the finite element mesh to automatically refine/coarsen a subportion of the mesh based on a user-defined accuracy level.

Here, a Continuous Galerkin (cG) FE method [34] is utilized for Elliptic/Parabolic PDEs and a Discontinuous Galerkin (dG) FE method [35] is employed for Hyperbolic systems. Details regarding the variational form and the discrete equations obtained by applying solution approximations for Elliptic/Parabolic and Hyperbolic systems are given in the following subsections.

### 3.1.1 Elliptic Systems: Continuous Galerkin Discretization

Consider a non-linear, second-order BVP given by the following Elliptic PDE

$$-\nabla \cdot D(\vec{r}, u) \nabla u + c(\vec{r}, u) u = q \quad \forall \vec{r} \in \Omega. \quad (3.1)$$

where $D(\vec{r}, u), c(\vec{r}, u)$ are smooth functions with $D(\vec{r}, u) \geq D_0 > 0, c(\vec{r}, u) \geq 0$ in $\Omega$ and $q \in L_2(\Omega)$ with appropriate boundary conditions specified at the boundary $\Gamma$. The Galerkin weak form of Eq. (3.1) is obtained by multiplying the equation with a test function $v$ and integrating by parts over domain $\Omega$ to obtain

$$\int_{\Omega} D(\vec{r}, u) \nabla u \cdot \nabla v + v(c(\vec{r}, u) u - q) d\Omega - \int_{\Gamma} v D(\vec{r}, u) \nabla u \cdot \vec{n} ds = 0 \quad \forall \vec{r} \in \Omega. \quad (3.2)$$
where Green’s theorem or divergence theorem given below is employed.

$$\int_{\Omega} \nabla \cdot (vD \nabla u) = \int_{\Gamma} vD \nabla u \cdot \vec{n} ds, \quad (3.3)$$

with $\vec{n}$ being the outward unit normal vector on the boundary.

The variational form of the above problem is to find the solution $u$ belonging to the Sobolev space $H^1$ such that

$$a(u, v) = (q, v), \quad \forall v \in \Omega. \quad (3.4)$$

where

$$a(u, v) = \int_{\Omega} (D(\vec{r}, u) \nabla u \cdot \nabla v + c(\vec{r}, u)uv) d\Omega - \int_{\Gamma} vD(\vec{r}, u) \nabla u \cdot \vec{n} ds \quad (3.5)$$

$$ (q, v) = \int_{\Omega} (qv) d\Omega \quad (3.6)$$

Now, for the purpose of finding the approximate numerical solution, a non-overlapping partition of the domain $\Omega$ is introduced such that $\bigcup_{K \in T} K = \Omega$ and $T$ is a Triangulation of $\Omega$. For simplicity, an assumption is made that the geometry is exactly represented by the sum of the parts of the finite partition. The discrete solution is sought in the finite dimensional trial space $S_h$ of piecewise continuous polynomial functions of order $p$. For Galerkin FE method, the trial space and the test space are the same but continuity requirements on the test space are usually less restrictive.

Expanding the numerical solution $u$ and the weight function $v$ in terms of basis functions $\Phi(r)$,

$$u(\vec{r}) \approx U(\vec{r}) = \sum_{i=1}^{K} U_i \Phi_i(\vec{r}), \quad (3.7)$$

$$v(\vec{r}) \approx V(\vec{r}) = \sum_{i=1}^{K} V_i \Phi_i(\vec{r}), \quad (3.8)$$
where $U_i$ and $V_i$ are the degrees of freedom for the FE discretization. If the basis functions $\Phi(r)$ are chosen to be interpolatory, e.g., Lagrange basis functions, then the degrees of freedom satisfy $U_i = U(\vec{r}_i)$ and $V_i = V(\vec{r}_i)$.

The finite dimensional form of the problem can now be restated as follows: Find $u_h \in S_h$ such that

$$a_h(u_h, v_h) = (q, v_h), \quad \forall v_h \in S_h.$$  (3.9)

Inserting Eq. (3.8) in Eq. (3.9), the following weak form is obtained.

$$\sum_{k=1}^{K} U_k a_h(\Phi_k(r), \Phi_i(r)) = (f, \Phi_i(r)) \quad \text{for} \quad i = 1, \ldots, K.$$  (3.10)

This discrete system of equations may be expressed in matrix-operator form as

$$f(U) = S + H - (K + M + B)(U) = 0,$$  (3.11)

where the $K(U)$, $M(U)$, and $B(U)$ are operators (vector functions) corresponding to the stiffness (diffusion), mass (reaction), and boundary terms, respectively; $S$ and $H$ are the volumetric load vector and boundary load vectors, respectively; and $U$ is the vector of unknowns that approximates the solution in the domain $\Omega$. If the operators are evaluated using an appropriate linearization, the Jacobian matrix for the non-linear equations system is simply

$$\tilde{J}_{\text{elliptic}}(U) = -(K + M + B),$$  (3.12)

where $K$, $M$, and $B$ are now the stiffness, reaction, and boundary matrices (evaluated at the linearization point). Appropriate preconditioners for diffusion or reaction dominated problems based on the knowledge of the physics can also be created based on the above description of the spatial discretization for elliptic problems.
3.1.1.1 Boundary Conditions: Essential and Natural Conditions

Most often in BVPs, three boundary conditions, namely Dirichlet, Neumann and Robin, are employed. To preserve generality, let boundary $\Gamma = \Gamma_D + \Gamma_N + \Gamma_R$. It is neccessary to understand how these conditions need to be included in the variational formulation itself in order to avoid inconsistency in the discretization. The derivation above for non-linear Elliptic/Parabolic problems is general and does not tie itself down to any specific boundary condition. In this section, we will discuss the methods to impose these various conditions for second order elliptic problems for the boundary integral term in Eq. (3.6) with a continuous Galerkin (cG) discretization.

3.1.1.2 Dirichlet BCs

On $\Gamma_D$, the Dirichlet essential boundary conditions are specified as follows

$$u(\vec{r}, t) = \alpha(\vec{r}, t), \quad \vec{r} \in \Gamma_D$$  \hspace{1cm} (3.13)

There are several ways Dirichlet boundary conditions can be imposed. A simple approach, which works for most interpolary bases like the standard Lagrange polynomials used in the current work for continuous Galerkin discretization, is to assign function values Eq. (3.13) directly to the degrees of freedom on the domain boundary $\Gamma_D$. This idea of imposing Dirichlet conditions directly on the solution is 'strong' in the sense that it does not change the Dirichlet solution as a function of the mesh discretization.

Dirichlet conditions can also be imposed with a "penalty" method. In this approach, essentially the $L_2$ projection of the boundary values are added to the linear system matrix. The projection is multiplied by some large factor so that, in floating point numeric arithmetic, the existing (smaller) entries in the matrix
and right-hand-side load vector are effectively ignored. This leads to modifying the boundary terms $B(U), H(U)$ in Eq. (3.11) as

$$B_i(U) = \int_{\Gamma_D} \Phi_i \sum_k \Phi_k U_k (1 + \rho \delta_{ik})$$  \hspace{1cm} (3.14)

$$H_i(U) = \rho \int_{\Gamma_D} \alpha(r, t) \Phi_i$$  \hspace{1cm} (3.15)

where $\rho$ is the penalty parameter, such that $\rho >> 1$.

3.1.1.3 Neumann BCs

On $\Gamma_N$, the Neumann natural boundary conditions are specified as follows

$$D(\vec{r}, u) \frac{\partial u(\vec{r}, t)}{\partial n} = \beta(\vec{r}, t), \hspace{0.5cm} \vec{r} \in \Gamma_N$$  \hspace{1cm} (3.16)

These conditions are called ‘natural’ because they are imposed as part of the variational formulation itself. Consider the boundary term in Eq. (3.6) and applying the conditions Eq. (3.16),

$$\int_{\Gamma} v D(\vec{r}, u) \bar{\nabla} u \cdot \bar{n} ds = \int_{\Gamma} v \beta(\vec{r}, t) ds,$$  \hspace{1cm} (3.17)

where $\bar{n}$ is the outward unit normal vector on the boundary.

This can be seen as the boundary $L_2$ inner product on $\Gamma_N$. This contribution is added to the boundary operator $H(U)$ and is imposed weakly on the variational form.
3.1.1.4 Robin BCs

On $\Gamma_R$, the Robin or mixed boundary conditions are specified as follows

$$D(\vec{r}, u) \frac{\partial u(\vec{r}, t)}{\partial \vec{n}} + \gamma u(\vec{r}, t) = \beta(\vec{r}, t), \quad \vec{r} \in \Gamma_R. \quad (3.18)$$

Imposing these mixed boundary conditions is very similar to that of the Neumann conditions since it requires same modifications on the variational formulation. Again, take the boundary term in Eq. (3.6) and applying the conditions Eq. (3.18),

$$\int_{\Gamma} vD(\vec{r}, u) \vec{\nabla} u \cdot \vec{n} ds = \int_{\Gamma_R} v(\beta(\vec{r}, t) - \gamma_R u(\vec{r}, t)) ds. \quad (3.19)$$

This boundary contribution is added to the operators $H(U)$ and $B(U)$.

3.1.2 Hyperbolic Systems: Discontinuous Galerkin Discretization

Consider a non-linear hyperbolic conservation equation with advection and reaction of the form

$$\vec{\nabla} \cdot \vec{G}(u, \vec{r}, t) + c(\vec{r}, u) u(\vec{r}, t) = q(\vec{r}, t) \quad (3.20)$$

where $\vec{G}(u), c(\vec{r}, u)$ are smooth functions with $c(\vec{r}, u) \geq 0$ in $\Omega$ and $q \in L_2(\Omega)$ with boundary conditions specified on the inflow boundary $u(\vec{r}, t) = \alpha(\vec{r}, t), \forall \vec{r} \in \Gamma_i$, where $\vec{G}(u) \cdot \vec{n} < 0$ and $\vec{n}$ is the outward unit normal vector on $\Gamma_i$.

Then the Galerkin weak form of Eq. (3.20) is obtained by multiplying the equation with a test function $v$ and integrating over the domain $\Omega$, like in the elliptic case, to obtain

$$\int_{\Omega} v(\vec{\nabla} \cdot \vec{G}(u, \vec{r}, t) + c(\vec{r}, u) u(\vec{r}, t) - q) d\Omega - \int_{\Gamma} \vec{G}(u(\vec{r}, t), \vec{r}, t) \cdot \vec{n} v d\Omega \quad \forall \vec{r} \in \Omega. \quad (3.21)$$
Let the numerical solution $u$ be expanded in terms of basis functions (Legendre polynomials) $\Phi(r)$ that are discontinuous functions of order $p$, defined on the mesh Triangulation $T$ of $\Omega$, $\bigcup_{K \in T} K = \Omega$.

$$u_k(\vec{r}) \approx U_k(\vec{r}) = \sum_{i=1}^{p} U_i \Phi_i(\vec{r}), \quad (3.22)$$

where $U_i$ are the degrees of freedom of the FE discretization.

Eq. (3.21) can now be rewritten as

$$\sum \left\{ - \int_K \{ \vec{G}(u) \cdot \nabla v \} + \sum_{K} \int_K \{ c(r, \vec{u}) uv \} + \int_{\partial K} \{ \vec{G}(u) \cdot \vec{n} v \} \right\} = \sum \int_K \{ v S \}, \quad (3.23)$$

Because of the discontinuous nature of the solution approximation, the true flux $\vec{G}(\vec{u}) \cdot \vec{n}$ is not defined at the cell’s boundaries and this quantity is usually replaced by a numerical flux $H_K(u^+, u^-, \vec{n})$ which approximates $\vec{G}(\vec{u})$. Here, $u^\pm$ represents the traces on the boundary edges from the interior/exterior of an element $K$.

With the introduction of the numerical flux, the weak form for the dG method can be rewritten as

$$\sum \left\{ - \int_K \{ \vec{G}(u) \cdot \nabla v \} + \int_K \{ c(r, u) uv \} + \int_{\partial K} \{ H_{LLF}(u^+, u^-, \vec{n}) v^+ \} \right\} = \sum \int_K \{ v S \}, \quad (3.24)$$

where $H_{LLF}(u^+, u^-, \vec{n})$ is the Rusanov, or Local Lax-Friedrichs (LLF), numerical flux given by

$$H_{LLF}(u^+, u^-, \vec{n}) = \frac{1}{2} \left\{ \vec{G}(u^+) \cdot \vec{n} + \vec{G}(u^-) \cdot \vec{n} + \lambda(u^+ - u^-) \right\}, \quad (3.25)$$

with $\lambda$ the largest eigenvalue (in absolute value) of the Jacobian matrix of $\vec{G}$. For the 1-dimensional non-linear conservation law used to model fluid flow for reactor applications, the eigenvalue $\lambda = \sup \{ v_x, v_x + c, v_x - c \}$ where $v_x$ is the velocity in
the direction of flow and $c$ is the sound speed that depends on the medium pressure, density and temperature.

Alternately, an Upwind flux can be used instead of the Rusanov flux, where the numerical flux function is given as

$$H_{\text{up}}(u^+, u^-, \vec{n}) = \begin{cases} \vec{G}(u^+) & \text{if } \vec{u} \cdot \vec{n} \leq 0 \\ \vec{G}(u^-) & \text{otherwise} \end{cases}$$

(3.26)

As an aside, it is interesting to note that in higher order accurate dG methods, the choice of Riemann solver is not that crucial to resolve the spatial scales correctly [36]. Hence, in the current work, the Upwind and Rusanov flux function are used [37,38] as the solver since it is easy and less expensive to implement, whereas many other choices such as the Godunov, Roe, Osher, HLL, HLLC, and HLLE solvers are available. Future tests to affirm the conclusions of these previous results for problems occurring in nuclear reactor analysis will be necessary to validate the current choice of Riemann solver.

In operator notation, Eq. (3.24) can be written in a general form as

$$f(U) = G(U) + B(U) + M(U) - S = 0$$

(3.27)

where the $G(U), M(U),$ and $B(U)$ are vector operators corresponding to the advection, reaction, and boundary terms, respectively; $S$ is the volumetric load vector; and $U$ is the vector of unknowns that approximates the solution in the domain $\Omega$. If the operators are evaluated using appropriate linearization, the Jacobian matrix for the non-linear equations system is simply

$$\tilde{J}_{\text{hyperbolic}}(U) = -\frac{\partial(G + M + B)}{\partial U},$$

(3.28)

where $\frac{\partial G}{\partial U}, \frac{\partial M}{\partial U},$ and $\frac{\partial B}{\partial U}$ are the partial Jacobian of advection, reaction, and boundary operators respectively, evaluated at the linearization point. Forming the Jacobian
for the conservation law with higher order dG discretization can be expensive and complicated. But recent work on steady state problems [39] emphasizes that accurate evaluation of the Jacobian matrix can be crucial to speed up convergence of the non-linear Newton iteration. Also, since the numerical flux functions can be arbitrarily chosen for a given problem, it is only required that the derivatives of the numerical flux $H_u^+$ and $H_u^-$ need to be calculated to assemble the Jacobian matrix correctly. Analytic forms for the derivative are sometimes not available directly but a numerical finite difference procedure can be performed to obtain these values. For the Upwind and Rusanov fluxes, these values are straightforward to compute and the analysis for other types will be left for future work.

Apart from directly computing the Jacobian from the dG residual in Eq. (3.27), approximate Jacobian matrix for preconditioning the linear system can be obtained based on the Implicit Continuous Eulerian (ICE) technique [31], in which a semi-implicit linearization treats the advection operators explicitly. The unknowns are then eliminated through a Gaussian elimination and substitution process, yielding a single pressure-Poisson equation [28]. This formulation is widely used for low Mach flow regimes as a solver by itself and thus could be quite effective when utilized as a preconditioner within the non-linear matrix-free framework used in the current work. Detailed description of the linearized Jacobian matrix obtained via perturbation of the numerical flux and the ICE preconditioner is provided in Section. (3.3.3).

3.1.2.1 Boundary Conditions: Inflow and Outflow

For 1-dimensional conservation laws that resemble the inviscid Euler equations, there are three characteristic speeds corresponding to the eigenvalues of $G'(u)$ [40], namely

$$\lambda_1 = v_x - c, \quad \lambda_2 = v_x, \quad \lambda_3 = v_x + c$$

(3.29)
According to the sign of the these characteristics, four different boundary conditions are usually employed at the inflow and outflow boundaries.

1. Subsonic Inflow: $\lambda_i < 0, \quad i = 1, 2 \text{ and } \lambda_3 > 0$

2. Subsonic Outflow: $\lambda_i > 0, \quad i = 2, 3 \text{ and } \lambda_1 < 0$

The supersonic inflow and outflow conditions have not been considered here since the regimes that are dominant in reactor analysis problems for fluid flows are primarily subsonic.

In order to provide details on the application of these boundary conditions, notations regarding the boundary faces need to be specified. Let us first subdivide the boundary $\Gamma$ into the inflow boundary $\Gamma_i$ and the outflow boundary $\Gamma_o$. Then split the element boundary terms into interior and boundary face terms such that $\sum K \int_{\partial K} = \sum K \int_{\partial K \setminus \Gamma} \bigcup \sum K \int_{\partial K \cap \Gamma}$. Now define the bilinear form of the weak statement to include the boundary face terms as follows:

$$a_{\Gamma}(u, v) = \sum_{K \in \Gamma} \int_{\partial K \cap \Gamma} H(u^+, u^-, n)v^+ds$$  \hspace{1cm} (3.30)

This boundary term consists of two parts in our case:

$$a_{\Gamma}(u, v) = a_{\Gamma_i}(u, v) + a_{\Gamma_o}(u, v)$$  \hspace{1cm} (3.31)

Depending on the domain boundary, these terms are specified in the weak form as follows:

1. At the inflow boundary $\Gamma_i$, the outer trace $u^-$ is replaced by the given boundary function $g$ as

$$a_{\Gamma_i}(u, v) = \sum_{K \in \Gamma_i} \int_{\partial K \cap \Gamma_i} H(u^+, g, \vec{n})v^+ds$$  \hspace{1cm} (3.32)
2. At the outflow boundary $\Gamma_o$, only one characteristic variable need to be imposed. In many cases, the outflow variable specified is pressure $p = p_o$. Hence on $\Gamma_o$, the outer trace $u^-$ is replaced by a modified solution $u^- = u_o^-(u)$. Then

$$a_{\Gamma_o}(u, v) = \sum_{K \in \Gamma_h} \int_{\partial K \cap \Gamma_o} H(u^+, u_o^-(u), \vec{n}) v^+ ds$$

(3.33)

Often the modified solution depends on the inner trace $u^+$ and prescribed pressure $p_o$ such that $u_o^- = (\rho, \rho v, \rho E(\rho e, p_o))$.

The specification and implementation of these boundary terms are different from that for elliptic PDE. Even though the Dirichlet conditions are specified for each of the solution variables in the inflow boundary, imposing these conditions occur naturally through the use of the numerical flux functions. Even time-dependent Dirichlet conditions do not require special treatment in order to be enforced consistently.

### 3.1.3 Spatial Coupling Error in Multi-mesh Approaches

Often times in multi-physics applications, each physics component is discretized on its own mesh, and the solution field from a given physics needs to be exported onto another mesh. $L_2$ projection or interpolation of the solution between the source and target meshes may cause non-negligible spatial error [41]. In order to minimize the spatial coupling error due to the data transfer between the different physics defined on non-overlapping meshes, several techniques have been developed [42]. Jiao and Heath [43] have derived rigorous cost estimates for different remapping methods along with the solution costs. The spatial coupling error due to, for instance, the use of different meshes, is still an ongoing topic of research [44].

In the current research work, we employ high order quadrature rules for the numerical integration of the terms residing on the target mesh, that approximates the spatial integrals to capture the multi-physics solution behavior. This idea is
applicable for arbitrary meshes, provided that the solution for a physics can be evaluated at any point based on the expansion of the solution in terms of the basis functions. Then, as the number of quadrature points is increased, the multi-mesh coupling error becomes ‘small enough’ as compared to the non-linear error that is not resolved in the coupled physics solution.

For illustration, let us consider two physics, indexed by 1 and 2. In the weak formulation, the non-linear residual of physics 1, \( f_1(y_1, y_2) \) is multiplied by a test function, \( b_1^j \). The following integral needs to be computed accurately for every cell \( K_1 \) of physics 1:

\[
\int_{K_1} f_1(u_1(x), u_2(x)) b_1^j(x) dx.
\]  
(3.34)

Expanding the solution fields onto the basis functions, \( u_1(x) = \sum_i b_1^i(x) \hat{u}_1^i \) and \( u_2(x) = \sum_i b_2^i(x) \hat{u}_2^i \), and replacing the integral by a numerical quadrature \((w_q, x_q)\) yield

\[
\sum_q w_q f_1 \left( \sum_i b_1^i(x_q) \hat{u}_1^i, \sum_i b_2^i(x_q) \hat{u}_2^i \right) b_1^j(x_q).
\]  
(3.35)

For identical meshes, the values \( b_1^j(x_q) = b_2^j(x_q) \) are simple to obtain: mapping \( K_1 \) onto a reference element is advantageous since the basis functions need only to be evaluated once at the quadrature points of the reference element. However, when the meshes are different, (1) the numerical integration needs to be carried out on the physical element \( K_1 \), and, (2) all the cells of physics 2 overlapping \( K_1 \) need to be retrieved and the basis functions \( b_2 \) need to be evaluated at the quadrature points, \((x_q)\). For general unstructured meshes, one cannot obtain straightforwardly \( b_2^j(x_q) \) in the reference element since this involves reverse lookups to find the correct target element for physics 2 containing the physical point. Hence, the numerical integration over a cell is carried out on the real geometry (the actual cell itself), and not on its mapped reference element. Here, high order quadrature rules for each physics are employed along with inverse mapping of the meshes in different physics in order to evaluate the basis functions at the given physical points. This computation
is necessary each time the residual for a given physics needs to be evaluated and an efficient linked list data-structure is created to store the required information in memory and speed up the integration over cells.

The current work does not delve indepth into the issues related to coupling the solution fields from several completely different legacy codes developed independently. These scenarios have solutions residing in meshes that conform to spatial scales of each physics and the a-priori determination of the number of quadrature points to perform the $L_2$ projection accurately is difficult. A workaround would be to create a ‘super’ mesh which is the union of all the individual physics meshes given by $\Omega_{Super}^h = \Omega_1^h \cup \Omega_2^h \cup \ldots \cup \Omega_N^h$. Then, the solution at all the mesh points can be interpolated, projected and used uniformly with affordable loss of accuracy.

It is also important to note that, making use of available degrees of freedom, certain quantities such as total mass and energy need to be conserved through these projections [42]. This needs special attention while devising schemes to project these variables on a different mesh to be coupled with another physics. Since this subject in itself involves considerable research, only the ideas have been proposed here and demonstrations using two physics will be presented in Section. (5).

3.2 Time Discretization

Tackling the whole coupled non-linear system provides tremendous flexibility to use high-order implicit time integrators. Implicitness is required for stability due to the great disparity in time scales of the various phenomena involved in the simulations. Even though traditional codes dealing with stiff individual physics systems tend to use semi-implicit (treat fast scales implicitly and others explicitly), these schemes might not be as effective when used in the context of coupled physics problems due to the introduction of time scales that cause increased stiffness. But since the temporal treatment in single physics problems are based on intimate knowledge
of the physics, these solution schemes and discretizations serve as excellent preconditioners for fully coupled physics problems.

Consider a vector valued non-linear system of equations \( f \), that is obtained after appropriate spatial discretization using cG or dG FEM for the different physics. This non-linear residual includes all the coupling details, i.e., the contributions from one physics to another is accounted correctly. The large system of time-dependent coupled non-linear ODEs describing the problem can be generally written as

\[
M \frac{dU}{dt} = f(t, U). \tag{3.36}
\]

where \( M \) is the mass matrix resulting from the spatial discretization of the temporal derivative term (the use of a finite difference technique in space or a lumped numerical quadrature results in \( M \) being the diagonal matrix whose entries contain the cell volume).

The initial BVP has an initial solution prescribed at some given time \( t_{\text{initial}} \). Let the 1-dimensional time domain \( \Theta = [t_{\text{initial}}, t_{\text{final}}] \) be partitioned into \( N \) steps with \( \sum_{n=1}^{N} \Delta t_n = t_{\text{final}} \).

Without loss of generality, consider a Runge-Kutta (RK) method for temporal discretization represented using the standard Butcher tableaux notation. Then, any RK method can be specified using the following notation:

\[
\begin{array}{c|c}
C & A \\
\hline
& B^T \\
\end{array} \tag{3.37}
\]
where $B = [b_1, \ldots, b_s]^T$, $C = [c_1, \ldots, c_s]^T$, and $A = (a_{ij})_{i,j=1,\ldots,s}$. Let $s$ be the number of intermediary stages for the RK method, and $\Delta t_n$ the size of step $n$. The application of the RK method to Eq. (3.36) yields the solution at $t_{n+1}$ as

$$U_{n+1} = U_n + \Delta t_n \sum_{i=1}^{s} b_i k_i,$$

(3.38)

where the intermediate vectors $k_i (i = 1, \ldots, s)$ are obtained by solving the following $s$ non-linear systems

$$M k_i = f \left( t_n + \Delta t_n c_i , U_n + \Delta t_n \sum_{j=1}^{s} a_{i,j} k_j \right),$$

(3.39)

The above equation shows that the computation of a solution at $t_{n+1}$ involves performing at least $s$ non-linear iterations for one single sweep and it is necessary to converge the stage vectors $k_i$ in order to obtain the time solution at the end of $n^{th}$ step. Since the derivation is still general, no assumptions have been made about the structure of the Butcher matrix $A$ to simplify the equations. This will be dealt with separately once we have a fully discrete system of equations.

Based on ideas by Hairer [45], a simple substitution of variables is introduced next. Let

$$Z_i = \Delta t_n \sum_{j=1}^{s} a_{i,j} k_j.$$

(3.40)

Substituting Eq. (3.40) in Eq. (3.39), the modified set of $s$ non-linear problems is

$$M k_i = f ( t_n + \Delta t_n c_i , U_n + Z_i ),$$

(3.41)

and, by recursion after simplification, this yields the modified non-linear ‘temporal’ residual equation defined by

$$F(Z) = (M \otimes I_s)Z - \Delta t_n A f( t_n + \Delta t_n C , U_n + Z ) = 0,$$

(3.42)
where $Z = \{Z_1, \ldots, Z_s\}$ and $f(Z) = \{f(Z_1), \ldots, f(Z_s)\}$.

Now that we have arrived at a final non-linear system, the solution to Eq. (3.42) for $Z$ can be obtained by some form of non-linear iteration, using either Picard or Newton method. Once $Z$ is found and converged for all $s$ stages, we can substitute in Eq. (3.38) to find the solution at end of time level $n$ using

$$MU_{n+1} = MU_n + \Delta t_n B^T f(t_n + \Delta t_n C, U_n + Z),$$

(3.43)

It is important to note that all derivations leading up to Eq. (3.43) are applicable to explicit and fully-implicit RK methods. Then, the selection of the appropriate RK methods that can handle stiff PDEs [46, 45, 28] is necessary in order to obtain high-order accurate solutions using the above discretization method. These choices are usually based on several optimal properties of the RK methods such as:

1. explicitness vs implicitness.

2. $A$-stability, (absolute stability) determines whether a method is conditionally stable or unconditionally stable for all time step sizes $\Delta t_n$ [47] (i.e., it is the domain $S \in \mathbb{R}(z)$ such that $S = \{z \in \mathbb{C}; |\mathbb{R}(z)| \leq 1\}$ where $\mathbb{R}(z)$ is the method’s characteristic polynomial applied to Dahlquist’s equation $y' = \lambda y$ and $z = \Delta t_n \lambda$). In coupled physics systems, the disparity in the time scales leads to stiff systems that require $A$-stable methods in order to resolve the behavior of the physics correctly.

3. $L$-stability, is an essential property that indicates the rate of damping of highly oscillatory modes independent of time step size [48] i.e., a method is $L$-stable if it is $A$-stable and $\lim_{z \to \infty} \mathbb{R}(z) = 0$. This property is crucial to determine the success of a given method for stiff systems since if all modes are not damped quickly over a transient, the solution procedure can become unstable due to oscillations, necessitating smaller time steps.
4. Efficiency: cost of solution method per time step. This is critical since there needs to be a balance in terms of cost per step (\( s \times \text{Average CPU cost per stage} \)) versus accuracy in solution (Local Truncation Error (LTE)) for solving the system.

A RK method of order \( p \) with \( s \) stages can be compared to the actual Taylor series expansion of a non-linear system, to derive the order conditions. For higher order methods, it gives the user great flexibility in deriving a scheme with optimal order and stability properties to fit the needs of the problem. This plasticity of the method and the ease of adjusting the coefficients to obtain embedded formulas make them attractive to adaptive time-stepping when needed.

Next, specializations for different families of RK schemes will be discussed and the specific changes in the non-linear equation Eq. (3.42) and step solution Eq. (3.43) will be shown.

### 3.2.1 Explicit-RK (ERK) Methods

If the Butcher coefficient matrix \( A \) is strictly lower diagonal, i.e., \( a_{i,j} = 0, \forall i = 1 \ldots , s, j \geq i \), then the RK method is said to be explicit. This is because the solution for any time step explicitly depends only on the previous solution and stages and hence these methods do not require any non-linear iterations.

All explicit methods are conditionally stable but due to the reduced cost in finding the solutions, they could be valuable when the physics dictates the usage of very small time steps to resolve the temporal scales. This ‘asymptotic regime’, when the user-specified tolerance for LTE dominates the solution, is suitable for the usage of such schemes.

\[
\mathbf{M} \mathbf{k}_i = f \left( t_n + \Delta t_n c_i , \mathbf{U}_n + \Delta t_n \sum_{j=1}^{i-1} a_{i,j} \mathbf{k}_j \right) \quad \forall i = 1 \ldots , s \quad (3.44)
\]

\[
\mathbf{U}_{n+1} = \mathbf{U}_n + \Delta t_n \mathbf{B}^T \mathbf{K} \quad (3.45)
\]
where \( K = \{k_1, \ldots, k_s\} \).

Hence ERK methods are easy to implement and have cheap computational cost per step since there are no non-linear iterations or Jacobian matrix solves other than the Mass matrix \( M \) at the end of each stage. However, they have poor stability properties and are unable to resolve very fast changing modes (explicit schemes are not suitable for stiff equations). To overcome this problem and to utilize the advantages of these one step schemes, modifications to the existing ERK schemes can be made, as shown by Eriksson et. al. [49], to extend the stability region.

In the current work, for the sake of completeness, we have chosen to implement Forward Euler (FE), a two stage ERK method of order 2 and the four stage ERK method by Kutta based on \( 3/8^{th} \) Quadrature Rule of order 4. Apart from these standard schemes, an embedded ERK method, DOPRI by Dormand-Prince [46] with stiffness detection, has been implemented as well.

The notation for naming each of the RK methods is usually given as RK\( p, p'(s) \) where \( p \) is the true order of the method, \( p' \) is the embedded order and \( s \) is the number of stages. With this notation, the Butcher Tableaux for each of the above methods are given below.

\[
\begin{array}{c|c}
0 & 0 \\
\hline
B^T & 1 \\
\end{array}
\]

FE 1(1) \hspace{1cm} (3.46)

\[
\begin{array}{c|c}
0 & 0 \\
\hline
\frac{2}{3} & \frac{2}{3} \\
\end{array}
\]

\[
\begin{array}{c|c}
B^T & \frac{1}{4} \quad \frac{3}{4} \\
\end{array}
\]

ERK 2(2) \hspace{1cm} (3.47)
In Eq. (3.49), $E^T$ is the error estimator coefficient that is useful in obtaining the Local Truncation Error (LTE) for the specified RK method. This is derived along with the optimal higher order ($p$) step coefficients $B^T$ for the embedded method. Then,

$$E^T = B^T - \tilde{B}^T$$

(3.50)

where $\tilde{B}^T$ are the coefficients for the lower order ($p'$) method. For brevity, $\tilde{B}^T$ have not been shown and can be easily obtained if necessary.
The LTE ($\epsilon$) for such an embedded method is

$$\epsilon_n = \Delta t_n E^T K + O(\Delta t_n^{p+1}) \quad (3.51)$$

A-priori estimates for the LTE are useful to create adaptive solution procedures that can change the step size $\Delta t_n$ and order $p$ of the method to reduce the local and global temporal error in the solution based on user specified tolerance. Based on principles in control theory, Gustafsson [50] introduced the PI controller and applied it to adaptive step-size selection for stiff ODE problems. Previous work for reactor problems [11] using these adaptive controllers were successful and hence have been used in the current research for use with embedded methods.

The PI controller predicts the new step size based on the evolution in LTE, the selection of previous step size and a user specified tolerance. Then,

$$\Delta t_{n+1} = \Delta t_n \left( \frac{Tol}{|\epsilon_n|} \right)^{\alpha} \left( \frac{|\epsilon_{n-1}|}{|\epsilon_n|} \right)^{\beta} \quad (3.52)$$

where $\alpha$ and $\beta$ are problem dependent constants. Gustafsson found after some numerical computation that $\alpha \approx \frac{0.7}{\min|p, p'|+1}$ and $\beta \approx \frac{0.4}{\min|p, p'|+1}$ are usually good choices for stiff problems. The paper cited above provides detailed derivation of the controller and the optimal parameters in Eq. (3.52).

### 3.2.2 Implicit RK (IRK) Methods

Implicit methods are either usually unconditionally stable ($A$-stable) or at least have much larger stability regions than ERK methods. Even if an IRK method is $A$-stable, it may not satisfy the required $L$-stability conditions that are essential to accurately resolve stiff systems of equations. We can also classify IRK methods based on the structure of the Butcher matrix $A$ into two broad categories. We will discuss
each family below along with the implication on the cost for obtaining a solution per time level.

3.2.2.1 Diagonally-Implicit RK (DIRK) Methods

For DIRK methods, the Butcher coefficient matrix $A$ is lower diagonal, i.e., $a_{i,j} = 0, \forall i = 1, \ldots, s, j > i$. Note that the diagonal term is non-zero and hence the solution at each stage requires an implicit non-linear solve, unlike with ERK methods.

The equations for the simplified non-linear system Eq. (3.42) at each stage can be modified as

$$
F(Z_i) = MZ_i - \Delta t_n \sum_{j=1}^{i} a_{i,j} f(t_n + \Delta t_n c_i, U_n + Z_i) = 0 \quad \forall i = 1, \ldots, s
$$

Then, if the Jacobian matrix $J(U)$ for the SS residual $f(U)$ can be computed approximately, the non-linear iteration to compute the solution update proceeds as

$$
\hat{J}(Z_l^i) \delta Z_l^i = -F(U_n + Z_l^i) \quad \forall i = 1, \ldots, s
$$

$$
Z_l^{i+1} = Z_l^i + \delta Z_l^i
$$

where $l$ is the non-linear iteration index and the transient Jacobian matrix $\hat{J}(Z)$ is

$$
\hat{J}(Z_l^i) = M - \Delta t_n a_{i,i} J(t_n + \Delta t_n c_i, U_n + Z_l^i) \quad \forall i = 1, \ldots, s
$$

Now that we have determined the necessary components to solve the transient non-linear system, the solution to Eq. (3.54), Eq. (3.55) for $Z = \{Z_0, \ldots, Z_s\}$ can be obtained. The use of either a Picard or Newton method as a non-linear solver will be discussed in the next section and the focus will be shifted to solve this system efficiently under constraints of memory and time.
Once $Z$ is found and converged for all $s$ stages, we can substitute in Eq. (3.38) to find the solution at end of time step $n$. This step involves inverting the Mass matrix $M$ which can be performed using a lumped-mass approach [51] that has been proven to be quite effective for several test problems.

Several DIRK methods possess unconditional stability and optimal properties that help improve the efficiency of solution procedure. For instance, it is advantageous to have the diagonal elements of the Butcher matrix $A$ to be the same i.e., $a_{i,i} = \gamma$. These DIRK methods are popularly called Singly-DIRK (SDIRK) methods. A variation of the SDIRK methods with an explicit first stage, Explicit SDIRK (ESDIRK), was investigated introduced by Kvaerno [52] and investigated further by Kennedy et al. [53] for advection-diffusion-reaction equations. These methods simplify the solution procedure to solving the non-linear system given in Eq. (3.54) since the transient Jacobian matrix $\hat{J}(t, U)$ that needs to be inverted is the same in all stages. Hence if a direct method such as $LU$ factorization can be used, then the factorization need be performed only once and utilized for all the stage computations. Note that in this case, the Jacobian is also lagged (computed at start of step).

Based on the analysis of the properties of DIRK methods, few of them have been chosen to be implemented: Backward Euler (BE), Implicit Midpoint (IM), SDIRK2(2), SDIRK3(2), SDIRK3(3) [45]. Note that BE, SDIRK2(2), SDIRK3(3) are $A-$, $L-$ stable schemes but IM2(1) and SDIRK3(2) are only $A-$stable and not $L-$stable. Since the provision for including arbitrary DIRK methods exists in the framework introduced thus far, any DIRK/SDIRK method that can be represented by a Butcher Tableau can be tested and used in the software implemented as part of the current work.

\[
\begin{array}{c|c}
1 & 1 \\
B^T & 1 \\
\end{array}
\]

BE 1(1)
\[
\begin{array}{c|cc}
0.5 & 0.5 \\
\hline
B^T & 1 \\
\end{array}
\]  
(3.58)

IM 2(1)

\[
\begin{array}{c|ccc}
\gamma & \gamma & \\
1 & 1 - \gamma & \gamma \\
\hline
B^T & 1 - \gamma & \gamma \\
\end{array}
\]  
(3.59)

SDIRK 2(2) with \( \gamma = 1 - \frac{1}{\sqrt{2}} \)

\[
\begin{array}{c|ccc}
\gamma & \gamma & \\
1 - \gamma & 1 - 2\gamma & \gamma \\
\hline
B^T & 0.5 & 0.5 \\
\end{array}
\]  
(3.60)

SDIRK 3(2) with \( \gamma = \frac{3 - \sqrt{3}}{6} \)

\[
\begin{array}{c|ccc}
\gamma & \gamma & \\
\frac{1 + \gamma}{2} & \frac{1 - \gamma}{2} & \gamma \\
1 & -\frac{6\gamma^2 + 16\gamma - 1}{4} & \frac{6\gamma^2 - 20\gamma + 5}{4} \\
\hline
B^T & -\frac{6\gamma^2 + 16\gamma - 1}{4} & \frac{6\gamma^2 - 20\gamma + 5}{4} \\
\end{array}
\]  
(3.61)

SDIRK 3(3) with \( \gamma = 0.435866521508459 \)

3.2.3 Fully-Implicit RK (FIRK) Methods

FIRK methods have a full Butcher coefficient matrix, i.e., \( a_{i,j} \neq 0, \forall i, j = 1 \ldots s \). One way to solve these systems would be to consider the full block non-linear system, all unknowns from the \( s \) stages, i.e., \( Z \) is the unknown instead of \( Z_i \) for individual
stages, and perform non-linear iterations on these. Due to memory restrictions for large scale fully discretized problems, this could be prohibitive.

Alternately, an outer iteration can be used in conjunction with ideas for solving DIRK methods, in order to converge the temporal step solution. This procedure is based on splitting the block matrix operator as $A = D + L + U$ where $D, L, U$ are the diagonal, strictly lower triangular and strictly upper triangular terms of the coefficient matrix. With this splitting, a Block Gauss-Seidel (BGS) iteration can be applied to obtain the residual as

$$ F(Z^l_{ibgs}) = MZ^l_{ibgs} - \Delta t_n (L + D) \otimes I_n f(t_n + \Delta t_n C, Z^l_{ibgs}) - \Delta t_n U \otimes I_n f(t_n + \Delta t_n C, Z^l_{ibgs-1}) = 0 $$

with the transient Jacobian matrix $\hat{J}(t, U)$ given by

$$ \hat{J}(Z^l_{ibgs}) = M - \Delta t_n (L + D) \otimes I_n J(t_n + \Delta t_n C_i, Z^l_{ibgs}) $$

where $ibgs$ is the BGS iteration number. This iteration can also be relaxed to improve the outer iteration convergence using block SOR scheme but due to the difficulty in determining the optimal relaxation factor for all multi-physics problems, this is left for future work.

Simply put, the solution procedure for FIRK methods involves performing multiple DIRK solves until convergence. Hence the cost of these methods is cost per DIRK step * Number of outer iterations. Due to the cost involved in computing the solution for FIRK methods, this is often not preferred unless extremely stiff problems are encountered.

Hairer [45] notes that collocation methods based on Gauss and Radau quadrature formulas can lead to FIRK methods with excellent stability properties. These methods are in general $A$- and $L$- stable and stiffly accurate (do not degrade convergence for stiff problems) [54].
An adaptive method using the RADAU5 scheme with a good error estimator was implemented previously \[11\] for coupled simulations using Point Reactor Kinetics Equations (PRKE) and lumped hydraulics models and the success of these methods in predicting sudden changes in temporal scales make them attractive. The use of such implicit adaptive techniques will be essential to capture complex waxing and waning of temporal scales from different physics during critical transients \[26\] and needs further investigation.

The Butcher matrix for the fourth order methods based on Gauss quadratures and third, fifth order methods based on RADAU IIA family are given below.

\[
\begin{array}{ccc}
\frac{1}{2} - \frac{\sqrt{3}}{6} & \frac{1}{4} & \frac{1}{4} - \frac{\sqrt{3}}{6} \\
\frac{1}{2} + \frac{\sqrt{3}}{6} & \frac{1}{4} + \frac{\sqrt{3}}{6} & \frac{1}{4} \\
\end{array}
\]

\[B^T\]

\[
\begin{array}{cc}
\frac{1}{2} & \frac{1}{2} \\
\end{array}
\]

Gauss 4(2)

\[
\begin{array}{ccc}
\frac{1}{3} & \frac{5}{12} & -\frac{1}{12} \\
1 & \frac{3}{4} & \frac{1}{4} \\
\end{array}
\]

\[B^T\]

\[
\begin{array}{cc}
\frac{3}{4} & \frac{1}{4} \\
\end{array}
\]

Radau IIA 3(2)

\[
\begin{array}{cccc}
\frac{4-\sqrt{6}}{10} & \frac{88-7\sqrt{6}}{360} & \frac{296-169\sqrt{6}}{1800} & \frac{-2+3\sqrt{6}}{225} \\
\frac{4+\sqrt{6}}{10} & \frac{296+169\sqrt{6}}{1800} & \frac{88+7\sqrt{6}}{360} & \frac{-2-3\sqrt{6}}{225} \\
1 & \frac{16-\sqrt{6}}{36} & \frac{16+\sqrt{6}}{36} & \frac{1}{9} \\
\end{array}
\]

\[B^T\]

\[
\begin{array}{ccc}
\frac{16-\sqrt{6}}{36} & \frac{16+\sqrt{6}}{36} & \frac{1}{9} \\
\end{array}
\]

Radau IIA 5(3)

Until now, the subject of obtaining the solution of a non-linear system was only briefly discussed. This is because the crux of the work in the temporal solution
procedure lies solely in this non-linear solve. Details regarding the usage of Picard or Newton iteration as non-linear solvers are provided next.

3.3 Methods for Solving Large-scale Non-linear Systems

This section discusses the numerical techniques employed for solving the non-linear equations arising from the fully discretized coupled physics system. By controlling how the non-linearities are resolved, a tight coupling or traditional loose coupling paradigm can be obtained. This allows testing existing coupling strategies and comparing to new tightly coupled methods in terms of accuracy and efficiency since all of these methods can be implemented within the same framework.

The basis for this idea stems from the fact that if the linear operator representing the Jacobian matrix used for solving the non-linear system is block-diagonal, it represents the decoupled treatment of the different physics and collapses to a Picard iteration strategy. This procedure can be iterated to any given tolerance as long as the spectral radius of the linearized operator is less than one i.e., $\rho(\hat{J}) < 1$. In other words, the convergence through Picard iterations for coupled physics problems is guaranteed if the eigenmodes due to the linearized terms are not dominant. Then, these iterations are a natural formulation for weakly coupled physics models. But if $\frac{\rho(J)}{\rho(\hat{J})} > 1$ where $J$ is the consistent fully coupled Jacobian matrix, then the physics are strongly coupled and much smaller time step sizes will be necessary in order to make the linearization valid. Hence, with a combination of time step control and appropriate linearizations, such iterative procedures over the different physics can produce tightly coupled solutions.

The current framework employs Picard or Newton methods (outer non-linear solves) and Krylov methods (inner linear solves) to solve the set of discrete non-linear equations effectively and accurately. The Matrix-Free (MF) approximation can be included such that the algorithm can be implemented without explicitly building the Jacobian matrix needed in the linear solve. Often, building the Jacobian matrix can
be costly in CPU time and memory, especially when different physics components reside in multiple codes. The MF nature of the solvers relies on (i) the fact that Krylov solvers build a solution subspace using only matrix-vector operations and (ii) these matrix-vector operations can be approximated using a finite difference formula that does not require knowledge of the matrix elements at all. Nevertheless, Krylov methods may require a certain number of basis vectors to be stored in order to find an accurate solution (i.e., the size of the subspace may be large). The Krylov space size and the overall computing time can be significantly reduced by the use of an appropriate preconditioner for the linear solves. Therefore, the MF non-linear algorithm consists of 3 levels of iterations:

1. Nonlinear iteration,

2. Linear iteration,

3. Preconditioner iteration.

Since the equations and the methods provided here are generic and are applicable to arbitrary non-linear systems, the same scheme can be utilized for solving linear, non-linear single- and multi-physics coupled systems. The following subsections provide details on the three levels of iterations that are part of the framework used to perform these multi-physics simulations.

3.3.1 Nonlinear Iteration Methods

Consider a system of non-linear equations of the form

\[ F(Z) = 0 \]  

(3.68)

obtained by space-time discretization of a problem with \( \xi \) physics components coupled non-linearly to each other, leading to a system of ordinary differential equations.
Let us apply the traditional Picard iteration and the Newton iteration introduced earlier to solve the fully-discrete non-linear problem.

3.3.1.1 Picard Iteration

Picard iteration, also known as Fixed Point Iteration (FPI), is a viable and an easy method to implement since it makes use of existing OS coupling paradigm to linearize the coupled physics solution terms. In solving differential equations, Picard iteration is a constructive procedure for establishing the existence of a solution to a discretized system of equations Eq. (3.68), that passes through the fixed point \((Z_0)\).

However, it is not very effective to iterate at every time step to converge the non-linearities in order to restore the higher convergence order. This is due to the fact that the Picard iterations are only linearly convergent and hence the scheme converges slowly to the true solution. Such a solution procedure takes a high iteration cost and usually requires longer computation times. Additional modifications could accelerate the rate of convergence for the vanilla non-linear Picard iterations in order to make it a viable candidate for reactor analysis problems. Schemes such as Steffensen \([55]\) and vector Wynn-Epsilon algorithm \([56]\) can be used to accelerate the convergence rate of the sequence of vectors found using Picard iterations.

Also, by the nature of the Picard linearization, the coupling between the different physics are treated explicitly. The system matrix arising from the space-time discretization of these physics reflect this weak coupling between different physics components. Let \(Z_P\) be the solution fields corresponding to a particular physics \(P\). Then, the non-linear residual equation describing the Picard linearization for each physics \(P\) can be written by splitting the non-linear contributions from each physics, as:

\[
F(Z^{\ell+1}, Z^{\ell}) = \{Z^{\ell+1}_P - N_{PP}(Z^{\ell+1}_P)\} - \sum_{\substack{P' = 1 \atop P' \neq P}}^\xi \tilde{N}_{PP}(Z^{\ell})
\] (3.69)
where \( \ell \) is the Picard iteration number, \( N_{PP}(Z_P) \) represents the non-linear residual describing the individual physics and \( \tilde{N}_{PP}(Z) \) represents the non-linear residual due to the coupling of physics \( P \) with physics \( P' \).

Since the diagonal coupled physics terms \( P' \) are computed at the previous Picard iterate, the new solution can be obtained by performing the following sequence of iterations:

\[
\begin{align*}
J_{FPI}(Z^\ell)\delta Z^\ell &= -F(Z^{\ell+1}, Z^\ell) \quad (3.70) \\
Z^{\ell+1} &= Z^\ell + \delta Z^\ell \quad (3.71)
\end{align*}
\]

where \( J_{FPI} \) is simply in this case,

\[
J_{FPI}(Z) = \begin{pmatrix}
N_{11} & 0 & \cdots & 0 \\
0 & N_{22} & \cdots & 0 \\
\vdots & \vdots & \ddots & 0 \\
0 & \cdots & 0 & N_{\xi\xi}
\end{pmatrix} \quad (3.72)
\]

Since the blocks \( N_{ii} \) require only the solution to the single physics itself, this fixed point iteration procedure can be continued to generate a sequence of solutions that converge to the true coupled physics fields \( Z_P \). This Picard iteration procedure has a Jacobian matrix that is Block-Jacobi structured and hence could be feasible to couple existing mono-physics codes. This OS coupling paradigm uses schematically represented by Fig. 1.1 in Section. (1).

The Picard iteration over multiple physics explained above is the least efficient and computationally expensive mode for performing multi-physics simulations although it is easy to implement for coupling existing legacy codes. Alternately, any level of tighter coupling can be enforced by accounting for the knowledge gained about the physics. These variations in Picard linearization involve simply evaluating the non-linear contribution \( \tilde{N}_{PP} \) from physics \( P' \rightarrow P \) at the current iterate.
solution $Z^{\ell+1}$ in Eq. (3.71) and correspondingly including the implicit contribution of the non-linear operators in the Jacobian matrix Eq. (3.72). These modified Picard variants are usually made such that the Jacobian matrix can be represented as a Block-Lower-Triangular or Block-Upper-Triangular matrix which would involve Block-Backward/Forward substitution respectively, in order to obtain the solution for the Picard iteration Eq. (3.71). A representation of the Block-Lower-Triangular Picard linearized matrix is given below.

$$ J_{FP1}(Z) = \begin{pmatrix} N_{1,1} & 0 & \cdots & 0 \\ N_{2,1} & N_{22} & \cdots & 0 \\ \vdots & \vdots & \ddots & 0 \\ N_{\xi,1} & \cdots & N_{\xi,\xi-1} & N_{\xi\xi} \end{pmatrix} \tag{3.73} $$

### 3.3.1.2 Newton Iteration

Instead of employing Picard iterations, one can apply Newton’s method to solve the non-linear system of equations in Eq. (3.68) and obtain the solution iteratively as follows:

$$ J(Z^{\ell})\delta Z = -F(Z^{\ell}) \tag{3.74} $$

$$ Z^{\ell+1} = Z^{\ell} + \delta Z \tag{3.75} $$

where $J(Z^{\ell}) = \frac{\partial F(Z^{\ell})}{\partial Z}$ is the Jacobian matrix of the system at the current Newton iterate $Z^{\ell}$, $\delta Z$ is the increment update, solution of the linear solve, and the next Newton iterate is given by $Z^{\ell+1}$.

It is clear that the Eq. (3.74) requires forming the Jacobian matrix explicitly in order to solve the system for $\delta Z$. In the case where the coupling between the different physics is complex and requires more memory storage, this option may
not be feasible. Also, the convergence of Newton’s method strongly depends on the consistency of the Jacobian matrix with respect to the residual description.

One may compute a numerical approximation to the Jacobian, based on a finite difference procedure by perturbing $F(Z)$. Provided that enough memory is available, $J$ can be built element by element or column by column. This is usually referred to as the numerical Jacobian. If recomputed at every Newton iteration, it is very expensive in terms of computational time, especially if the size of the non-linear system $N$ is quite large since $F(Z)$ needs to be perturbed at least $N$ times. The cost of this numerical Jacobian is hence $O(N)$ non-linear residual function evaluations.

Alternately, when storing the entire Jacobian is not feasible due to memory constraints or when the computational cost of forming the numerical Jacobian itself is prohibitive, a matrix-free approach is preferred. Based on the ideas by Brown and Saad [22], the Jacobian-free approach can be used to efficiently tackle the non-linear system where the linear solve can be performed with only the action of the Jacobian matrix on a given vector. Using only this defined operation, the linearized system in Eq. (3.74) can be solved using an efficient linear solver.

Generally speaking, the action of the Jacobian on a given vector $v$ can be computed using the following finite-difference approximation:

$$ Jv \approx \frac{F(Z + \epsilon v) - F(Z)}{\epsilon} \quad (3.76) $$

where $\epsilon$ is a parameter used to control the magnitude of perturbation.

Note that the accuracy of the approximation depends strongly on the choice of $\epsilon$. A typical simple choice is usually the square root of machine precision $\epsilon^2 = \Upsilon \approx 1E - 16$. Other optimal equations for choosing the perturbation parameter $\epsilon$ have been derived in the reference papers [22, 23]. For completeness, this optimal form of $\epsilon$ is given by

$$ \epsilon = \frac{\sqrt{(1 + \|Z\|)}\Upsilon}{\|v\|} \quad (3.77) $$
Further analysis done on the optimization of this finite difference parameter by 
Xu [57], in the context of coupled multi-physics problems, can also be useful to
determine the error in the approximation and increase the efficiency of the algorithm
explained above.

Other types of finite difference procedures such as, two-sided difference formulas
instead of the one-sided difference formula used in Eq. (3.76), can increase the ac-
curacy of the approximation. But such modifications involve extra computational
work and increase the number of function evaluations needed for better estimations.
Hence, we have only considered the one-sided difference approximation in this cur-
cent work and the applicability of these alternate Jacobian-free approximations can
be analysed in the future.

The exact Newton method involves solving the linear system in Eq. (3.74) exactly,
i.e., to a tight tolerance at every Newton iteration. This is a waste of computational
effort when the solution to the non-linear problem is far away from the bowl of
asymptotic convergence. Hence, an adaptive technique to change the linear tolerance
in the Newton iteration based on the non-linear residual amplitude can decrease the
CPU cost during the initial stages of the iteration. Such a formulation is super-
linearly convergent and approaches quadratic convergence in the asymptotic regime.
The linear tolerance for this inexact Newton iteration can be generally chosen as

\[
||\text{LinearResidual}|| = ||J(Z^\ell)\delta Z^\ell + F(Z^\ell)||_2 < \gamma ||F(Z^\ell)||_2
\]  

(3.78)

where \(\gamma\) is a forcing term, generally chosen to be smaller than unity. Generally the
choice of \(\gamma\) results in a tradeoff in the number of non-linear iterations versus linear
iterations since too large a value results in more Newton iterations or even divergence
and too small a value results in more time spent in the linear solver. Several strategies
for optimizing the computational work with a variable ‘forcing term’ \(\gamma\) are given in
the work by Eisenstat and Walker [58]. Due to the potential savings in this inexact
Newton strategy coupled with the Jacobian-free formulation, this non-linear iteration
scheme to solve the coupled non-linear multi-physics problem will be used as the primary solver algorithm in the current work. Note that as $\gamma \to 0$, one recovers the exact Newton algorithm.

In addition to the basic inexact Newton iteration, line search strategies to obtain the global solution satisfying the non-linear system can be used. Such modifications can avoid local stagnation and helps to stabilize Newton’s method by scaling the update appropriately. This modification is of the form

$$Z^{\ell+1} = Z^\ell + d^\ell \delta Z^{\ell+1}$$

where $d^\ell$ is the scaling factor that restricts the update. The standard Newton algorithm is recovered when $d^\ell = 1$. Further reading regarding these global line search methods is available in [22, 59, 58]. The methods for linear systems arising in Eq. (3.71) and Eq. (3.74) are discussed next.

### 3.3.2 Krylov Methods for Solving Linear Systems

The linear system obtained from the Picard or Newton linearization applied to the non-linear equation Eq. (3.68) can be efficiently solved using a Krylov method in which an approximation to the solution of the linear system is obtained by iteratively building a Krylov subspace of dimension $m$ such that

$$K(v, J) = \text{span}\{v, Jv, J^2v, J^3v, \ldots, J^{m-1}v\}$$

where $v$ is the initial Krylov vector.

Most coupled multi-physics problems produce linear systems that are block unsymmetric, even if the individual blocks may be symmetric due to the type of spatial discretization, e.g., Continuous Galerkin for elliptic problems. Hence robust Krylov methods are needed to tackle these unsymmetric systems. Previous studies on the
usage of GMRes (Generalized Minimum RESidual), BiCGStab (Bi-Conjugate Gradient Stabilized) and Transpose-free Quasi Minimal Residual (TFQMR) methods to tackle such systems [60] in the context of non-linear multi-physics problems suggest the feasibility of these choices.

In the current work, since a general framework is required to solve large-scale coupled linear systems, the robustness of the linear solver is an important factor in determining the total computational time of the algorithm. It is also necessary that the linear solver used be insensitive to the numerical roundoff and finite difference errors that are created as part of the approximations used in the Jacobian-free formulation. It is worthwhile to note that the use of Arnoldi-type of Krylov iterative methods yields the best convergence since complete orthogonalizations of all the subspace vectors aids in correcting the numerical errors introduced by the finite difference approximation. Although it is not possible to select one efficient linear solver for all types of unsymmetric problems, such an Arnoldi based GMRes solver is expected to be reliable and provide monotonically decreasing residuals.

The success of the GMRes iterative method, introduced by Saad and Schultz [61], and its popularity due to its efficiency in solving nonsymmetric system of equations make it attractive for the usage in tightly coupled multi-physics systems. The GMRes algorithm generates a sequence of orthogonal vectors, and because the matrix being inverted is not symmetric, short recurrence relations cannot be used as in the case of the Conjugate Gradient algorithm. Instead, all previously computed vectors in the orthogonal sequence have to be retained. In current study, the modified Gram-Schmidt algorithm for orthogonalization is used instead of the classical Gram-Schmidt algorithm in order to create a stable solver that is insensitive to roundoff errors. In the GMRes algorithm, one matrix-vector product is required per iteration and the matrix-free approximation introduced earlier in Eq. (3.76) can be used to obtain the action of the Jacobian matrix on any vector. Detailed information on
the exact numerics and implementation of GMRes in the MFNK framework can be found in [23].

The cost of the GMRes algorithm strongly depends on the size of its Krylov subspace that is created through the matrix-vector products. The memory cost increases linearly with every iteration and the number of Inner-Products (IP) required for orthogonalization increases quadratically. Hence, when solving large systems of equations, it is necessary to limit the size of Krylov subspace used. To limit the Krylov subspace size, a restarted variant of GMRes algorithm, GMRes\(r\), where \(r\) is the size of Krylov space, can be employed.

Flexible versions of the restarted GMRes algorithm, FGMRes\(r\), are useful in cases where the matrix-vector products are computed inexactly, and a need for robust Krylov solvers that can provide monotonic convergence to the solution is necessary. FGMRes\(r\) algorithm differs from the standard preconditioned GMRes\(r\) implementation by allowing variations in preconditioning at each iteration. This is especially important since the preconditioned solve at each Krylov iteration is performed inexactly (varying number of iterations or tolerance for each preconditioner solve). Because of these advantages, in the current research, FGMRes\(r\) is the preferred linear solver for unsymmetric systems of equations.

Optimizations beyond restricting the size of the subspace \(r\) due to memory reasons involve reducing the total number of linear iterations through the use of appropriate preconditioners. A discussion of the preconditioner implementations and the options available for different kinds of physics is provided next.

3.3.3 Preconditioners for the Linear Iteration

The preconditioner \(P\) is usually a good approximation of the Jacobian and should be easier to form and solve as compared to the Jacobian matrix itself. The inherently two-step process for this stage requires the computation of the action of \(P^{-1}\) on any vector \(v\), rather than actually forming the preconditioning matrix itself. This
algorithm can be made strictly Matrix-Free and studies for real-world problems previously [27] have shown possible increased efficiency when using this approach.

The right-preconditioned Matrix-Free Nonlinear-Krylov (MFNK) algorithm which involves using Eq. (3.76) for the Jacobian-vector products and an appropriate numerical or physics-based preconditioner results in a modified form of the non-linear iteration. The right-preconditioned non-linear equation is given by

$$(JP^{-1})(P\delta Z) = -F(Z). \quad (3.81)$$

The application of the right preconditioner requires only the action of $JP^{-1}$ on any Krylov vector $v$ and has to be performed at each Krylov iteration. This is realized in a two-step process:

1. First, apply the preconditioner and solve for $w$

$$JP^{-1}w = -F. \quad (3.82)$$

2. Next, the update is obtained by solving the linear system

$$P\delta Z = w. \quad (3.83)$$

The right-preconditioned version of Eq. (3.76) is used to solve Eq. (3.82) and is expressed as follows

$$JP^{-1}v \approx \frac{F(Z + \epsilon P^{-1}v) - F(Z)}{\epsilon} \quad (3.84)$$

where $v$ is any GMRes vector. Upon convergence of the linear solve in Eq. (3.82), one more preconditioner application is necessary using Eq. (3.83) to obtain the true Newton update for the non-linear iteration.

Up until now, the algorithm has been described in a general fashion, in the sense that the non-linear residual can be obtained after space-time discretization, the
approximate action of the Jacobian on a vector can be computed using Eq. (3.84) and finally an appropriate preconditioner $P$ can be chosen to reduce the conditioning number of the true Jacobian matrix.

Generally, preconditioners can be subdivided into two broad categories: Algebraic and physics-based preconditioners. The former deals with creating approximate sparse inverse factorizations, using numerical strategies to reduce the spectral radius of the linear system being solved. Some examples of such preconditioners include Incomplete Cholesky($\ell$) factorization, Incomplete-LU($\ell$) factorization along with reverse Cuthill-Mckee (RCM) reorderings, Sparse Approximate Inverses (SPAI) [62], Block-Jacobi splitting, Additive-Schwartz methods and Algebraic multigrid [63]. Algebraic preconditioners are often times also referred to as ‘numerical’ preconditioners. Algebraic methods are often easier to develop and use, and are particularly well suited for irregular problems that arise from discretizations involving unstructured meshes of complicated geometries. Furthermore these Algebraic methods can be fine tuned to make use of multi-processor architectures intricately in order gain improved scalability in the solution procedure.

Alternately, with intuitive understanding of the governing physics PDEs, the geometry, boundary conditions and details of the discretization for the problem under consideration, specialized preconditioners, usually based on physics-based OS linearizations, can be devised and used as very efficient preconditioners to damp the dominant modes, thereby leading to a well conditioned system. Multilevel methods usually fall in this category since they solve ‘nearby’ problems based on lower order discretizations. Few examples in this category of physics-based preconditioners include multigrid preconditioners [27], the method of Diffusion-Synthetic-Acceleration (DSA) [64] often used as a preconditioner for Transport equation and Implicit Continuous Eulerian (ICE) [31] for near-incompressible fluid flow problems. These problems are usually optimal for specific types of problems and might not be effective as generic preconditioners for all scenarios.
Note that in these physics-based preconditioners, the use of Algebraic preconditioners themselves is most often seen and hence such Algebraic preconditioners can be considered as building blocks for more advanced preconditioners. In the current work, both these approaches will be used in a mixed fashion, depending on the problem being solved in order to reduce the total number of linear iterations in Krylov solves. Also, care is needed while using preconditioners in a multi-processor architecture since traditional sequential preconditioners may sometimes fail in these scenarios. Hence, scalable preconditioners that can be used in both sequential and parallel linear Krylov solvers are preferred. A thorough survey of many state-of-art preconditioning methods used in computational physics problems was presented by Benzi [65].

Below, a brief description at some specific physics-based preconditioning techniques used in this research is provided and the reader is referred to previous work on these techniques for further details.

3.3.4 Physics-based Preconditioners

Legacy codes written to tackle the mono-physics models typically contain approximations for specific problems that usually result in increased efficiency even with a little loss of generality. A physics-based preconditioner is usually derived by the linearization of the non-linear physics components, in both the Elliptic and Hyperbolic equations based on semi-implicit treatment of the stiff terms. Such intricate knowledge of the physics systems for problems of interest can considerably improve the efficiency of the simulation. In the context of utilizing the MFK framework introduced earlier, these algorithms that currently exist in such codes can accelerate the linear solver convergence, thereby preserving man-years of testing and verification.
3.3.4.1 Linearized Jacobian for Elliptic Systems

Consider the SS terms in the non-linear Elliptic system shown in Eq. (3.11), linearized about the last non-linear iteration (*) as

\[ f(u^{n+1}) = q - (-\nabla \cdot D(\vec{r}, u^*)\nabla u^{n+1} + c(\vec{r}, u^*)u^{n+1}) \quad \forall \vec{r} \in \Omega. \] (3.85)

Let us define a new variable as \( \delta u = u^{n+1} - u^* \) which represents the true update for the non-linear iteration. Then if the physics-based preconditioner \( P \) approximates the Jacobian matrix for the non-linear Elliptic system, the preconditioner solve is

\[ P(\delta u) = -f \] (3.86)

Substituting this definition into Eq. (3.85), we obtain

\[ f(u^{n+1}) = q - (-\nabla \cdot D(\vec{r}, u^*)\nabla (\delta u + u^*) + c(\vec{r}, u^*)(\delta u + u^*)) \quad \forall \vec{r} \in \Omega. \] (3.87)

Expanding and simplifying Eq. (3.87), results in a modified residual equation of the form,

\[ f(u^{n+1}) = f(u^*) - (-\nabla \cdot D(\vec{r}, u^*)\nabla (\delta u) + c(\vec{r}, u^*)(\delta u)) = 0 \quad \forall \vec{r} \in \Omega. \] (3.88)

Hence in a Nonlinear-Krylov iteration framework, the coefficients \( D(r, u) \) and \( c(r, u) \) are evaluated about the linearized point to yield a linear elliptic equation system and the forcing function (source) for this linear equation for \( \delta u \) is \( f(u^*) \). Applying the Continuous-Galerkin FE discretization to Eq. (3.88) results in the standard stiffness and mass matrices along with appropriate boundary conditions applied to the solution. Hence, the preconditioner iteration is simply, in this case,

\[ (K^* + M^* + B^*)(\delta u) = f(u^*), \] (3.89)
and the preconditioner matrix \( P = (K^* + M^* + B^*) \). Once this matrix is formed, a Krylov method such as Conjugate Gradient (CG) or GMRes with appropriate Algebraic preconditioners can be used to effectively find the update for the solution \( \delta u \).

The linearized Jacobian matrix is an effective preconditioner when the linearization point (*) is closer to the true non-linear solution. The use of Incomplete-Cholesky and ILU factorization for symmetric and unsymmetric systems respectively can considerably reduce the total cost of the preconditioner solve itself. The current work utilizes such a linearized Jacobian matrix in conjunction with Algebraic preconditioners for non-linear scalar/vector elliptic/parabolic equation systems in order to reduce the total cost of FGMRes(\( r \)) Krylov solves.

3.3.4.2 Nearly Incompressible, Low-Mach Fluid Flow Systems

Fluid flows in reactor analysis problems fall under the low Mach (\( Ma \)) flow regime. In the conservative variable formulation, as the flow velocity of the fluid decreases, it is very difficult or almost impossible to solve low-speed flows with a conventional compressible algorithm because of slow convergence. The difficulty in solving the compressible equations for low Mach numbers is associated with the large disparity between the acoustic wave speed and the waves propagating at the fluid speed, which is called eigenvalue stiffness. To overcome this difficulty, several ideas have been proposed. In the current study, we will specifically use the Implicit Continuous Eulerian (ICE) scheme [31,32] for solving these low-speed problems. Some theoretical asymptotic analysis on the semi-discrete Euler equations using the ICE scheme using the implicit Backward-Euler method is shown in this section. The extension to higher order ERK/IRK methods is trivial.
Let us start with the non-linear inviscid Euler-like equations for unsteady fluid flow in the conservative form. For generality, a source term is also included. The equation system is given as:

\[
\frac{\partial U(x)}{\partial t} + \frac{\partial}{\partial x} F(U) = S(U),
\]

(3.90)

where

\[
U = \begin{bmatrix}
\rho : \text{Density} \\
\rho v : \text{Momentum} \\
E : \text{Total energy}
\end{bmatrix}; \quad F(U) = \begin{bmatrix}
\rho v \\
\rho v^2 + P \\
vE + vP
\end{bmatrix},
\]

and the source term \( S(U) \) is non-zero when solving manufactured problems (for verification studies), when the effects due to friction and gravity are included or when the fluid equations are coupled to energy transfer from a heated surface (conjugate heat-transfer). The pressure \( P \) is usually given by the closure relation, the Equation of State (EOS), in a linearized form as

\[
P = P_0 + \left. \frac{\partial P}{\partial \rho} \right|_0 \rho + \left. \frac{\partial P}{\partial E} \right|_0 E.
\]

(3.91)

The spatial discretization is performed using the Discontinuous Galerkin method with appropriate numerical flux functions (Upwind flux or Rusanov flux). Higher order spatial discretizations can be obtained by increasing the polynomial order of the Legendre basis functions.

For simplicity, let us redefine the momentum variable as \( M = \rho v \). Then the semi-discrete form of the equations, which are essentially the non-linear residual for the continuity, momentum and energy equations, can be written as:

\[
r_C(n + 1) : \quad \frac{\rho^{n+1} - \rho^n}{\Delta t} + \partial_x (M)^{n+1} = S_C (\rho, M, E)^n,
\]

(3.92)
\[ r_M(n + 1) : \\
\frac{M^{n+1} - M^n}{\Delta t} + \partial_x \left( \frac{M^2}{\rho} \right)^n + \partial_x P^{n+1} = S_M(\rho, M, E) |^n, \quad (3.93) \]

\[ r_E(n + 1) : \\
\frac{E^{n+1} - E^n}{\Delta t} + \partial_x \left( \frac{M^{n+1} E^n + P^n}{\rho^n} \right) = S_E(\rho, M, E) |^n, \quad (3.94) \]

where \( S_C, S_M \) and \( S_E \) are the continuity, momentum and energy source terms.

(Eq. (3.92))-(Eq. (3.94)) are the non-linear residual functions about the point \((n + 1)\). Now, choose a linearization point (*) that is typically chosen as the last non-linear iteration, about which a change in the state variable can be defined. This can then be given as

\[
\begin{align*}
\delta \rho &= \rho^{n+1} - \rho^* \quad (3.95) \\
\delta M &= M^{n+1} - M^* \quad (3.96) \\
\delta E &= E^{n+1} - E^* \quad (3.97) \\
\delta P &= P^{n+1} - P^*. \quad (3.98)
\end{align*}
\]

Substituting these new variables in (3.92)-(3.94), the following conservation equations for the delta form of the state variables are obtained.

\[
\begin{align*}
\frac{\delta \rho}{\Delta t} + \partial_x (\delta M) &= -r^*_C \\
\frac{\delta M}{\Delta t} + \partial_x \delta P &= -r^*_M \\
\frac{\delta E}{\Delta t} + \partial_x \left( \delta M \left( \frac{E + P}{\rho} \right)^* \right) &= -r^*_E,
\end{align*}
\]

(3.99)-(3.101)
where the linearized discrete residuals evaluated at the linearization point (*) are

\[ r_{C}^{*} = \frac{\rho^{*} - \rho^{n}}{\Delta t} + \partial_{x} (M)^{*} - S_{C}(\rho^{*}, M^{*}, E^{*}), \quad (3.102) \]

\[ r_{M}^{*} = \frac{M^{*} - M^{n}}{\Delta t} + \partial_{x} \left( \frac{M^{2}}{\rho} \right)^{*} + \partial_{x} P^{*} - S_{M}(\rho^{*}, M^{*}, E^{*}), \quad (3.103) \]

\[ r_{E}^{*} = \frac{E^{*} - E^{n}}{\Delta t} + \partial_{x} \left( \frac{M^{*} E^{*} + P^{*}}{\rho^{*}} \right) - S_{E}(\rho^{*}, M^{*}, E^{*}), \quad (3.104) \]

Note that the advection terms in the momentum Eq. (3.100) and energy Eq. (3.101) conservation equations and along with the source terms have been linearized about the point (*).

Rearranging the momentum equation Eq. (3.100), we obtain

\[ \delta M = -\Delta t \partial_{x} \delta P - \Delta t r_{M}^{*}. \quad (3.105) \]

This expression can then be substituted in the continuity Eq. (3.99) and the energy Eq. (3.101) equations to obtain a system of equations in \( \delta \rho, \delta E \).

\[ \delta \rho = -\Delta t \partial_{x} (\delta M) - \Delta t r_{C}^{*}, \quad (3.106) \]

\[ \delta E = -\Delta t \partial_{x} \left( \delta M \left( \frac{E + P}{\rho} \right)^{*} \right) - \Delta t r_{E}^{*}. \quad (3.107) \]

Now using the linearized Equation of State (EOS) introduced in Eq. (3.91), we can then substitute

\[ \delta \rho = \frac{1}{\frac{\partial P}{\partial \rho}(\rho)|_{0}} \left( \delta P - \frac{\partial P}{\partial E}(\rho)|_{0} \delta E \right), \quad (3.108) \]

Substituting the above equation in Eq. (3.106), we get

\[ \delta P = \frac{\partial P}{\partial E}(\rho)|_{0} \delta E - \frac{\partial P}{\partial \rho}(\rho)|_{0} \left( \Delta t \partial_{x} (\delta M) + \Delta t r_{C}^{*} \right). \quad (3.109) \]
Rearranging the above equation and substituting Eq. (3.107) and Eq. (3.105) for $\delta M$ and $\delta E$ respectively, we get the semi-discrete form of the pressure Poisson equation, given as

$$
\delta P = \Delta t \frac{\partial P}{\partial E} \bigg|_0 \left( \Delta t \partial_x \left( \partial_x \delta P + r^*_M \left( \frac{E+p}{\rho} \right)^* \right) - r^*_E \right) \\
+ \Delta t \frac{\partial P}{\partial \rho} \bigg|_0 \left( \Delta t \partial_x \left( \partial_x \delta P + r^*_M \right) - r^*_C \right),
$$

(3.110)

This system shown in Eq. (3.110) can be solved for $\delta P$ and back-substituted to obtain $\delta M$ from Eq. (3.105), $\delta E$ from Eq. (3.107), $\delta \rho$ from Eq. (3.106) respectively. It is important to note that the new system expressed as an elliptic pressure equation is exactly same as the original semi-discrete ICE linearized Euler equations. It is quite clear that by doing the algebraic manipulation shown in Eq. (3.108) for EOS and substitution of Eq. (3.105) into the continuity and energy equations, an equivalent Gaussian elimination on a system of size $4N (\delta \rho, \delta E, \delta M, \delta P)$ has been performed analytically to convert it to a block upper triangular form that is solved by back substitution. Hence solving the original ICE system Eq. (3.99) – Eq. (3.101) and the elliptic pressure equation Eq. (3.110) do yield the exact same result as long as the spatial discretization of the PDE’s are consistent in both cases. Since the pressure waves are resolved with an ICE solve, it results in eliminating all the dominant eigenmodes occurring due to the pressure waves, i.e., acoustic scales in the medium. Hence, the resulting system has a smaller spectral radius, especially for low-Mach flows where the spread between the eigenvalues in the original non-linear fluid flow equations is the quite large.

The gain in computational time when using ICE as a preconditioner and as a solver by itself has been shown previously in [66]. Now let us consider the advantages and disadvantages of the ICE preconditioner of size $N$ introduced earlier (denoted hereafter as $N$-ICE).
Pros

1. The elliptic pressure matrix in the fully discrete form is clearly only $N \times N$ while the original equation system was a $3N \times 3N$ hyperbolic system. The gain in terms of reduction in the size of the system, without any major approximations or loss of accuracy therein makes this a valuable method in low-Mach regime flow calculations.

2. Additional cost savings in terms of forming and solving the modified linear system in Eq. (3.110) using Algebraic preconditioners can significantly decrease total Krylov iterations for solving the Jacobian matrix.

Cons

1. It is difficult to maintain the consistency of the fully discrete ICE system w.r.t. the original dG discretization of the non-linear hyperbolic system, due to the requirements to evaluate the derivatives of momentum residuals in the right hand side in Eq. (3.110). Care is needed if a consistent preconditioner is to be created from the $N$-ICE system.

The $N$-ICE solver can typically be used as a solver by itself but the semi-implicit treatment leads to conditional stability only. However, when used as a preconditioner, the updates provided by such a linearization approximate the updates necessary for the outer Newton iteration. Hence in low-Mach regimes, these schemes are valuable to resolve the stiffness in the linear system quickly and, hence, act as efficient preconditioners to reduce the total linear iterations, thereby requiring fewer actions of the Jacobian on a Krylov vector.

3.4 Closing Remarks

In this section, we have covered the space-time discretization methods for different PDE systems and described the process along with the constraints to resolve
the different spatial and temporal scales in multi-physics computations. Based on these discretizations, the algorithm for a Matrix-free non-linear iteration method based on finite-difference approximations was introduced. The available options for using robust linear solvers along with different kinds of preconditioning techniques to increase overall efficiency of the algorithm were presented.

The ability to precondition Newton-type iteration methods with Picard linearized matrix falls under the category of multilevel preconditioning. This idea is at the core of the proposed MFNK framework wherein consistent actions of a Jacobian matrix on a vector are obtained through finite-difference approximations and inexpensive physics-based linearizations open up the possibility to make use of existing legacy code algorithms on top of powerful and scalable Algebraic preconditioners.

The current research implements all of these algorithms with help of some external software, to accurately couple multi-physics models in a computationally efficient unified code system.
4. A NON-LINEAR MULTI-PHYSICS COUPLED CODE SYSTEM

‘The function of good software is to make the complex appear to be simple.’
– Grady Booch

The methods for spatio-temporal discretization of different physics and the methods for tackling the non-linear system of coupled equations arising from the discretization were introduced in Section (3). Here, we describe the implementation of the MFNK framework, from a software perspective.

Software engineering of a coupled multi-physics code involves several considerations in the design and implementation of the interaction between different parts of a code. Even though the numerics laid out in Section (3) have a well defined structure regarding coupling multi-physics models tightly, without careful planning in the software design, even loosely coupled physics using the OS paradigm can be quite complicated to implement. Hence, utilizing the different numerics and physics models strongly depends on creating a software framework that is flexible, extendable and follows a plug-in architecture that can evolve as new or better methodologies for coupling multi-physics components are devised.

Some of the software requirements for a coupled multi-physics code framework include:

1. To re-use existing libraries to minimize development time, and to base the framework on already well verified discretization and non-linear/linear solver libraries. This thought stems from the basic Object-Oriented (OO) philosophy in avoiding code replication and modularizing implementation to accelerate development and testing phases.

2. To provide flexible data containers and physics objects to facilitate and simplify the evaluation of the non-linear residuals for different physics components.
3. To be able to use coarse grain physics models for rapid prototyping, testing and verification and the functionality to interchangeably use higher fidelity physics models to describe the physical phenomena in a straightforward fashion through common API contracts, as and when required by problem constraints.

4. To be able to use within the same architecture, different kinds of multi-physics coupling strategies with minimal changes from an end-user perspective. For instance, using OS with simultaneous or staggered updates, or using OS with Picard iterations to converge the coupling between physics, or employing MFNK tight coupling approaches side-by-side without changing how the non-linear residual describing the discretized PDEs is evaluated.

5. To have the flexibility to add different types of preconditioners, both Algebraic and physics-based, for each of the different physics component models and the option to choose how they are applied to reduce the total cost of linear iterations.

6. To use of recent advances in computer engineering for state-of-the-art multi-core, multi-processor parallel shared memory architectures that can significantly reduce run times for simulation of a physical phenomena.

7. To make the coupled physics code system independent from any specific spatio-temporal discretization. This involves the usage of different spatial discretization with any temporal discretization allowing the possibility to verify the implementation of the same equation system through more than one use-case.

The philosophy behind the software framework for multi-physics applications is to “solve tightly coupled phenomena using a loosely coupled software methodology”. The loosely coupled architecture is primarily made possible by requiring a software contract or a defined set of methods to be implemented. This is often called the API (Application Programming Interface) and needs to be defined clearly to allow future extensions.
In order to verify the numerical algorithms proposed in the current work, the need for a new code system was inevitable. Efforts to address this has led to the development of the KARMA framework (K(ode for Analysis of Reactor and other Multi-physics Applications).

4.1 KARMA

KARMA is a fully implicit, non-linearly coupled multi-physics eigenvalue and transient analysis test-bed code written completely in C++ programming language. Its primary intended application is to analyze and model coupled problems for nuclear reactor applications although it is not only limited to this family of problems. KARMA makes extensive use of the advanced OO concepts such as abstraction, encapsulation and inheritance, to create loosely coupled objects that allow seamless integration of new physics and numerics models.

The plug-in architecture employed in KARMA makes it straightforward to modify/add any number of coupled physics components. It also serves as a framework to conduct experiments on code architectures and software design for the next generation of consistent coupled multiphysics codes. The framework can be used to seamlessly integrate such physics models with consistent numerical algorithms that were introduced for non-linear PDE systems. In creating such a flexible framework, one of the prime concerns is the ability to achieve high levels of efficiency while still maintaining the ease of development, testing and maintenance. Careful planning of the computational domain has led to a decision to use well tested linear algebra data-structures and methods in order to reduce the overhead in re-implementing these standard algorithms, thereby eliminating the possibility of introducing errors in these basic building blocks for the numerical algorithms proposed in the current work. This also follows closely the OO principles and code re-use whenever possible, thereby preserving man-years of effort pertaining to code verification.
The requirements enumerated earlier are at the core of the design of the KARMA framework. These abilities in a multi-physics physics code framework are considered representative of current and future trends in solving coupled problems. Similar motivations have also led to the recent development of other coupled multi-physics codes like MOOSE [67].

KARMA is built on top of the state-of-the-art scientific library PETSc, the Portable, Extensible Toolkit for Scientific computation [68] from Argonne National Laboratory (ANL), for fast, scalable and robust data-structures and solvers. It provides tools for the parallel (and serial) numerical solution of PDEs that require solving large-scale, sparse non-linear systems of equations. It includes non-linear and linear equation solvers that employ a variety of Newton-type methods with line-search techniques and Krylov subspace methods. It also offers several parallel vector formats and sparse matrix formats, including compressed row, block compressed row, and block diagonal storage. The primary advantage of using PETSc is that well tested black-box methods and codes that can tackle non-linear systems arising from discretization of Parabolic/Hyperbolic system of equations are obtained implicitly by just linking with with the library. Also, usage of several different kinds of home-grown and external Algebraic preconditioners are obtained by interfacing KARMA with PETSc. Since PETSc is designed to facilitate extensibility, users can incorporate customized solvers and data structures when using the package. PETSc also provides an interface to several external software packages, including Matlab, PVODE, and SPAI and is fully usable from C and C++. Due to the advanced design, users can create complete application programs for the parallel solution of non-linear PDEs without writing much explicit message-passing code themselves. Parallel vectors and sparse matrices can be easily and efficiently assembled through the mechanisms provided by PETSc. Furthermore, PETSc enables a great deal of runtime control for the user without any additional coding cost. The runtime options include control over the choice of solvers, preconditioners and problem parameters as well as the generation
of performance logs. Since KARMA uses PETSc, all programs using the framework benefit from these ubiquitous options to control the program at a fine-grained level. For instance, options can be specified whether to run a program using a completely matrix-free approach with ‘-snes_mf’, where the action of the Jacobian is found through Eq. (3.76) and no preconditioner is used to solve the coupled system. This approach can lead to large number of Krylov iterations and hence adversely affect the CPU time. Alternately, if an approximation to the Jacobian is available, then a tightly coupled solution procedure can be used with the option ‘-snes_mf_operator’, where the action of Jacobian is again through Eq. (3.76) but the preconditioner matrix is created using the approximate Jacobian matrix representation which is usually some form of linearization about the last Newton iteration (physics-based preconditioner). Additional options using ‘-pc_type’ can be specified for the mode of solving the preconditioner itself which can either be an Algebraic variation (ILU, ICC, AMG) or using a much lower fidelity representation of the Jacobian matrix. Note that any level of recursion in the level of preconditioning, depending on the problem, can be implemented using such a MFNk technique based framework.

Efficient spatial discretizations using cG and dG FE methods along with FD and FV methods can be implemented for each of the physics PDEs. KARMA currently uses the general FEM library, libMesh [69]. It is written in C++ and provides support for first and second order Lagrange, arbitrary order \( C^0 \) hierarchic, \( C^1 \) continuous and discontinuous finite element bases. libMesh also facilitates writing dimension-independent code assembly of the non-linear residual and the Jacobian matrix for each of the physics component, which greatly simplifies the verification process for complex non-linear problems. libMesh has interfaces to the parallel vector, matrices, linear algebra data-structures provided by PETSc, and hence reduces the overhead to write distributed algorithms that are capable of utilizing the features inherently provided by PETSc.
KARMA also supports several different input and output formats that are convenient to generate the correct geometry, assign material region attributes and specify boundary markers. For convenience, in the current work, Gmsh [70] is used as the primary mesh generator. Gmsh uses the popular Delaunay mesh generators namely Triangle in 2-d and TetGen in 3-d. The parallel decomposition of the mesh can be performed using ParMETIS [71] to minimize the net communication time for a given geometry.

The chosen output format for writing out the solution fields is the VTK format [72] which is supported by several visualization packages.

 Optionally, KARMA can also be linked with a suite of eigensolvers exposed through the SLEPc library [73] that is based on PETSc. These state-of-art eigensolvers can handle symmetric and unsymmetric generalized eigenvalue problems that arise from the discretization of the elliptic and hyperbolic systems. For instance, the eigenvalue problem to find the fundamental mode in nuclear reactor design calculations is typically solved using the traditional Power Iteration method but we can also employ one of the eigensolvers provided by SLEPc e.g., use the more efficient Krylov subspace methods. The application of these solvers will be discussed in Section. (6).

Several other utility codes can also be optionally used to deal with XML and CSV input/output formats. For instance, TinyXML [74], a small C++ library that can handle reading, manipulation and writing of XML data with very little memory overhead can be used for specifying input options through a file which is forwarded to PETSc. And CSVParser can be employed as a parser to read and write Comma Separated Values (CSV) to aid in reading data from spreadsheets like Excel or data exported from MATLAB (csvwrite, csvread).

A schematic diagram showing these different parts of the KARMA framework and their interaction with the above mentioned packages is shown in Fig. 4.1.
Fig. 4.1. Schematic Diagram of KARMA Framework
4.2 Modules in KARMA Framework

The KARMA framework has four modules that are responsible for the implementation of the numerical schemes discussed in Section (3). Brief details regarding each of these modules is given below.

1. INTERFACE: This module is the heart of KARMA and is responsible for managing the different physics components and applying the numerical discretizations seamlessly to the coupled non-linear problem. Apart from maintaining a uniform interface to all the physics, it serves as the primary rendezvous point for all user interactions to obtain the coupled non-linear residual or the approximate Jacobian matrix. It also provides ‘C’ wrappers that serve as function pointers in order to interface with the PETSc and SLEPc solvers for the non-linear/linear and eigenvalue solvers respectively.

2. PHYSICS: This module contains all the physics descriptions, including the non-linear residual, the approximate Jacobian matrix and preconditioners, if any, that are spatially discretized forms of the corresponding physics PDE. All the different Physics objects derive from KARMAPhysicsBase that specifies the required methods that need to be implemented by all physics. Since this contract is known a-priori, a generic code to solve the physics can be written in the INTERFACE, making use of this polymorphic behavior for generating non-linear residuals and preconditioners.

3. NUMERICS: This module comprises of the necessary spatial and temporal discretization objects that are used by the PHYSICS module to provide the discretized form of the PDE. The use of libMesh in the current work eliminates additional overhead for spatial discretization. All necessary definitions of temporal integration methods in the form of a Butcher Tableaux are available along with generic integrators for ERK, DIRK and FIRK methods with adaptive time-stepping capability. This module also contains all the necessary
higher level wrappers necessary to use the non-linear, linear, eigenvalue solvers and preconditioners provided by PETSc and SLEPc libraries. These higher level objects make use of the wrappers provided by INTERFACE internally and hence the API provided by these objects remain the same, immaterial of whether, say, FGMRes or CG, is used as a solver.

4. IO: As the name of the module suggests, it contains all the necessary interfaces to the parsers and data writers to read/write the mesh format (msh), CSV, XML, and VTK for input data processing and output data manipulation in a generic fashion.

4.3 Solving a Non-linear Coupled Elliptic Problem

In this section, a step by step example of creating and solving a coupled elliptic problem involving two physics components Phy₁ and Phy₂ is provided below.

4.3.1 Adding a New Physics

A new physics model can be added in a straightforward manner by just deriving from the KARMAPhysicsBase class and implementing primarily three operators that are essential to solve any physics component. These operators are given below.

1. SystemOperator: This operator implements the steady-state non-linear residual definition, which is essentially the spatially discretized PDE representing the physics evaluated at a given time \( t \). This operator may also optionally provide an approximate Jacobian matrix, if it is easy to form. This matrix can be invoked and used during a solve with the `'-snes_mf_operator'` option. The implementation of this operator completes the description of the SS form of the problem.
2. **MassOperator**: This operator represents the mass matrix or a lumped version of it, resulting from the spatial discretization of the time derivative term for the Implicit Differential Equation Eq. (3.36). Note that this operator can either be time dependent itself (property dependent mass matrix or mesh changes with time) or be static, in which case it is only necessary to compute it once.

3. **PreconditionerOperator**: Any physics object can contain an array of preconditioner operators. These can be viewed as multi-level preconditioners where one could use $P_0$ to precondition the approximate Jacobian matrix $J$ and $P_1$ to precondition the solve for $P_0$, and so on. In practice, this sort of recursion might not be very efficient unless care is taken, for each physics, to resolve the stiff components first and systematically reduce the modes that are responsible for the high condition number of the true Jacobian matrix.

Once a physics system implements these three operators, the framework has all the necessary information to solve the system. All the material properties are provided through a problem context as function pointers and hence facilitate the use of arbitrary user-specified properties based on table lookups or correlations.

### 4.3.2 Writing a Non-linear Residual Function

The non-linear residual function that is part of the **SystemOperator** is at the heart of any physics description since it represents the discretized PDE for the physics model. Based on a sample implementation using **libMesh** library, a snippet C++ code is given for a single element residual assembly.

Let $e$ be the finite element under consideration that is a subset of the discrete mesh $\Gamma_h$. The local residual contribution can be computed based on the family of basis functions used to discretize the solution field, the points and location of the quadrature for element integration, and the degrees of freedom for the local solution
unknowns. For a diffusion-reaction physics system, the different components of the residual are obtained using Code Snippet 4.1.

```cpp
for (unsigned int iqp=0; iqp<qrule.n_points(); iqp++)
{
    qp = quadrature_point[iqp] ;

    qp_solution = 0. ;
    for (unsigned int i=0; i<phi.size(); i++)
        qp_solution += phi[i][iqp] * solution(dof[i]) ;

    // evaluate the properties at quadrature point with qp_solution
    diffusion_coefficient = properties.Diffusion(qp, time, qp_solution) ;
    reaction_coefficient = properties.Reaction(qp, time, qp_solution) ;
    source_term = properties.Source(qp, time, qp_solution) ;

    for (unsigned int i=0; i<phi.size(); i++)
    {
        Se(i) += JxW[iqp] * source_term * phi[i][iqp] ;
    }

    for (unsigned int j=0; j<phi.size(); j++)
    {
        Ae(i,j) += JxW[iqp] * ( 
            diffusion_coefficient * dphi[i][iqp] * dphi[j][iqp] +
            reaction_coefficient * phi[i][iqp] * phi[j][iqp] 
        ) ;
    }
}
```

Code Snippet 4.1 Element Residual Components

where $JxW$ is the Jacobian for the element transformation multiplied by the quadrature weight and the diffusion, reaction and source terms are computed at every
quadrature point, based on the non-linear solution at those points. The variational form of the diffusion-reaction problem yields the stiffness and mass matrices, which are stored in $A_e$, while the source term is assembled and stored in the load vector $S_e$.

With these computed contributions, the local SS residual is simply obtained by performing a local assembly followed by multiplication with the local solution dofs, as shown in Code Snippet 4.2.

```cpp
for (unsigned int i=0; i<phi.size(); i++)
{
    Re(i) = Se(i) ;
    for (unsigned int j=0; j<phi.size(); j++)
        Re(i) -= Ae(i,j)*solution(dof[i]) ;
}
```

Code Snippet 4.2 Residual Computation

It is obvious that all of the above steps for a single element are independent of the next element and hence provide a great deal of inherent parallelism. Also, since the basis functions and quadrature weights are calculated based on the problem’s dimension, the residual contribution code is dimension-independent.

If the above physics was coupled to a solution from another physics, it is then necessary to compute projection of the coupled physics solution on to the physical quadrature points used in the element assembly. This is greatly simplified if the coupled physics components use the same mesh since the projection operator is its own interpolant but, in the case of multi-mesh scenarios Section. (3.1.3), the $L_2$ projection of the solution with increased quadrature points will be necessary in order to reduce the spatial coupling errors.

It is important to note that none of the above code snippets mandate any specific discretization method to be used for the physics. A FD or FV method could have
been implemented as well instead of the FE method shown above, as long as the non-linear PDE is discretized accurately.

4.3.3 Obtaining Coupled Global Residuals

Once the individual physics components are computed, the driver code invokes the residual function in the KARMAInterface object which then calls the non-linear residual routines that are part of SystemOperator for each physics, as illustrated in Code Snippet 4.3.

```c
// distribute the provided solution vector to each physics
synchronize_physics_solution(X);

// loop over all the physics and compute the residuals
// and assemble it in the interface residual
for (; pos != end; ++pos)
{
    // Get a reference to the current physics system using the iterator
    KARMAPhysicsBase* physics = pos->second;

    KARMAVector& phy_residual = physics->get_residual();
    KARMAVector& phy_solution = physics->get_local_solution();

    // call the residual function for the physics
    physics->system_operator->residual(phy_solution, phy_residual);
}

// assemble the computed residual to the output vector
synchronize_interface_residual(R);
```

Code Snippet 4.3 Coupled Multi-physics Non-linear Residual Computation
The two synchronization routines namely `synchronize物理学_solution` and `synchronize_interface_residual` merely copy data back and forth between the physics object and the interface that maintains a global vector (for all physics). Hence, these residuals could be computed in a ‘black-box’ code and the KARMA framework could be utilized to compute high-order coupled temporal solution, provided that the coupled physics components are treated consistently (as modeled in the PDE).

4.3.4 Summary

In order to compute the non-linear solution for the coupled problem, only the non-linear residual vector is necessary. However, due to the inefficiency in this solution procedure, a linearized form of Jacobian matrix can be computed, with a similar implementation as 4.3.2 based on the nature of the physics.

Temporal discretization can be performed with any one of the ERK, DIRK or FIRK methods (using constant time-stepping or adaptive time-stepping controllers) available as part of KARMA package. With the SS non-linear residual $\mathbf{f}$, Jacobian matrix $\mathbf{J}$ and the MassOperator, Eq. (3.43) can be used to compute the time evolution of the solution.

A pseudo-code for the calling stack from the driver to solve the coupled multiphysics problem is shown below. This is general and can be extended easily to several physics components:

1. Create discrete meshes for the physics: $\Omega_1$ and $\Omega_2$
2. Create physics objects: $\text{Phy}_1$ and $\text{Phy}_2$
3. Add coupled physics reference to one another
4. Initialize KARMAInterface and all physics objects $\text{Phy}_i$, $\forall i = 1, 2$
5. Set initial temporal solution
6. Invoke the TemporalIntegrator to compute final step solution

- initialize and allocate intermediate stage vectors
- loop while time \( \leq \) final\_time
- solve the coupled non-linear problem at each stage
  - use finite difference approximation for action of Jacobian on a vector using Eq. (3.76)
  - if option -snes_mf\_operator is used, the approximate global Jacobian is built based on the Jacobian matrix of the individual physics by either ignoring the coupling terms in a Block-Jacobi fashion Eq. (3.72) or variations of this operator with additional coupled terms in order to resolve more stiff components. This problem-dependent preconditioner formulation can be quite effective to reduce the number of FGMRes iterations.
  - if additional preconditioners are available for each physics, apply them to the linear solves in the MFNK solution procedure
  - perform non-linear Newton or Picard iteration until convergence
- if adaptive, compute new time step
- if requested, write solution fields to VTK file
- end

7. Write final solution fields to VTK file

Since command-line options are provided by KARMA to choose the type of coupling strategy (loose (OS) or tight (MFNK)), comparisons on the performance of each coupling strategy can be analyzed without changing any of the user code implemented. This enables computation of sensitivity of a numerical method wrt the coupling strategy for multi-physics problems.
4.4 Closing Remarks

KARMA is designed with extensibility in mind and provides a flexible API to couple an arbitrary physics modules together. These modules are written either as part of the library itself or provided through an external code. Since different kinds of coupling methods can be tested under one code system, KARMA serves as a very valuable test-bed code to gain intuition on the optimal strategy for a particular multi-physics simulation. Adaptive techniques in both space and time have already been tested for few non-linear physics and these preliminary results show the feasibility of extending these ideas to multi-physics coupled problems.
5. RESULTS

‘What most experimenters take for granted before they begin their experiments is infinitely more interesting than any results to which their experiments lead.’
– Norbert Wiener

In previous sections, the numerical methods for coupled multi-physics problems and the code framework implemented based on these methods to solve the non-linear system of equations arising from the discretization of physics models have been given. First, the verification of the methods and the code is necessary, in order to understand the efficiency of these multi-physics coupling methods and to stress the need for tightly coupled solution methods. Then, the efficacy of these tightly coupled schemes will be analyzed for some problems that have multiple time scales.

The results section is organized as follows: First, each of the physics models are tested for spatial and temporal consistency (convergence order); Next, verification studies are performed for a conjugate heat-transfer model using semi-analytical techniques that will be discussed in subsequent sections. For problems with widely varying time scales, an efficacy (efficiency in terms of computational time versus accuracy of the solution fields) study is performed to analyze the computational gain due to tightly coupled methods. The coupled treatment of neutronics/heat-conduction physics is simulated next with problems that verify the implementation and quantify the uncertainty propagation due to errors in cross-section data. In the same setting, a stiff transient benchmark problem for the coupled neutronics/heat-conduction physics is used to present the advantages in terms of accuracy and stability of using MFNK tight coupling methods in contrast to traditional OS schemes.
5.1 Solution Verification

Typically, the accuracy and convergence of numerical models are established by using simplified problems for which analytical solutions are available. For time-dependent coupled multiphysics problems, analytical solutions are more difficult to obtain and we need to resort to (1) the Method of Manufactured Solutions (MMS) [75] (2) semi-analytical methods [76] and (3) formal convergence order studies to check behavior of discretization errors. In the current Dissertation, MMS problems are used extensively to analyze and prove code correctness. The basic philosophy behind MMS is as follows: an exact solution $U_{\text{ref}}$, with enough smoothness in space and time, is chosen a-priori and substituted into the continuous form of the PDE to obtain a suitable forcing function (i.e., right-hand-side of the PDE) that is then employed in the numerical simulation. Therefore, the numerically obtained values can be compared to the exact ones, providing a measure of the error. Since the discretization of the source terms are performed consistently in both space and time, the discrete solution can be driven towards the exact solution as the mesh and time step sizes are reduced. This procedure can be applied also for non-linear and coupled physics problems and is an useful tool for verification purposes. The second leg of the code verification is performed by using well established benchmarks involving the physics component that need be tested.

The global error in a numerical solution $U_{\text{num}}$ is usually measured in the $L_2$ norm,

$$
\|Error\|_2 = \|U_{\text{ref}} - U_{\text{num}}\|_2 = \sqrt{\frac{1}{|\Omega|} \int_{\Omega} (U_{\text{ref}}(r) - U_{\text{num}}(r))^2 \, dr}.
$$

A method is of order $p_s$ in space and order $p_t$ in time if the error varies as $O(\Delta r^{p_s})$ and $O(\Delta t^{p_t})$, respectively. The order of convergence in space is measured by computing the global error in a transient simulation for which the spatial mesh is successively refined. A small temporal grid is necessary to ensure that the temporal error is small enough so that the error observed is due to the spatial grid only. A similar procedure
holds for the temporal error calculation, where the spatial mesh is fine enough so that the spatial errors do not pollute global error and successive simulations are performed with uniformly refined time step sizes. The measurement of the error and the comparison of the space/time accuracy orders obtained with expected convergence rates prove that the code implementation is consistent with the mathematics and that the numerical solution converges to the true solution of the PDE.

The MMS forcing functions used in the simulations are given in the Appendix for different coupled physics scenarios. For further reading regarding the MMS, refer to [75, 77].

5.2 Verification of Individual Physics Models

In order to verify the implementation of the physics and the numerics models, a step-by-step process is adopted. For verifying a coupled multi-physics code, we first verify the single physics models, some of which are can be non-linear by themselves. To perform this step, analytical solution methods and/or the MMS techniques explained earlier are utilized.

5.2.1 Nonlinear Scalar Parabolic Problem

A general scalar non-linear Parabolic PDE is considered and spatial/temporal discretization of the equations are performed using cG with Lagrange basis functions and ERK, IRK methods respectively. The verification studies for problems involving such systems is shown here.
5.2.1.1 Verification

Using the MMS technique, a dimension-independent non-linear heat conduction problem is modeled. The thermal conductivity is chosen to be the non-linear functional \( k(T) = T^2 \). The exact solution is taken to be

\[
T(\vec{r}, t) = \tanh(t) \prod_{i=1}^{\text{dim}} \sin(\pi r_i),
\]

where \( \vec{r} = \{x, y, z\} \) and \( \text{dim} = 1, 2, 3 \). Because the exact solution is known, the spatial and temporal error discretization can be readily quantified using Eq. (5.1).

This test is presented with linear and quadratic Lagrange basis functions for spatial discretization, in a two-dimensional domain. As theoretically expected, second and third order spatial convergence rates are observed for linear and quadratic Lagrange basis functions respectively, see Fig. 5.1(a). Using a fine spatial mesh, the temporal order of accuracy for BE1(1), IM2(1) and the SDIRK3(2) schemes are obtained and plotted in Fig. 5.1(b). We can clearly note that the non-linear solution method based on the MFNK framework is high-order accurate in space and time and presents the expected theoretical orders of accuracy based on the space/time discretization.
5.2.2 Nonlinear Fluid Flow Problem

Next, the discretization of the problem arising from hyperbolic fluid flow equations are verified. We will consider a manufactured solution to verify accuracy and discretization errors. After these verification studies, the efficiency of the ICE preconditioner for the conservative equations is shown.

5.2.2.1 Verification

Using the MMS, profiles for the state variables are assumed and corresponding forcing functions for the continuity, momentum and energy equations are derived.
The exact solutions assumed for density, velocity and total energy [28] are shown below

\[
\rho = \rho_{\text{min}} + (\rho_{\text{max}} - \rho_{\text{min}}) \text{sech}\left(\frac{x - \omega t}{\delta}\right) \quad (5.3)
\]

\[
v = v_{\text{min}} + (v_{\text{max}} - v_{\text{min}}) \text{sech}\left(\frac{x - \omega t}{\delta}\right) \quad (5.4)
\]

\[
E = E_{\text{min}} + (E_{\text{max}} - E_{\text{min}}) \tanh\left(\frac{x - \omega t}{\delta}\right). \quad (5.5)
\]

The closure relation for the pressure equation was chosen to be the ideal gas equation represented by

\[
P = \rho e(\gamma - 1), \quad (5.6)
\]

where \(\gamma = \frac{c_p}{c_v}\). Using these exact solutions and the equation of state, the source terms for each conservation equation were obtained and the spatial and temporal convergence orders were computed.
The 1-d Navier-Stokes equations are solved in a fully implicit manner using the MFNK method. The spatial order of accuracy was measured for different polynomial orders of Legendre basis functions with dG FEM discretization. The obtained accuracy orders are as expected theoretically and prove that the spatial treatment of the 1-d fluid equations are consistent. With a fine spatial mesh and second order dG finite elements with Legendre basis functions, the temporal order of convergence for density solution for a final time $t = 2$ sec. was obtained and plotted on Fig. 5.2. The convergence plot for different methods is shown in Fig. 5.3.
(a) Temporal accuracy in $\rho$

(b) Temporal accuracy in $\rho U$

(c) Temporal accuracy in $\rho E$

Fig. 5.3. Fluid Flow Problem: Temporal Accuracy

5.3 Verification of Coupled Physics Models

Now that individual physics models have been verified, the next step involves verifying the numerical solution for coupled problems based on a combination of these single-physics models. The results obtained from these studies is shown in the following subsections.

5.3.1 Coupled Conjugate Heat Transfer Example

Coupled conjugate heat-transfer problems are common in all thermal-hydraulic calculations. These coupled phenomena are primarily boundary condition based and
hence lead to locally strong coupling effects. First, the MMS is applied to verify the implementation of such a coupled system of equations and then the efficacy of using different kinds of coupling schemes for these problems is presented.

5.3.1.1 Verification

To verify conjugate heat transfer between a conducting solid and a fluid, we employed a manufactured solution. A Matlab script was written to obtain the forcing functions based on the following assumptions. The script is given in Appendix A.1.

The flow is assumed to be uni-directional along the \( z \)-direction with a constant inlet mass flux. The Blasius correlation in the turbulent regime is used to compute the friction factor \( f_w \):

\[
 f_w = \frac{0.3164}{Re^{0.25}},
\]  

where \( Re \) is the Reynolds number of the fluid.

Thermal conduction of energy in the bulk fluid is assumed to be absent and energy is added to the fluid only at the wall surface (fluid-solid interface). The solid heat conduction model is basically a \( x-z \) slab in Cartesian co-ordinate system which convects the heat generated to the bulk fluid. The exact solutions for the fuel and fluid temperatures, \( T_{fuel} \) and \( T_c \), respectively, are given below.

\[
 T_{fuel}(x, z, t) = r_F \left( 1 + \tanh(C_{tf} t) \right) \left( \frac{1}{2} + \sin \left( \frac{\pi x}{2L_X} \right) \right) \\
 \left( 1 + \tanh \left( \frac{2w}{3} - \frac{w z}{L_Z} \right) \right) + T_{f0}
\]  

(5.8)

\[
 T_c(z, t) = r_T \left( 1 + \tanh(C_{tc} t) \right) \left( a + b \tanh \left( -cw + \frac{w z}{L_Z} \right) \right)
\]  

(5.9)

\[
 \rho(z, t) = \rho_c + f \left( 1 - \frac{T(z,t)}{T_{co}} \right) + g \sqrt{1 - \frac{T(z,t)}{T_{co}}},
\]  

(5.10)
The internal energy and total energies are given by

\[ \rho e = \rho C_v T_c \]  \hspace{1cm} (5.11)
\[ \rho E = \rho e + \frac{1}{2} \frac{G^2}{\rho}, \]  \hspace{1cm} (5.12)

where \( r, T, T_f, r_f, T_c, C_{tc}, w, a, b, c, f, g \) are parameters to control the magnitude and time scales of the solution. \( C_v \) is the specific heat at constant volume, \( G \) is the mass flow rate.

The Equation of State employed to close the system of equations is a linearized relation, dependent on density and temperature.

\[ P = P_0 + \alpha (\rho - \rho_0) + \beta (T_c - T_0), \]  \hspace{1cm} (5.13)

where \( \alpha, \beta \) are the linearization constants. Note that \( \alpha \) is actually related to the speed of sound in the flowing medium and provides a simple way to manually change the Mach number in the calculations, apart from varying the mass flux \( G \) itself.
Based on this manufactured solution, the forcing functions have been generated and a convergence order study has been carried out for various levels of spatial and temporal discretizations. The numerical solutions approach the true solutions as the spatial and temporal meshes are refined, as expected. The convergence results, shown in Fig. 5.4 and Fig. 5.5, prove that the implementation of the physics is verified and demonstrate that higher-order accuracy can be obtained using the MFNK technique.
The conjugate heat transfer problem introduced earlier with MMS is used again to test the efficacy of the coupling methods. By varying the temporal scales of the exact solutions i.e., change the constants as $C_{tf} = 1000C_{tc}$. This forces the evolution of the fuel temperature to occur at a much faster rate than the transient in the fluid equations. In order to resolve this stiffness, we use the traditional OS coupling strategy and the MFNK method with the $L$–stable SDIRK2(2), SDIRK3(3) methods.
The accuracy results obtained for a given spatial mesh is plotted against the total computational time for the different coupling techniques in Fig. 5.6. From the efficacy plot for the total energy field Fig. 5.6(b), it is evident that for any given user-specified tolerance, the total cost for the solution procedure using OS strategy with first order BE method is higher by several orders of magnitude. But since the temporal evolution of fuel temperature occurs at a much faster rate, resolving the physical time scales required reducing the time step sizes for all the methods. Once the asymptotic region was reached, the higher order MFNK methods quickly reduce the error in the solution. For the temperature variable, for the same computational cost, the loosely coupled scheme is less accurate than MFNK based tightly coupled solution by two orders of magnitude. This result indicates that for stiff problems, once the dynamical physical scales are resolved, the higher order temporal accuracy in the tightly coupled schemes do provide considerable computational gain. It is also interesting to note that using a second order temporal method (CN) with OS coupling saves computational due to the fact that there is only 1 implicit stage while the SDIRK2(2) method has two stages per time step. These results indicate that it is possible for OS schemes with high-order temporal methods to be feasible in terms of efficiency, as compared to tightly coupled methods, when the time step sizes are well below dynamical time scales of the individual physics.
5.3.2 Coupled Neutronics-Thermal Conduction Example

Neutronics and Heat conduction physics solution fields are strongly coupled to each other through temperature dependent cross-section values and heat generation from fission in the fuel. Since the coupling between these two physics models is strong,
it is necessary to test the effectiveness of the coupling methodology to accurately resolve the varying time and length scales. In this section, the coupled neutronics and heat conduction physics models are verified by means of MMS. In the first case, the same spatial discretization are applied to both physics; in the second case, different (non-embedded) spatial meshes are used to quantify the effect of non-conforming meshes.

5.3.2.1 Verification: Identical Spatial Meshes

Making use of the MMS technique once again, a test problem is used to verify convergence of the method to exact solutions. Since the coupling between neutronics and conduction is non-linear and due to the fact that the conduction physics is non-linear by itself, obtaining manufactured solutions can become quite intricate. A Matlab script has been written to obtain the forcing functions based on the following assumptions.

Two delayed neutron precursors groups are considered along with two energy groups and one scalar temperature field. Hence, the total number of solution fields is five. We give below the exact solution for the fields for the 2D test case:

\[
\begin{align*}
\phi_1(x, y, t) &= A_\phi(1 + \tanh(r_\phi t)) \sin(\pi x) \sin(\pi y) xy \\
\phi_2(x, y, t) &= \phi_1(x, y, t) \times \frac{\Sigma_{s,1-2}}{(\Sigma_{rem,2} + D_2 B_2^g)} \\
C_i(x, y, t) &= C_i(x, y, 0)e^{-\lambda_i t} + \int_0^t ds \ e^{\lambda_i(s-t)} \sum_{g=1}^{g=2} \beta_{i,g} \nu \Sigma_{f,g} \phi_g(x, y, s) \\
T(x, y, t) &= A_T(1 + \tanh(r_T t)) \sin(\pi x) \sin(\pi y),
\end{align*}
\]

where \( B_g^2 = \left(\frac{\pi}{L_X}\right)^2 + \left(\frac{\pi}{L_Y}\right)^2 \) is the geometric buckling term, \( L_X \) and \( L_Y \) are the domain sizes in the \( x \) and \( y \) dimensions. \( A_\phi, r_\phi, A_T \) and \( r_T \) are constant parameters. Using
the exact solutions for the fluxes, the exact solutions for the precursors, $C_i$, can easily be obtained and are given by

$$C_i(x, y, 0) = \frac{1}{\lambda_i} \sum_{g=1}^{g=2} \beta_{i,g} \nu \Sigma_{f,g} \phi_g(x, y, 0), \quad i = 1, 2. \quad (5.18)$$

Doppler feedback is accounted in the neutronics model through the removal cross-section of group 1:

$$\Sigma_{rem,1}(T) = \Sigma_{rem,1}(T_0) + \left( \frac{\partial \Sigma_{rem}}{\partial \sqrt{T}} \right)_0 (\sqrt{T} - \sqrt{T_0}). \quad (5.19)$$

The following equation is employed to described the temperature-dependent conductivity:

$$k(T) = k_0 + \left( \frac{\partial k}{\partial T} \right)_0 (T - T_0). \quad (5.20)$$

The Matlab script used to obtained the forcing functions is given in Appendix A.2. Note that here only the fast energy group’s removal cross-section is affected by temperature variations. Additional temperature dependencies can be included in the removal and fission cross-sections for both groups using the provided script. With the knowledge of the exact solution profiles and the corresponding forcing functions, a space/time convergence study is carried out. The convergence plots for this example are shown in Fig. 5.7 and Fig. 5.8.
It is clear from the results that the solution fields for all physics components are high-order accurate in space and time and agree well with theoretical convergence rates. By varying the coupling coefficient $\frac{\partial \Sigma_{\text{rem}}}{\partial \sqrt{\frac{T}{T}}}$ in Eq. (5.19), and other free parameters such as $r_\phi$, $r_T$, $1/v_1$, $1/v_2$, stiffer transients were created and convergence to the true solution was observed still. The higher order temporal schemes are efficient and the stiffly-accurate SDIRK schemes prove to be superior than traditional low-order BE scheme.
Fig. 5.8. Coupled Neutronics/Heat Conduction Problem: Temporal Accuracy

5.3.2.2 Verification: Non-embedded Spatial Meshes

Multi-physics applications often require that different physics components employ different spatial meshes, which resolve the spatial scales occurring in the specific physics. Even though a complete study of this question is not covered in this Dissertation and requires extensive work all by itself, an example is provided here for the coupled neutronics/heat conduction physics utilizing different spatial meshes; see Fig. 5.9. Note that the meshes are not embedded (when meshes are embedded, projection and interpolation operators are simpler to define).
As detailed in Section 3.1.3, the spatial coupling error that may occur due to inadequate projection/interpolation in between source/target meshes can be avoided by employing a ‘high-enough’ numerical quadrature in the finite element setting. It is important to verify that mitigation of the interpolation errors is possible by using
high order quadratures. In this example, the same test problem with MMS from Section 5.3.2.1 is considered but different spatial meshes are employed.

The exact solution profiles for the two physics components show that their spatial scales vary differently. Without the use of high-order quadratures, the interpolation error starts to dominate and corrupt the numerical solution from reaching asymptotic spatial convergence orders. In order to eliminate this, high-order quadrature rules are employed to overkill the spatial errors due to solution interpolation and projection on a target mesh, as indicated in Section 3.1.3. Fig. 5.10 shows that high-order spatial convergence is recovered, as expected, for the MMS by using more quadrature points to resolve the variation in the coupled solution. This is only a preliminary study with multi-mesh for multi-physics software verification and further analysis is needed to test the efficiency of the technique employed in the Dissertation.

![Graphs showing spatial accuracy for Fast Flux and Fuel Temperature](image)

Fig. 5.10. Coupled Neutronics/Heat Conduction Problem with Non-conforming Meshes: Spatial Accuracy

5.3.2.3 Efficacy

A rod ejection benchmark problem from the ANL problem book [78] (Identification: 14-A1) is considered here to verify the coupling method described in this
Dissertation. This is a super-prompt critical transient with adiabatic heating and Doppler feedback for a thermal reactor model. In contrast with the previous MMS test cases, the time scales change quite drastically over the transient and it is expected that higher order time stepping schemes would outperform low-order schemes. In this simulation, the total power level changes by over 10 orders of magnitude. The adiabatic heating assumption is also numerically challenging: since no dissipative terms are present, the errors introduced in computing the temperature field can grow undamped with inconsistent numerical treatment, thereby requiring accurate coupling and time stepping methods.

Based on the described geometry, the domain was discretized using second-order triangular Lagrange finite element, as shown in Fig. 5.11. The initial steady state eigenvalue obtained is \( k_{\text{eff}} = 0.996364153180855 \) for the given spatial mesh and it closely matches the reference value. The transient calculation was performed using various combinations of coupling and temporal discretization schemes: OS-BE1(1), OS-IM2(1), MFNK-BE1(1), MFNK-IM2(1), MFNK-SDIRK3(2). The reference solution was calculated with a fine spatial mesh and a 3rd order \( L \)-stable SDIRK scheme SDIRK3(3) with \( \Delta t = 0.001 \) secs. The results obtained from these the reference simulation for the fast and thermal flux distributions are shown in Fig. 5.12 and the transient evolution in Fig. 5.13. It is interesting to note that the profile peak power solution does not match the benchmark solution but considering that low order discretizations with coarse spatial meshes were used in the benchmark, it is possible that the true converged solution is the one given here. Ryosuke et al. [79] noticed similar temporal evolution of the solution fields and the code to code verification of KARMA with the PRONGHORN software gives confidence in the results obtained here.
All the tightly coupled simulations (MFNK-based solves) use $\Delta t = 0.004$ secs. The OS simulations had to employ a much smaller time step of $\Delta t = 10^{-4}$ secs in order to converge; indeed, the explicit linearization of the OS schemes led the solution to diverge with large time steps. Fig. 5.14 shows a closer view of the power peak due to the rod ejection transient.
It is clear that the first-order BE1(1) scheme, even with tight coupling, i.e., MFNK-BE1(1), wrongly computes the magnitude and time of occurrence in the power peak. This trend with BE1(1) can also be seen with the OS coupled strategy, even with the much smaller time step used for OS schemes. The higher order schemes, IM2(1) and SDIRK3(2), approach the reference solution consistently as time step sizes are reduced and predict the occurrence of the peak power accurately.
These results emphasize the usefulness of employing higher-order temporal integration schemes to predict the behavior of stiff transients that can occur in nuclear
reactors. The usage of a tight coupling strategy allows to use larger time steps than OS schemes, due to the fully resolved non-linearities at each time step, which can be used in combination with adaptive time-stepping techniques to reduce the total computational times.

5.4 Uncertainty Quantification for Multi-physics Problems

Uncertainty quantification is necessary even for single-physics simulations and it is imperative to perform this analyses for multi-physics problems in order to gain better understanding on weak versus strong coupling between the physics. Aleatory and Epistemic uncertainty contributions can occur from a variety of sources. They can stem from physical model errors, data uncertainty errors, numerical errors due to inaccurate geometry descriptions, resolution of spatial and temporal scales, numerical errors from coupling methodologies to name a few, apart from Epistemic errors that are inherent to the system being modeled. In the current work, uncertainty propagation in the solution specifically due to material properties which are usually obtained through experiments and/or higher fidelity models with necessary approximations and homogenizations, is analyzed. Since the properties used in numerical simulations depend heavily on closure relations and empirical correlations, the propagation of error and uncertainty in the solution fields need to be ascertained. It is also important to note that the uncertainties introduced in the solution obtained from simulations are irreducible, even with a fine resolution of spatial or temporal scales when there is an uncertainty in the data.

Uncertainty quantification in the solution due to these variables following certain distributions can then be performed using several mature methods. One popular and less intrusive option is the Generalized Polynomial Chaos (GPC) method [80]. GPC is in essence a decomposition method where an uncertain output variable of
interest $Z$ is expressed in terms of a basis $\Phi$ of a stochastic space consisting of random variables $\zeta = \{\zeta_1 \ldots, \zeta_n\}$,

$$ Z(t, \zeta) = \sum_n Z_n(t) \Phi_n(\zeta) $$

(5.21)

where $\zeta$ can be described by either a Gaussian, Gamma or Uniform distribution. For most problems, where the true distribution is not known, a Gaussian distribution can be chosen to represent the random variable. In this case, Hermite polynomials are used as basis functions $\Phi_i$ and are given by

$$ H_n(x) = \Phi_n(x) = (-1)^n e^{x^2} \frac{d^n}{dx^n}(e^{-x^2}) $$

(5.22)

where $n$ is the order of the Hermite polynomial and $x$ is the random variable of interest. GPC methods have exponential convergence as the order of the polynomials $p$ is increased. Further information can be obtained from references [81, 82].

The application of a GPC method of order $p$, GPC($p$), to the coupled neutronics-conduction model problem is studied here. The introduction of the dependency on the randomness of the properties adds another dimension to the system of equations. Hence, the coupled physics solution fields are also functions of the stochastic variation in these random input variables. In other words, Eq. (5.21) can be rewritten in terms of random variables $\zeta$ as

$$ Z(t, \zeta) = C_0 + \sum_j C_j(t) H_1(\zeta_j) + \sum_j C_j(t) \prod_k H_2(\zeta_k) + \ldots $$

(5.23)

where $C_j$ are the coefficients of the expansion and can be computed making use of the orthogonality property of Hermite polynomials. In Eq. (5.23), the coefficients $C_j$ are essentially the moments of the $j^{th}$ order Hermite polynomial. Note that the zero’th moment $C_0$ represents the mean value of the physics solution $Z$ when all the random variables $\zeta$ are at their mean values. In order to obtain the coefficients,
Eq. (5.23) can be multiplied by a Hermite polynomial of certain order and integrated from $-\infty$ to $\infty$, and using orthogonality relation, we have

$$C_i = \int_{-\infty}^{\infty} d\zeta e^{-\sum_j \zeta_j^2} Z(t, \zeta) H_n(\zeta)$$

(5.24)

Note that in Eq. (5.24), the solution $Z(t, \zeta)$ is obtained from code results using a random sample for the input variable based on its distribution. The actual integral in Eq. (5.24) is computed using numerical quadrature rules, leading to non-intrusive uncertainty quantification methodology. Gauss-Hermite quadrature rules are used for the numerical integration and the order of the quadrature chosen should be high enough to compute the integral exactly. For instance, to compute the GPC(2) coefficients, a Gauss-Hermite quadrature rule that can exactly integrate a fourth order polynomial (three point quadrature rule, $qp=3$) is required. Hence the total number of simulations necessary, after optimizations to eliminate recurrent cross-term effects, is in the order of $n \ast p \ast qp$ where $n$ is the number of input random variables. Once the required simulations are computed, the coefficients $C_i$ can be computed and the output variables of interest can be found using the basis Eq. (5.23).

Alternatively, Monte-Carlo based sampling procedure are efficient at providing accurate statistical information regarding the uncertainty in the models. But since Monte-Carlo based sampling methods require large number of samples (or model runs), it is prohibitive to directly perform the simulation for every sample. We can instead use the basis obtained through GPC($p$) polynomial and then use this as a surrogate model to obtain the required statistical data for each output variable of interest, along with their respective Probability Distribution Functions (PDF). Using such surrogate modeling methodology designed specifically for propagating uncertainty from model inputs to model outputs, a quantitative analysis can be performed and better confidence in simulation predictions can be gained.

A super prompt-critical transient problem is considered here, along the lines of the previous ANL benchmark. This 2-d, two energy-group, two delayed-group neutron
diffusion problem with Doppler feedback has a variation in the thermal absorption cross-section defined by the relation:

\[ \Sigma_{rem,2}(T) = \Sigma_{rem,2}(T_0) + \left. \frac{\partial \Sigma_{rem}}{\partial \sqrt{T}} \right|_0 (\sqrt{T} - \sqrt{T_0}) + \left. \frac{\partial \Sigma_{rem}}{\partial T} \right|_0 (T - T_0). \] (5.25)

The geometry of the problem essentially consists of a rodded fuel element surrounded by unrodded fuel which is encompassed in a lattice of reflector assemblies. This is shown in Fig. 5.15. The material properties are obtained from MATPRO correlations for UO\(_2\) fuel with 95% theoretical density [83]. The transient is initiated by changing the thermal fission cross-section for the rodded fuel element through a ramp duration of 0.1 seconds. The solution profiles for the fast and thermal fluxes at SS conditions is shown in Fig. 5.16.

Fig. 5.15. Uncertainty Quantification Test Problem: Geometry and Computational Mesh
Four random input variables are chosen with the following Gaussian distributions:

\[
\begin{align*}
\Sigma_{\text{rem},2}(T_0)_{\text{rod+fuel}} & = N(0.2231, 5\%) \\
\Sigma_{\text{rem},2}(T_0)_{\text{fuel}} & = N(0.09194, 5\%) \\
\frac{\partial \Sigma_{\text{rem}}}{\partial \sqrt{T}} & = N(0.002043, 10\%) \\
\frac{\partial \Sigma_{\text{rem}}}{\partial T} & = N(5 \times 10^{-6}, 10\%)
\end{align*}
\] (5.26) (5.27) (5.28) (5.29)

and the uncertainty in three specific solution fields namely, \( k_{\text{eff}} \), the total maximum power in the transient, and the peak fuel temperature, are sought after.

A second order GPC expansion \( (p = 2) \) is considered along with the possible cross-terms to account for propagation of errors due to complex interplay of different physics. In order to efficiently compute the analytical integrals arising from the GPC basis expansions to compute the coefficients, a high-order Gauss-Hermite quadrature rule is used to eliminate any integration errors.
Once a reference spatial mesh is determined, the mean solution fields with all input random variables at their probable mean values are found. Then, each random variable is perturbed by a factor of $10^{-6}$ sequentially and the corresponding change in the solution fields is found. This measure of the global sensitivity of the solution fields w.r.t these random variables is given in Table 5.1. Since the current test problem is

Fig. 5.16. Uncertainty Quantification Test Problem: Initial Solution Profiles
at hot zero power conditions (initial fuel temperature = 300 C), the eigenvalue $k_{eff}$ is not dependent on the coupling with temperature in the fuel. But all the transient parameters namely, the peak power and maximum fuel temperature, are strongly sensitive to the doppler coefficient $\frac{\Sigma_{rem}}{\sqrt{T}}$. The values themselves in Table 5.1 are specific for the problem considered here and can change depending on the mean, the deviation and the chosen input distribution.

Table 5.1

| Output variable      | $\Sigma_{rem,2}(T_0)|_{rod+fuel}$ | $\Sigma_{rem,2}(T_0)|_{fuel}$ | $\frac{\partial \Sigma_{rem}}{\partial \sqrt{T}}$ | $\frac{\partial \Sigma_{rem}}{\partial T}$ |
|----------------------|---------------------------------|---------------------------------|---------------------------------|---------------------------------|
| $k_{eff}$            | -0.3173834                      | -8.4336155                      | 0                               | 0                               |
| Peak Total Power     | -12.156938                      | 28.4867                         | -551.53479                      | -11.8566                        |
| Peak Temperature     | -6.74126                        | 15.06638                        | -1.046647E6                     | -2.61168E4                      |

For the transient simulated here, the stability limit for traditional OS coupling was found to be around $\Delta t = 0.02$ by experiment i.e., the OS strategy needed time step sizes smaller than $\Delta t = 0.02$ to obtain a stable numerical solution. We use both the tightly coupled and OS based loosely coupled methods for the same transient with this stability limit step size. Based on the simulation results, the output variable expansion coefficients were computed and a surrogate model based on the expansion in Hermite polynomial basis was created. This surrogate model was then used with Monte-Carlo sampling techniques with $5 \times 10^6$ samples to obtain the mean and deviation in the output variables given in Table 5.2 and the corresponding Probability/Cumulative Distribution Functions shown in Fig. 5.17 for both OS and tight coupling based solutions.
Fig. 5.17. Uncertainty Quantification Test Problem: Distribution Functions for Output Variables

(a) Probability Distribution Function

(b) Cumulative Distribution Function
It is evident from the peak total power distribution plots that the loosely coupled schemes have a different distribution than the true solution distribution (tightly coupled). As the time step size is decreased, this disparity does vanish since the OS linearization for coupled physics terms becomes more accurate. It is possible to use Richardson extrapolation technique to compute the time step independent coefficients $C_j$ to eliminate the corruption of temporal errors in estimating the uncertainty due to input random variables, which provides the new possibility to obtain better prediction of the output uncertainty with OS coupling schemes.

Table 5.2
Mean and standard deviations for output variables

<table>
<thead>
<tr>
<th>Coupling</th>
<th>$k_{eff}$</th>
<th>Peak Total Power</th>
<th>Peak Temperature</th>
</tr>
</thead>
<tbody>
<tr>
<td>OS</td>
<td>N(0.9999867970, 0.019539)</td>
<td>N(177685.35,23178.69)</td>
<td>N(3968.08,295.6)</td>
</tr>
<tr>
<td>MFNK</td>
<td>N(0.9999867970, 0.019539)</td>
<td>N(167900.83,20936.29)</td>
<td>N(3939.17,290.3)</td>
</tr>
</tbody>
</table>

Since traditional multi-physics codes use loose coupled schemes, the predicted mean and the standard deviation for the range of input distribution are conservative (mean power and temperature are higher). In reactor design and safety calculations, the margins calculated using these can then restrict the design and reduce the total thermal power output. Even though the problem posed here is not a full reactor core configuration, the results obtained suggest that improvements based on tightly coupled methods can lead to improved margin characterization. Further studies are needed using higher order GPC expansions and different time step sizes in order to reduce the total uncertainty in the output variables and to gain better intuition on uncertainty quantification of multi-physics problems.
‘If you wish to advance into the infinite, explore the finite in all directions.’
– Johann Wolfgang von Goethe

In this section, the application of the KARMA framework to a different variety of non-linear multi-physics problems, other than those that occur in nuclear reactors transients, is explored. The flexibility of the framework and the versatility of the applications of the implemented code are demonstrated using these problems. First, the application of the Newton’s method for eigenproblems is shown followed by a stiff system of coupled equations occurring in radiation-diffusion problems. The theory and results obtained from these problems are shown here.

6.1 Criticality Eigenproblem and Modal Analysis

Large algebraic eigenvalue problems often arise in nuclear engineering applications. In reactor physics, criticality problems typically require the calculation of the fundamental eigenvalue and associated eigenmode to determine cycle lengths, reactivity margins and power distributions. For analyzing the unstable patterns encountered in Boiling Water Reactors (BWR) [84,85], often modal analysis is performed that requires solutions to large eigenvalue problems. Reactor instabilities are local power variations occurring due to periodic flow oscillations and can sometimes grow undamped. In stability studies, not only the fundamental mode but also higher-order eigenmodes are needed in order to predict core behavior accurately.

Fast and accurate numerical methods to obtain the fundamental mode and to perform modal analysis for reactor instabilities are an ongoing topic of research; see, e.g., [86,87,88,89,90]. In the current section, we underscore the link between all eigenproblems with constrained non-linear optimization problems and take advantage of this reformulation to use the Newton’s method for solving the eigenproblems.
This lends the possibility to use the existing MFNK framework introduced earlier in Section. (3).

Now let us consider the eigenproblems encountered in reactor physics based on the Multi-group Neutron Diffusion (MGND) equation given below:

\[-\vec{\nabla} \cdot D^g(\vec{r}) \vec{\nabla} \Phi^g(\vec{r}) + \Sigma_r^g(\vec{r}) \Phi^g(\vec{r}) - \sum_{g' \neq g} \Sigma_{s}^{g' \rightarrow g}(\vec{r}) \Phi^{g'}(\vec{r})\]
\[= \frac{1}{\lambda^g} \sum_{g'=1}^{G} \nu \Sigma_{f}^{g'}(\vec{r}) \Phi^{g'}(\vec{r}) \quad \forall g = 1, \ldots, G, \text{ for } \vec{r} \in \mathcal{D}. \quad (6.1)\]

In addition to Eq. (6.1), which is defined in the reactor domain $\Omega$, boundary conditions are supplied on the domain’s boundary $\partial \Omega$. The boundary conditions are typically of homogeneous Dirichlet-type (zero flux) or of homogeneous Neumann-type (symmetry lines).

After spatial discretization using cG FEM, the above multigroup equations can be recast in a discrete operator form

\[L \phi_m = \frac{1}{\lambda_m} F \phi_m, \quad (6.2)\]

where $L$ is the multigroup loss operator containing leakage, absorption and scattering terms, $F$ is the multigroup production operator containing the fission terms, $\lambda_m$ is the $m^{th}$ eigenvalue and $\phi_m$ is the $m^{th}$ (multigroup) eigenmode associated with the
The pair \((\lambda_m, \phi_m)\) will be denoted hereafter as the \(m^{th}\) eigenpair. The block structure of the \(L\) discrete operator is

\[
L = \begin{bmatrix}
L_1 & 0 & \ldots & 0 \\
-L_{2,1} & L_2 & 0 & \ldots & 0 \\
\vdots & \vdots & \ddots & \ddots & \ddots \\
-L_{t,1} & -L_{t,2} & \ldots & L_t & -L_{t,t+1} & \ldots & -L_{t,G} \\
\vdots & \ddots & \ddots & \ddots & \ddots & \ddots & \ddots \\
-L_{G1} & \ldots & -L_{G,G-1} & L_G
\end{bmatrix}
\] (6.3)

where the row index \(t\) denotes the first thermal energy group, i.e., the first group for which up-scattering is present. The diagonal blocks \(L_i\) represent the discrete form of \(-\nabla \cdot D \nabla + \Sigma^i\) whereas the off-diagonal blocks \(L_{i,j}\) represent down-/up-scattering operator \(\Sigma^{j-i}_s\). Similarly, blocks \(F_{i,j}\) are the discrete representation of \(\chi^i \nu \Sigma^j_f\). Each block in \(L\) and \(F\) is a real and sparse matrix of size \(n \times n\) with \(n\) denoting the number of spatial unknowns. It is also important to note that matrix \(L\) is not symmetric due to the non-symmetric scattering processes in between energy groups.

The problem dimension \(N = n \times G\) is large (for moderately coarse 3D 2-group diffusion calculations, \(N\) is of the order of, at least, 50,000 unknowns.) The \(m\)-th eigenfunction is a multigroup vector: \(\phi_m = [\phi^1_m, \ldots, \phi^G_m]^T\), where each component \(\phi^g_m\) is a vector of length \(n\).

The following ordering of eigenvalues is chosen: \(|\lambda_1| > |\lambda_2| \geq |\lambda_3| \geq \ldots \geq |\lambda_{N-1}| \geq |\lambda_N|\), where we have indicated the uniqueness of the largest eigenvalue in neutronics applications [91]. Out of the \(N\) eigenmodes, the first (fundamental) mode predicts the critical core configuration (in neutronics, it is customarily to let \(\lambda_1 = k_{eff}\), where \(k_{eff}\) is the effective multiplication factor) and higher modes dictate the behavior of the system during a transient. Since higher modes decay away in comparatively shorter time periods as the mode index number increases, only a subset of the higher-order modes are needed in practical applications.
Eq. (6.2) can be symbolically transformed by left multiplication with $L^{-1}$ to obtain a standard unsymmetric eigenvalue problem of the form

$$M \phi_n = \lambda_n \phi_n,$$

with $M = L^{-1}F$.

The generalized eigenvalue problem of Eq. (6.2) is only reformulated as Eq. (6.4) for notational simplicity and $M$ is never directly computed and stored due to the large memory requirements (i.e., $L^{-1}$ is never computed explicitly.) To simplify notations, the standard eigenvalue problem, Eq. (6.4), will be used hereafter, although it is important to note that each matrix-vector (matvec) product of the form $z = Mx$ requires (i) one matvec operation $y = Fx$ followed by (ii) a linear solve $Lz = y$ for $z$. Traditionally, the solve of the loss operator $L$ is performed by recursively sweeping through the energy groups in a Block-Gauss-Seidel (BGS) fashion until convergence. Hence, we will consider one such BGS sweep as the primary work unit and performance parameter for the different algorithms tested in computing several eigenmodes.

In most existing reactor analysis codes, the Power Iteration (PI) [92] method has been used to obtain the largest eigenvalue. By using deflation techniques or spectral shifting techniques [92] with the PI method, higher eigenmodes can be obtained, albeit with decreasing accuracy. This is due to the fact that performing deflation very accurately is computationally expensive and any inaccuracy introduced in this process, i.e., in the previous modes, can result in the corruption of the deflated system for higher-modes, thereby making it difficult to recover the true eigenmodes.

Recently, Krylov-subspace iteration methods have been proposed as alternatives to the PI method to compute several dominant eigenmodes, e.g., the subspace projection method with locking of converged eigenvectors [93], the explicit Arnoldi iteration [73, 90], the implicit restarted Arnoldi method (IRAM) [94, 95, 96, 97, 98], the Jacobi-Davidson method [99, 88, 100]. Subspace methods for eigenproblems have
been described and reviewed by several authors in the mathematics and numerical analysis communities; see, for instance, [101, 102].

The approach employed in the current work aims to employ the existing framework based on MFNK technique in order to find multiple eigenmodes accurately. This methodology is based on hybridizing the traditional techniques for solving eigenvalue problems with a variant of inexact Newton’s method. In order to underscore the link between eigenproblems with optimization problems tackled using Newton’s method, consider the following Rayleigh quotient functional (a non-linear functional) for a given Hermitian matrix $A$, defined as:

$$ R(\phi) = \frac{\phi^T A \phi}{\phi^T \phi} \quad \text{for} \quad \phi \neq 0. \quad (6.5) $$

The minimization of the Rayleigh quotient requires satisfying the optimality condition

$$ \nabla R(\phi) = 0 = \frac{2}{\phi^T \phi} (A \phi - R(\phi) \phi). \quad (6.6) $$

This optimality condition is solved using Newton’s method

$$ [\nabla^2 R(\phi^l)] \delta \phi = - \nabla R(\phi^l) \quad (6.7) $$

$$ \phi^{l+1} = \phi^l + \delta \phi, \quad (6.8) $$

where $\nabla^2 R$ is the Hessian matrix of the Rayleigh quotient functional, or equivalently the Jacobian matrix of the optimality condition, Eq. (6.6). These simple lines of algebra may facilitate establishing the relationship between eigenproblems and Newton’s method, which will be discussed in section 6.1.2, where we propose a hybrid scheme that combines Newton’s method and subspace iteration eigensolvers in order to compute, with high accuracy, a large number of eigenmodes for nuclear reactor analysis problems. Furthermore, the non-linear solves of Newton’s method will be performed in a matrix-free fashion to avoid computing the possible expensive matrix
Finally, recall that matrix $M$ is never formed, and only the action of the linear system on a vector is realized. Hence for improved efficiency, preconditioned versions of the matrix-free Newton’s method are applied.

6.1.1 Review on Existing Schemes to Compute Multiple Eigenmodes

Various subspace iteration eigensolvers applying a Rayleigh-Ritz projection have become popular over the recent years to determine several dominant eigenmodes in nuclear reactor applications; several references of such work were given in the introduction. In their simplest forms, they generate Krylov subspaces by repeatedly multiplying matrix $M$ with a basis vector, and in that sense can be thought of generalizations of the power iteration method. To reduce the size of the Krylov space needed to compute several eigenvalues near a portion of the eigenspectrum, some appropriate shifting strategies can be employed. Similar shifting strategies can also be used in the simple PI method to obtain eigenvalues with a faster convergence. A brief review of the PI and Krylov-subspace family of methods are given below.

6.1.1.1 Standard Power Iteration

In most reactor design and analysis codes currently in use, some form of modified/accelerated Power Iteration (PI) technique is employed to find the fundamental eigenvalue ($\lambda_1$) and the associated eigenmode. Such a procedure is known to be slowly convergent when the dominance ratio ($\frac{\lambda_2}{\lambda_1}$) is close to 1, resulting in a large number of operator applications, requiring many inversions of the multigroup loss operator $L$.

Common schemes to accelerate the power iteration technique are the Chebyshev acceleration [103] and the Wiedlandt shift [92].
6.1.1.2 The Shifted Inverse-Power Iteration

The Shifted Inverse-Power Iteration (SIPI) [101] is a popular and improved variant of the PI method where a guess (shift) of the eigenvalue is known reasonably well.

In the SIPI method, the eigenspectrum is shifted by a constant value $\sigma$ while preserving the corresponding eigenvectors. Hence, when a reasonable guess for the eigenvalue is known a-priori, the SIPI iteration converges to the eigenpair efficiently. Alternately, the Rayleigh quotient can be used as an improved shift at each iteration, leading to a locally quadratic asymptotic convergence rate for unsymmetric problems [104]. It has also been shown by Geltner that a good initial guess of the eigenvector is necessary for local convergence of an eigenmode, particularly when the loss matrix is not positive definite. The Rayleigh shift resembles the Rayleigh quotient shown in Eq. (6.5) and is given by

$$
\sigma_i = \frac{\phi_i^T M \phi_i}{\phi_i^T \phi_i},
$$

and at each SIPI iteration the following linear system needs to be solved

$$(M - \sigma_i I) \phi_{i+1} = \phi_i.$$

It should be noted that, as the iteration converges to the solution eigenpair, the shifted matrix operator $(M - \sigma_i I)$ becomes ill-conditioned and singular. This is one of the primary disadvantages in using the SIPI method.

The linear system $(M - \sigma_i I)$ is usually solved iteratively (inner iterations) to a given linear tolerance. It is possible to set this linear tolerance adaptively based on the norm of the residual of the eigenvalue problem, leading to an inexact-SIPI method, akin to the inexact-Newton method in philosophy, which is computationally less expensive than the exact SIPI variant. The motivation behind this is that the linear solves need not to be extremely accurate, while the eigenvalue residual is still large.
The SIPI procedure is suitable to find one eigenpair. When more than one eigenpair is sought, e.g., when performing modal analysis, additional work needs to be performed to remove the contribution of converged eigenmodes. In the work by Demaziere [90], the traditional PI method is used to calculate several eigenmodes with Wielandt deflation but this procedure involves the necessity for calculating the forward and adjoint eigenvector for each mode. The accuracy of the subsequent modes may be affected if the computed modes have a large residual error that can propagate in the computation of subsequent higher modes and corrupts the deflated linear system (in [90], a tolerance close to machine precision was used).

Alternately, a suitable locking technique through orthogonalization can be performed in order to remove the contribution of an already computed mode from the linear system [105]. In the our implementation of Rayleigh quotient iteration, this procedure is utilized for computing several dominant eigenmodes.

6.1.1.3 Krylov Subspace Iteration Methods

The basic philosophy of the PI method is to build a subspace by the repeated action of operator $M$ on a starting vector $v$, yielding a $\ell$-dimensional subspace whose span is $V = \{v, Mv, M^2v, ... M^{\ell-1}v\}$. As the size $\ell$ of the subspace becomes large, the power series spanned by the subspace does not form an appropriate basis to extract the eigenvalue and eigenvector approximations [102]. The columns of this subspace matrix $V$ are not orthogonal. In Krylov subspace iteration, an orthogonal basis is sought and the resulting Krylov matrix $\hat{V}$ provides approximations to the eigenvectors corresponding to $\ell$ dominant eigenvalues of $M$. Some advanced Krylov subspace methods for non-symmetric system of equations include the explicit Arnoldi [73] and Implicitly Restarted Arnoldi [95], the Jacobi-Davidson [99] and Krylov-Schur [73] methods.

The original Arnoldi algorithm [101] is a powerful extension of the subspace iteration in that it builds an orthonormal basis of the Krylov subspace and factorizes
the matrix $M$ into an upper Hessenberg matrix. The central idea behind the Arnoldi factorization is to construct eigenpairs of the original matrix from eigenpairs of the factorized, much smaller, Hessenberg matrix. Sorensen [106] improved the Arnoldi method by introducing several shifted QR iterations on the Hessenberg matrix; this reduces the overall number of required matvec operations. Several other variations and optimizations have been added to this Arnoldi method and have been successfully implemented in the library ARPACK [95]. The Implicitly Restarted Arnoldi Method (IRAM) [106] is considered to be one of the state-of-the-art schemes to compute several dominant eigenpairs for large, sparse linear systems. Several researchers [97, 107, 98, 90] have considered the use of an Arnoldi eigensolver for $k$-eigenvalue problems and found it to be a promising alternative to obtain the fundamental mode.

6.1.2 Newton Iteration Based Hybrid Algorithm

Instead of approaching the eigenvalue problem with traditional iterative techniques, we recast the eigenproblem as a non-linear problem to be solved by means of Newton’s method. Peters and Wilkinson [108] proved that the inverse iterations are equivalent to a Newton’s iterative scheme. Saad [101] also considered the suitability of using non-linear Newton type iteration schemes for large symmetric eigenvalue problems and proposed several variations for the definition of the non-linear residual function.

Let us consider the eigenvalue problem in Eq. (6.4) and formulate it as a $(N+1)$-dimensional unconstrained optimization problem (recall that (i) $M = L^{-1}F$ is a matrix of size $N$ by $N$ and that (ii) it is never explicitly formed).

Based on a formulation proposed by Wu et al. [102] for the standard eigenvalue problem, the non-linear residual function can be written as follows

$$
\mathcal{F}(y) = \begin{bmatrix}
(M - \lambda I) \phi \\
-\frac{1}{2} \phi^T \phi + \frac{1}{2}
\end{bmatrix} = 0
$$

(6.11)
where $\mathcal{F}(y)$ is the non-linear Newton residual vector, of size $N + 1$. The first $N$ components of $\mathcal{F}$ represent the linear system for the eigenproblem in Eq. (6.4) and the last component of $\mathcal{F}$ is simply a 2-norm normalization of the eigenvector. The $N + 1$ dimensional solution vector $y$ contains the eigenmode and the eigenvalue and is given by

$$y = \begin{bmatrix} \phi \\ \lambda \end{bmatrix}$$  \hspace{1cm} (6.12)

The eigenmode and the eigenvalue solution are then obtained using Newton’s method

$$\delta y^l = -[J(y^l)]^{-1}\mathcal{F}(y^l)$$  \hspace{1cm} (6.13)

and

$$y^{l+1} = y^l + \delta y^l$$  \hspace{1cm} (6.14)

where the Jacobian matrix is given by

$$J(y) = \begin{bmatrix} M - \lambda I & -\phi \\ -\phi^T & 0 \end{bmatrix}$$  \hspace{1cm} (6.15)

Peters [108] proved that this augmented matrix is non-singular even when $(\lambda, \phi)$ is the true eigenpair being sought. Saad [101] confirmed the proof for the symmetric case and extended it to the non-symmetric matrix case which is considered here.

The choice of the optimization problem given in Eq. (6.11) avoids nearly singular, ill-conditioned Jacobian matrices and hence is convergent locally to an eigenpair without the numerical difficulties (e.g., large condition numbers) often observed in techniques such as the SIPI method near an eigenpair solution [109]. The quadratic convergence of the Newton scheme can be utilized to create a robust eigensolver as long as a suitable initial guess for the eigenmode is available. It is also known that the above Newton-type procedure for eigenvalue problems is equivalent to the Rayleigh Quotient iteration, whose convergence behavior is well understood [102,
109]. Nonetheless, a few points have to be addressed in order to use Newton’s method to compute the eigenmodes:

1. the need for an appropriate initial guess close to the desired eigenpair;

2. the cost of computing the Jacobian matrix $J$ and memory requirements for storing it are prohibitive for large problems;

3. the need of efficient preconditioning technique to reduce Krylov subspace required for inner linear solves.

The next paragraphs address these. First, Newton’s initial guesses to compute the dominant eigenmodes will be obtained from either a few SIPI iterations or a Krylov subspace iteration technique, with coarse tolerances. The Newton iteration acts as the eigensolver once it has been “bootstrapped” using a standard eigensolver used to provide an initial guess to focus Newton’s search eigenspace around one or several desired modes. Second, we will rely on matrix-free approaches to avoid the computation of the Jacobian matrix. Finally, preconditioning techniques will be employed. Details regarding the proposed algorithm are discussed below.

6.1.2.1 Starting the Newton Iteration

It is well known that Newton’s method converges quadratically to the solution as long as the initial condition is inside the “ball of convergence”. If we are to obtain a certain number of dominant modes, then proper initial guesses must be provided. For our purpose, we are not interested in just any eigenmode and a procedure to enrich the Newton’s search directions for the dominant eigenmodes is necessary in order to focus the Newton’s search space.

One option is to perform first a few SIPI iterations (with coarse tolerance) to obtain an approximation to the fundamental eigenmode, thus providing a starting vector for Newton’s method. We refer to this scheme as the “SIPI-Newton” hybrid
scheme. Note that if the shift parameter is set to 0, the standard PI is then used to focus the search space.

A second option is to employ a subspace iteration method (we have chosen the IRAM scheme) to bootstrap the eigenpair search with Newton’s method. This option is more appealing than the “SIPI-Newton” scheme because the Arnoldi iteration readily provides approximations to several dominant eigenpairs at once. They can be obtained with a coarse tolerance *without* having recourse to explicit deflations or the need for an initial eigenvalue estimate as in the SIPI method. Additionally, the subsequent Newton iterations to solve for the different modes are completely independent and this stage could be performed in parallel. Hereafter, we denote this scheme, where coarse-tolerance estimates from the IRAM technique are employed as initial iterates for the Newton solves, as the “Arnoldi-Newton” hybrid scheme.

Finally, we note that a fine tolerance is not necessary at the start of Newton solve in Eq. (6.13) when the search direction is far away from true solution. Then, the Jacobian solve can be performed ‘inexactly’ in the sense that the linear iteration tolerance can be made to depend on the non-linear function residual Eq. (6.13). This Inexact-Newton iteration method results in the following convergence criteria for the linear solve.

\[ \| J(y') \delta y' + F(y') \| \leq c^l \| F(y') \|, \]

where \( c^l \) is a parameter chosen to tighten the linear solve convergence as the non-linear residual is reduced. For more information on Inexact-Newton schemes and the forcing factor \( c^l \), we refer the reader to [59].

6.1.2.2 Matrix-free Technique

In Jacobian-free variants of Newton’s method, the explicit computation of the Jacobian matrix \( J \) is not required, which is particularly useful in our case since the
Jacobian matrix contains matrix $M = L^{-1}F$, which we do not want to compute explicitly nor store. Since the Jacobian-free method is at the core of the tightly coupled solution algorithm introduced for solving multi-physics problems, much of the infrastructure for implementing the hybrid eigenvalue solver is in place.

Let the $(N + 1)$-dimensional Krylov vector $v$ be written as $(\varphi, \mu)^T$ where $\varphi$ is the portion of the Krylov vector corresponding to the eigenfunction and $\mu$ is a scalar corresponding to the eigenvalue. Then, the action of the Jacobian matrix on a Krylov vector $v$ can be obtained as

$$ Jv \simeq \frac{F(y + \epsilon v) - F(y)}{\epsilon} = \begin{bmatrix} M\varphi - \lambda\varphi - \mu(\phi + \epsilon \varphi) \\ -\phi^T\varphi - \frac{1}{2}\epsilon\varphi^T\varphi \end{bmatrix} $$  \hspace{1cm} (6.17)

where it should be remembered that $z = M\varphi$ is actually the following linear solve: $Lz = F\varphi$. Eq. (6.17) is based on a finite difference approximation, which can be avoided here: since the definition of Jacobian is exactly available, albeit not explicitly, a matrix-free solve using GMRes, with no memory allocated for the Jacobian matrix itself, can be carried out and the (exact) matrix-vector operator is given by

$$ Jv = \begin{bmatrix} M\varphi - \lambda\varphi - \mu\phi \\ -\phi^T\varphi \end{bmatrix} $$  \hspace{1cm} (6.18)

In Eq. (6.18), the effect of the perturbation $\epsilon$ has vanished. Both Eq. (3.76) and Eq. (6.18) are Jacobian-free approaches (the Jacobian matrix is not formed) and both forms require the same linear solve $Lz = F\varphi$. In our implementation, we have chosen the exact matrix-vector operation, i.e., Eq. (6.18), over the finite difference approximation of Eq. (3.76).
6.1.2.3 Preconditioners for the JFNK Technique

In order to limit the size of the Krylov subspace in the linear solve at each Newton iteration, effective preconditioning techniques are necessary. The form of outer (Newton) – inner (Krylov) iteration for eigenvalue problems resembles the inverse iteration subspace family of methods and also is closely related the Jacobi-Davidson scheme [108, 109] when the secondary equation being used as a preconditioner is Point-Jacobi. The power of the MFNK scheme is that any number of preconditioners can be employed as long as (i) the cost of computing the preconditioner itself is cheaper than the cost of computing the true operator \( M \) and (ii) the preconditioner collapses the eigenspectrum so that the spread in eigenvalues is reduced. Note that every preconditioner solve can also be performed in a matrix-free fashion rather than actually forming the preconditioning matrix \( P \), if memory requirements necessitate that.

Some details on the preconditioning methods for the simple eigenproblem in Equation (6.4) are now discussed.

6.1.2.4 Block Gauss-Seidel Preconditioner

The Block Gauss-Seidel (BGS) form of the preconditioner, where a block is defined as one energy group, is a natural choice of preconditioner to invert \( L \). In traditional reactor analysis codes using PI method, the BGS method is typically used to invert the loss operator solve. In the current JFNK context, we use a single BSG sweep over all groups as a preconditioner. Since each diagonal block \( L_i \) is symmetric (when discretized using a standard finite element or finite difference method), efficient Krylov linear solvers such as Conjugate Gradient can be used to solve each one-group equation. Here, the individual blocks are themselves preconditioned with
a level-0 Incomplete Cholesky decomposition, IC(0). Therefore, the preconditioner $P$ is given by

$$
P = \begin{bmatrix}
\tilde{L}^{-1} F - \lambda I & -\phi \\
-\phi^T & 0
\end{bmatrix}
$$

(6.19)

where $\tilde{L}$ is the lower triangular block of $L$, hence discarding any upscattering terms. A preconditioner solve requires the following solves:

$$
L_i z_i = (F \varphi)_i - \sum_{j<i} L_{i,j} z_j \quad \text{for } i = 1, \ldots, G
$$

(6.20)

Other preconditioners such as multigrid, multilevel methods could be utilized instead of the BGS scheme. Another idea would be to use traditional PI technique as a preconditioner for the Newton solve, as suggested by Knoll [110].

6.1.3 Implementation

The methods presented here have been implemented in the KARMA code framework which interfaces to external libraries such as PETSc [68], SLEPc [73] and ARPACK [95] to make use of existing eigenvalue solvers for comparison purposes and to bootstrap the Newton iteration by providing appropriate initial guess.

The PETSc-based Scalable Library for Eigenvalue Problem Computations (SLEPc) library has several standard eigenvalue solvers such as PI, SIPI, Explicitly restarted Arnoldi method and Krylov-Schur methods. Apart from the built-in solvers, SLEPc also provides interfaces to the ARPACK eigensolver package. The IRAM algorithm employed as bootstrapping in our hybrid method was taken from ARPACK.

For the requested number dominant eigenmodes, ($nev$), the computational cost of IRAM behaves as $N \ell^2$ where $N$ is the problem rank of matrix $M$ and $\ell$ is the size of the subspace. Hence for fine tolerances and large problem sizes ($N$), a bigger span of Krylov space may be needed, increasing the total memory cost. Instead, the proposed hybrid Arnoldi-Newton scheme can be implemented in a completely
matrix-free fashion using JFNK, with IRAM to deliver the initial eigenmodes to coarse tolerances (thus requiring only a smaller subspace) and Newton’s method to drive the residual of the eigenproblem to a tight precision. This scheme can compute several dominant eigenpairs (in parallel) with low memory overhead and high accuracy with existing frameworks of PI implementation.

The hybrid SIPI-Newton scheme can also be used to converge to the dominant eigenpairs but usually it is expected to be inferior in terms of performance in comparison to advanced Krylov iteration techniques like Arnoldi to find multiple eigenvalues. This is due to the fact that the choice of the shift parameter affects the convergence of this scheme significantly and eigenpairs can only be found one at a time, i.e., sequentially. Nevertheless, the low memory requirements and the relative ease in implementation of this scheme as compared to the Arnoldi-Krylov method may make it attractive, especially since traditional reactor analysis codes already have most of the necessary framework in place. In order to make the results shown here independent of a user-provided shift, 20 iterations of power iteration are performed to obtain an eigenvalue estimate to 1e-2 tolerance. This then is used as the initial shift for all SIPI runs thereby making this hybrid scheme automated to some extent.

6.1.4 Results

Numerical results using the different eigensolvers introduced in the previous sections are presented here for three typical eigenvalue problems found in nuclear reactor analysis. The first case is a 2-D IAEA benchmark problem that is used to analyze the convergence of the implemented scheme for the first few eigenmodes. The next problem considered is a homogenous 2-group, 2-d problem to compare the traditional PI, state-of-art IRAM and the current hybrid algorithms in terms of efficiency in computing the fundamental mode, as a function of the dominance ratio.
6.1.4.1 Case 1: IAEA 2-D Benchmark Problem

For the purpose of determining the accuracy and efficiency of the proposed numerical scheme, the well known IAEA 2-D benchmark problem (ANL, 1977) with modifications to the cross-section to account for z-leakage is used as a test problem. This problem is a two group model of a PWR quarter core with reflective boundary conditions on left and top sides and vacuum boundary on the remaining sides (zero incoming current). Details on the used cross-sections for different materials and the lattice representation is given in Appendix (B.1).

The 2-D domain is discretized using triangular mesh elements. The spatial discretization of the generalized eigenvalue problem is performed using piecewise quadratic, Lagrange elements on triangles. The reference results presented below for the eigenvalue computation were from a discretization with 81940 elements and 123937 unknowns/group.

The thermal flux profiles for the first five modes of the benchmark problem obtained using the hybrid Arnoldi-Newton iteration are shown in Fig. 6.1 - Fig. 6.5.

![Thermal Flux Profile](image.png)

**Fig. 6.1.** IAEA 2D Benchmark Problem: First Eigenmode for Thermal Flux.
Fig. 6.2. IAEA 2D Benchmark Problem: Second Eigenmode for Thermal Flux.

Fig. 6.3. IAEA 2D Benchmark Problem: Third Eigenmode for Thermal Flux.
Fig. 6.4. IAEA 2D Benchmark Problem: Fourth Eigenmode for Thermal Flux.

Fig. 6.5. IAEA 2D Benchmark Problem: Fifth Eigenmode for Thermal Flux.
In Table (6.1), the eigenvalues for the first 10 modes are shown for the benchmark problem along with the L2 norm of the residual error $M \phi_i - \lambda_i \phi_i$. The eigenvalues from the IRAM-Newton hybrid scheme were compared to the solution from a fine tolerance (1E-14) IRAM run that is implemented in ARPACK, interfaced through SLEPc. The eigenvalues computed from both schemes match exactly and hence this test case proves the convergence of the hybrid scheme to all the desired modes. In the above run, an IRAM iteration provides a reasonable guess of the eigenmodes to a coarse tolerance of only 1E-3 with maximum subspace size of 15 to bootstrap the inexact Newton iteration.

<table>
<thead>
<tr>
<th>Mode</th>
<th>IRAM-Newton</th>
<th>Residual Error</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1.02958492118978</td>
<td>3.925623517e-14</td>
</tr>
<tr>
<td>2</td>
<td>1.00262113845152</td>
<td>1.665425359e-14</td>
</tr>
<tr>
<td>3</td>
<td>0.99162639339383</td>
<td>4.511323676e-14</td>
</tr>
<tr>
<td>4</td>
<td>0.93909498902443</td>
<td>3.756085782e-14</td>
</tr>
<tr>
<td>5</td>
<td>0.91382020547476</td>
<td>4.287420904e-13</td>
</tr>
<tr>
<td>6</td>
<td>0.90139826550461</td>
<td>1.052507002e-13</td>
</tr>
<tr>
<td>7</td>
<td>0.89045692528088</td>
<td>3.088887872e-14</td>
</tr>
<tr>
<td>8</td>
<td>0.82718775002535</td>
<td>1.762787579e-13</td>
</tr>
<tr>
<td>9</td>
<td>0.82499156741152</td>
<td>3.749614193e-14</td>
</tr>
<tr>
<td>10</td>
<td>0.81562246518003</td>
<td>4.042513635e-14</td>
</tr>
</tbody>
</table>

The results shown in Table (6.1) indicate that the new numerical scheme computes the eigenpairs with machine precision accuracy for all the requested modes (nev=10) of the model problem. Hence the new convergence of the Newton based hybrid scheme to the true numerical eigenmode is guaranteed as long as the initial guess provided is in the ‘ball of convergence’.
Next, the same problem is solved using different eigen-methods and the cost of computation in terms of number of BGS operator applications is listed in Table (6.2). Also the size of the subspace indicates the memory cost requirements needed for solving the problem. The results indicate that the SIPI based schemes are unfeasible in terms of the total cost due to the chosen method for determining the initial shift. If a better algorithm is available for this, the higher eigenmodes can be computed with lesser cost. On the other hand, IRAM and IRAM-Newton hybrid scheme show very efficient performance and the hybrid scheme only requires half the memory requirement as compared to pure IRAM run. The results show the strength in terms of convergence with lower memory cost of the hybrid scheme based on Arnoldi iteration.

<table>
<thead>
<tr>
<th>Method</th>
<th>Number of operators</th>
<th>Subspace size</th>
</tr>
</thead>
<tbody>
<tr>
<td>SIPI</td>
<td>8115</td>
<td>1</td>
</tr>
<tr>
<td>SIPI-Newton</td>
<td>6507</td>
<td>1</td>
</tr>
<tr>
<td>IRAM</td>
<td>2180</td>
<td>20</td>
</tr>
<tr>
<td>IRAM-Newton</td>
<td>2116</td>
<td>10</td>
</tr>
</tbody>
</table>

The preliminary results show that there is an optimal subspace-size for IRAM which minimizes the total number of BGS operators. But a-priori, this size is not known and can only be determined by experiments on the problem of interest. Also, as the subspace-size increases, the memory requirements increase linearly and can become prohibitive unless distributed systems are utilized. The IRAM-Newton scheme alternately uses IRAM only to start the iteration and hence could gain immensely from the usage of a much lesser subspace size to achieve coarse tolerances. And if auxillary systems representing the Jacobian matrix can be used as preconditioners, the total cost of the linear Krylov solve can also be reduced considerably.
6.1.4.2 Case 2: Homogenous Infinite Medium, 2-Group Problem with Up-scattering

Consider a 2-D, two energy group problem with non-zero, fast-to-thermal and thermal-to-fast scattering cross-section. Hence, the Loss matrix $L$ is a full block matrix, unlike the IAEA 2D case which was in block-lower triangular form. The cross-section data used for the problem are given in Appendix (B.2). The implementation was verified by checking the spatial convergence of the eigenvalue to the exact eigenvalue, as the mesh is refined.

In the infinite medium limit, the dominance ratio for the discretized, generalized MGND eigenvalue problem can be made to approach unity. This translates to a large number of iterations that is needed in order to converge to the fundamental eigenpair. If several dominant eigenpairs are desired in this context, traditional power iteration techniques are practically infeasible.

Better Preconditioners such as multigrid, multilevel methods for such elliptic systems will significantly improve the performance of the linear solves and hence reduce the number of operator applications for the Inexact-Newton based schemes. Such possibilities are left for future investigation and currently only the BGS preconditioner is used currently to resolve the neutronic system with upscattering.

In this setting, the problem is solved using power iteration, IRAM, SIPI-Newton hybrid iteration and Arnoldi-Newton hybrid iteration for various dominance ratios. The results shown below arise from the discretization of the homogenous problem using piecewise quadratic Lagrange basis functions on a structured grid with 400 QUAD8 elements and 1281 dofs/group. The cost results obtained from various test runs with a tolerance of 1E-10 are shown in Table (6.3). All the columns list the total number of BGS operator applications since the total cost is a function of this parameter.

It is quite evident that if the subspace size is increased for the Arnoldi iteration, the number of operator count will decrease until the optimal size is used. This was
shown by [90]. Since it is hard to calculate this optimal subspace size a-priori and that larger memory requirements prohibit storing many vectors, the situation where limited subspace size is prescribed has been considered here.

Table 6.3

<table>
<thead>
<tr>
<th>Length</th>
<th>DR</th>
<th>Power</th>
<th>CSIPI</th>
<th>SIPI-Newton</th>
<th>IRAM</th>
<th>IRAM-Newton</th>
</tr>
</thead>
<tbody>
<tr>
<td>12.255</td>
<td>0.3500</td>
<td>83</td>
<td>75</td>
<td>68</td>
<td>47</td>
<td>62</td>
</tr>
<tr>
<td>23.127</td>
<td>0.5000</td>
<td>147</td>
<td>121</td>
<td>96</td>
<td>73</td>
<td>89</td>
</tr>
<tr>
<td>85.95</td>
<td>0.9001</td>
<td>980</td>
<td>285</td>
<td>216</td>
<td>180</td>
<td>153</td>
</tr>
<tr>
<td>126.38</td>
<td>0.9500</td>
<td>1916</td>
<td>443</td>
<td>394</td>
<td>276</td>
<td>188</td>
</tr>
<tr>
<td>291.1</td>
<td>0.9900</td>
<td>7317</td>
<td>1133</td>
<td>563</td>
<td>353</td>
<td>284</td>
</tr>
<tr>
<td>926.5</td>
<td>0.9990</td>
<td>74270</td>
<td>5396</td>
<td>786</td>
<td>1488</td>
<td>692</td>
</tr>
<tr>
<td>2931</td>
<td>0.9999</td>
<td>604400</td>
<td>8478</td>
<td>1069</td>
<td>1899</td>
<td>782</td>
</tr>
</tbody>
</table>

The shift parameter for SIPI is found by performing several power iterations to coarse tolerance. The hybrid SIPI-Newton and Arnoldi-Newton schemes use a coarse tolerance of 1E-3 to obtain the initial guess and the eigenpair is converged to the user specified tolerance by the inexact Newton algorithm. For all calculations involving Arnoldi based schemes, the size of the subspace has been set at a fixed value of 10 for the current problem.

As expected, the results prove that as the dominance ratio approaches unity, the number of power iterations increase exponentially. The Arnoldi iteration converges to the dominant pair in much lesser iterations but the true power of this scheme, finding multiple eigenpairs simultaneously, is not utilized here. On the other hand, the hybrid SIPI-Newton scheme performs the quite well unlike the IAEA 2-D benchmark problem and this depends on the methodology to choose the initial shift parame-
ter. Hence, when a guess for the dominant eigenvalue is available, the SIPI-Newton scheme can be the optimal method of choice and but otherwise, the hybrid Arnoldi-Newton scheme provides a gain of at least factor of 2 as compared to just using IRAM scheme.

Clearly, results in Table (6.3) show the better convergence of the hybrid schemes for higher dominance ratio problems. This test result proves the feasibility of such schemes to resolve the fundamental eigenmodes for even strongly coupled (in terms of energy groups) problems. Further studies are necessary to improve the bootstrapping procedure and implementation of more efficient preconditioners for these block-symmetric system of equations.

6.1.5 Closing Remarks

The hybrid technique proposed using inexact Newton iteration is proven to be quite effective for the problems considered and delivers performance and convergence on-par to the state-of-art IRAM scheme. Future work is necessary to gauge the applicability of various kinds of preconditioners, other than the ones shown in this section, to improve computational efficiency. The flexibility of the MFNK framework to include these ideas apart from solving tightly-coupled multi-physics simulations, affirms that KARMA framework code can tackle linear, non-linear, eigenvalue and transient problems with different preconditioning approaches and coupling methods under "one roof".
6.2 Non-equilibrium Radiation Diffusion Physics Problem

The Boltzmann transport equation describes the behavior of radiation traveling through a material. It is generally considered as the most accurate model for the statistical average density of particles in a system, with very few assumptions. Non-equilibrium radiation transport physics deals specifically with the transport of photon energy and its coupling with a background material. It occurs at the heart of stars and several high energy physics systems [111], where there are steep energy and temperature gradients.

Several approximations that can be applied to the Boltzmann equation since solving the transport equation is computationally quite expensive. In the current work, we use the gray diffusion approximation with flux limiters [112]. The system of equations describing the radiation and material energy fields used here are given as

\[
\frac{\partial E}{\partial t} - \nabla \cdot (D_r \nabla E) = \sigma_a (T^4 - E), \tag{6.21}
\]

\[
\frac{\partial T}{\partial t} - \nabla \cdot (D_t \nabla T) = \sigma_a (E - T^4), \tag{6.22}
\]

where \( E, T \) are the radiation energy and material temperatures respectively, \( D_r, D_t \) are the radiation and temperature diffusion coefficients and \( \sigma_a \) is the photon absorption coefficient.

The definitions for these material properties [113] are

\[
\sigma_a = \frac{z^3}{T^3}, \tag{6.23}
\]

\[
D_r(T) = \frac{1}{3\sigma_a + (1/E)\left|\frac{\partial E}{\partial x}\right|}, \tag{6.24}
\]

\[
D_t(T) = kT^{5/2}, \tag{6.25}
\]

with \( z \) being the material atomic number and \( k \) is a constant.
Recently, the use of tightly coupled methods for non-equilibrium radiation diffusion with Jacobian-Free Newton Krylov methods has been analyzed [27, 20, 2] and its superiority over Operator-Split loosely coupled methods are provided. Based on these ideas, the KARMA framework is used in the current work to solve problems with different discretizations and coupling methods in a single code system.

Since radiation diffusion problems in general have strong gradients, traditional cG FEM spatial discretization of the elliptic operator in Eq. (6.22) is unstable without additional stability preserving terms added to the weak form. These family of methods, generally referred to as Stabilized Finite Element Methods (SFEM) [114] have enjoyed much success in advection dominated problems. The above set of equations can also be discretized with a dG(0) formulation or with the traditional Finite Volume Method (FVM) to conserve the radiation energy solution field throughout the transient. Preliminary studies with both these spatial treatments have shown that SFEM necessitates the selection of an optimal parameter in order to avoid unphysical solutions and is not guaranteed to be absolutely stable while FVM requires the usage of fine mesh resolution in order to gain better solution accuracy. Hence, all results shown in the current section will use FVM for spatial discretization with high number of elements to capture the energy and temperature profiles accurately.

Once the spatial scales are resolved by an appropriate spatial treatment, the temporal integration can be performed using $L$–stable schemes such as SDIRK2(2) and SDIRK3(3). It is also possible to use adaptive time-stepping strategies to choose a time step based on the dominating dynamical time scale of the problem. In order to correctly resolve the solution fields from both the radiation and material temperature accurately, a minimum of the time scales for the solution evolution is computed and used based on the following controller.

**Dynamical time scale controller:**

$$
\tau_{\text{dyn}}^n = \frac{S}{\frac{1}{\phi} \frac{\partial \phi}{\partial t}}_n \approx \Delta t_n \frac{S}{\phi_{n+1} - \phi_n} \frac{\phi_{n+1} - \phi_n}{\phi_{n+1}}
$$

(6.26)
where \( \phi \) is the solution field for current physics and \( S \) is a safety factor.

### 6.2.1 One-dimensional Problem

In order to analyze the different space-time discretization and coupling methods, we consider a one-dimensional model problem that consists of a unit radiation flux impinging on an initially cold slab of unit depth. This results in with Robin boundary conditions for the radiation equations at \( x = 0 \) and \( x = 1 \). These conditions are

\[
\frac{1}{4} E - \frac{1}{6\sigma_a} \frac{\partial E}{\partial x} = 1, \quad x = 0, \quad (6.27)
\]
\[
\frac{1}{4} E + \frac{1}{6\sigma_a} \frac{\partial E}{\partial x} = 0, \quad x = 1. \quad (6.28)
\]

The material temperature has homogenous Neumann conditions imposed at the boundaries. The initial solution fields are

\[
E(x, 0) = 10^{-5}, \forall x = [0, 1], \quad (6.29)
\]
\[
T(x, 0) = E(x, 0)^{1/4} \approx 0.0562. \quad (6.30)
\]

The material atomic number \( z \) for the homogenous medium is taken to be 1.0. Taking \( k = 0.1 \), we replicate the results provided by Mousseau et al. [113].

### 6.2.2 Results

The problem shown above was discretized with FVM using 800 elements and SDIRK3(3) method in both space and time, respectively, and solved using both weakly and strongly coupled methods. The solution profiles at a final time of \( T=2.5 \) seconds obtained using an adaptive time step controller are shown in Fig. 6.6.
Fig. 6.6. Non-equilibrium Radiation Diffusion Test Problem: Radiation and Material Temperature Profiles.

Note that the results given by Mousseau in [113] for the same test problem do not match the solutions shown here. The wave propagation speed of the energy source due to the left boundary condition is resolved differently in Fig. 6.6 but it converges consistently to a reference solution as the safety factor $S$ is reduced.

Next, we will look at the different tight coupling strategies (Picard versus Newton), based on the linearized physics-based Jacobian that is lagged at the previous step along with ILU(0) as an algebraic preconditioner. The results for the computational cost of the tightly coupled methods with constant time stepping using different step sizes are shown below.
Table 6.4
CPU time for one-dimensional problem using Picard vs Newton iteration

<table>
<thead>
<tr>
<th>$\Delta t$</th>
<th>Picard-Krylov</th>
<th>Newton-Krylov</th>
</tr>
</thead>
<tbody>
<tr>
<td>$4 \times 10^{-3}$</td>
<td>–</td>
<td>78.3</td>
</tr>
<tr>
<td>$2 \times 10^{-3}$</td>
<td>236.624</td>
<td>127.143</td>
</tr>
<tr>
<td>$1 \times 10^{-3}$</td>
<td>354.773</td>
<td>144.921</td>
</tr>
</tbody>
</table>

From Table 6.4, we can infer that the cost of the Newton-Krylov solution procedure increases sub-linearly as the number of steps is increased. Also due to the usage of the linearized Jacobian as only a preconditioner, higher time step sizes can be used to solve the system of equations. This is not the case for Picard iterations using the linearized Jacobian as the operator with ILU(0) as its preconditioner, since the solution starts to diverge for large time step sizes.

6.2.3 Closing Remarks

The flexibility of using the KARMA multi-physics code system is evident from the experiments conducted for the problems involving radiation diffusion physics using different spatial and temporal discretization schemes, along with a variety of consistent coupling methods. Also, using high-order methods with time adaptivity enables the ability to accurately capture the solution evolution based on its dynamical time scales. Further investigation is necessary to ascertain the reasons for the disparity in the results obtained here compared to Mousseau.
7. CONCLUSIONS

‘Computers are useless. They can only give you answers.’
– Pablo Picasso

Numerical simulation of multi-physics problems are difficult due to the need to resolve the stiff variations in spatial and temporal scales. This Dissertation used tight coupling methods for multi-physics problems in order to retain high-order spatio-temporal accuracy in the computed solution fields for stiff transients occurring in nuclear reactors and other fields (radiative transfer). The lessons learnt from using tight coupling methods for these problems are summarized below along with envisioned extensions to the KARMA framework that would provide further intuition for the development of efficient methods for coupled multi-physics problems.

7.1 Lessons Learned

1. Existing coupling methods using Operator-Splitting (OS) strategies were analyzed and the deficiencies in these methods including conditional stability and degradation of the higher order temporal accuracy were shown. In contrast, tightly coupled methods with either Picard or Newton iterations do restore the accuracy in the numerical solution, thereby reducing the total computational time where stability limits do not dominate the dynamical physical scales.

2. The tightly coupled methods can be unified under a single Matrix-free Nonlinear-Krylov (MFNK) framework that is based on a finite-differenced expression to obtain the action of the Jacobian matrix representing the coupled system matrix on a vector. Existing OS strategies offer intuitive knowledge regarding the important length and time scales to be tackled which can be used to create specialized schemes that serve as good preconditioners to reduce the total computational time in the MFNK technique. Note that usually such precon-
ditioners are used as solvers themselves if the coupling strength between the physics terms is weak.

3. The efficacy of the tightly coupled schemes are superior as compared to the loosely coupled OS schemes. Slight modifications on existing coupling strategies by introducing Picard-like iterations or performing fewer Newton iteration can improve the stability regions of the simple OS strategies.

4. The variations in time scales during the course of a multi-physics transient problem are often not dominated by a single physics component alone. The complex interplay of these temporal scales requires adaptive techniques in order to resolve the solution fields accurately in amenable wall-clock times. Tightly coupled schemes with adaptive time stepping procedures are excellent candidates to attain better efficacy using higher order, $L$–stable, temporal methods.

5. Method of Manufactured Solutions (MMS) and analytical methods are effective tools for the verification of non-linear multi-physics problems. Even though high fidelity physical models were not used in this Dissertation, the application of these techniques to even complicated cases is possible using symbolic mathematical toolboxes.

### 7.2 Future Work

The non-linear solution methods for multi-physics problems given in this Dissertation offers tremendous scope for future extensions. Few of the research areas that need to be focussed in the near future are listed below.

1. Create a multi-channel analysis code within the existing KARMA framework in order to apply the code system to problems in design and safety analysis of nuclear reactors.
2. Use the ability to create an array of varying fidelity physics models with the KARMA framework in order to verify the application of efficient, fidelity-independent coupling strategies.

3. Use spatial adaptive capabilities from hp-FEM ideas and adjoint based error estimators. Similar principles apply to time adaptivity using adjoint based temporal adaptivity.

4. If rigorous a-priori estimates for determining the strength of the coupling terms can be obtained based of $Jv$ and $\tilde{J}v$, where $\tilde{J}$ is the OS Jacobian matrix, an adaptive solution procedure selection can be made to choose from using either OS coupled strategy as the solver itself or as a preconditioner for the tightly coupled solution technique. This has tremendous potential to reduce the total runtimes in regions of the transient when the dynamical time scales are themselves well below the stability limits for OS coupling. Such algorithms can create computationally efficient, high accuracy schemes while making use of the unified MFNK framework proposed here.

5. The KARMA framework has extensive parallel capabilities since it is based on PETSc for all the relevant data-structures. Preliminary results for single-physics non-linear diffusion problems and coupled neutronics/thermal-conduction problems have shown linear speedup for up to 32 processors. Further studies need to be performed to measure the scalability of tightly coupled methods and for finding efficient parallel solve/preconditioning techniques for problems of interest in reactor analysis.

6. Finally, the KARMA framework is implemented with a flexible API. With only definitions of the semi-discrete non-linear residual that is consistent with the actual PDE for the physics model, the MFNK can be used to solve the non-linear, time-dependent problem. Improved efficiency can also be obtained if an external code can provide a suitable preconditioner to reduce the total Krylov
iterations. This lends the possibility to use existing single-physics codes in order to attain tightly coupled solutions, when necessary, thereby preserving numerous man-years of development efforts on these codes.
REFERENCES


A.1 MMS Script for Coupled Conduction/Fluid Problem

```matlab
clear all; clc;

% |=========| | yMAX
% | | Tf |
% y | 2-D | Tc (1-D: y)
% | | (x,y) |
% |=========| | 0
% 0 --x-- xMAX

syms x y t rF rT Ctf Ctc Tf Tc rho rhoU rhoE mu hc Dh
syms xMAX yMAX w del GAMMA Cp Cv RCONST Qconst G
syms dpdrho dpde kfluid Fw a b c p0 rho0 Tf0 Nu0 mu0

% Heat conduction PDE

% Ctf and Ctc are time constants -> Basically, you could
% make Ctf >> Ctc and this will introduce fast time scales
% due to heat conduction and slower scales due to heat
% removed by fluid.

% Fuel temperature exact solution
Tf = Tf0 + (1+tanh(Ctf*t)) * rF * ((0.5+ sin(pi/2*y/yMAX))' * (1+tanh(w+2/3-w*x/xMAX)));

% Coolant temperature exact solution
Tc = (1+tanh(Ctc*t)) * rT * (a-b*tanh(c+w-y/yMAX));

% viscosity variation with temperature
mu0 = 1.2075e-006 * subs(subs(Tc, t, 0), y, 0);```
% Density profile
rhoc = 219 ; f = 275.32 ; g = 511.58 ; Tc0 = 2503.7 ;
rho = rhoc + f*(1-Tc/Tc0) + g*(1-Tc/Tc0).^0.5 ;
inte = Cv * Tc ;

% Velocity constant in space-time
v = G./rho ;

% momentum = mass flux
m = rho .* v ;

% Total energy
e = rho.*inte + 0.5*m.*m./rho ;

% linearized EOS
p = p0 + dpdrho * (rho-rho0) +
    dpde * (Tc-sub(subs(Tc, t, 0), x, 0)) ;
M = v ./ sqrt(dpdrho) ;

% viscosity variation with temperature
mu = 1.2075e-006 * Tc ;

% http://www.cheresources.com/convection.shtml
Nu = Nu0 * (mu./mu0).^0.14 ;

hc = Nu * kfluid / Dh ; % Heat Transfer Coefficient
Re = G * Dh / mu ; % Reynolds number
Fw = 0.3164 / Re^0.25 ; % Blasius friction factor

% Thermal conductivity for fuel.
k = 6400.0./Tf^2 * \exp(-16.35/Tf) ;

\% \% \% \% \% \% \% \% FORCING FUNCTIONS \% \% \% \% \% \% \% \% \% \%

STf = \text{diff}(Tf, t) - \text{diff}(k \times \text{diff}(Tf, x), x)
- \text{diff}(k \times \text{diff}(Tf, y), y)
+ hc \times \text{subs}(Tf, x, xMAX) - Tc) ;

Scont = \text{diff}(rho, t) + \text{diff}(m, y);

Smom = \text{diff}(m, t) + \text{diff}(m \times v, y) + \text{diff}(p, y)
+ Fw \times v \times \text{abs}(v) ;

Sener = \text{diff}(e, t) + \text{diff}((e+p) \times v, y)
- hc \times \text{subs}(Tf, x, xMAX) - Tc) ;

disp(’STf’); ccode(STf)
disp(’Scont’); ccode(Scont)
disp(’Smom’); ccode(Smom)
disp(’Sener’); ccode(Sener)

Code Snippet A.1 MMS Script for Coupled Conduction/Fluid Problem
A.2 MMS Script for Coupled Neutronics/Conduction Problem

```plaintext
syms x y z Lz t s T egroups dgroups k avgnu Bg2
syms LX LY CT CF rT rF PI normalization_const
syms DEFAULT_FUEL_TEMP doppler_coeff k0 k1 rho cp
syms beta_tot PHI1 PHI2 xsrem1 xsrem2 xsfiss1 xsfiss2
syms xsrem01 xsrem02 xsnufiss1 xsnufiss2
syms xsdiff1 xsdiff2 energy_per_fission1 energy_per_fission2
syms invvell invvel2 xsscat12 xsscat21

sx = sin(x/LX*PI) ;
sy = sin(y/LY*PI) ;

T = CT*(1+tanh(rT*t))*sx*sy ;
k = k0 + k1*(T - DEFAULT_FUEL_TEMP) ;

egroups = 2 ; % energy groups
dgroups = 2 ; % delayed precursor groups

xsnufiss1 = avgnu * xsfiss1 ;
xsnufiss2 = avgnu * xsfiss2 ;

xsrem1 = xsrem01 + doppler_coeff*(sqrt(T) - sqrt(DEFAULT_FUEL_TEMP)) ;
xsnufiss2 = xsrem02 ;

beta(1) = sym('beta[1]') ;
beta(2) = sym('beta[2]') ;
lambda(1) = sym('lambda[1]') ;
lambda(2) = sym('lambda[2]') ;
beta_tot = beta(1) + beta(2) ;

% 2-D geometric buckling
Bg2 = PI*PI*(1/LX^2+1/LY^2) ;
```
\[ \text{PHI1} = CF \times (1 + \exp(rF \times t)) \times x \times y \times (x/LX \times y/LY) \]

\[ \text{PHI2} = \text{PHI1} \times xscatt12 \div (xsrem2 + xsdiff2 \times Bg2) \]

\textbf{FOR} \ dg = 1 : dgroups

\% Initial precursor concentration
\begin{verbatim}
PREC_0(dg) = beta(dg) \times (subs(xsnufiss1 \times PHI1 +
    xsnufiss2 \times PHI2, t, 0)) / lambda(dg) ;
\end{verbatim}

\% Analytically solve the precursor ODE
\begin{verbatim}
PREC(dg) = ( PREC_0(dg) + beta(dg) \times int(
    subs(
        (xsnufiss1 \times PHI1 +
        xsnufiss2 \times PHI2) \times \exp(lambda(dg) \times s),
        t, s),
    s, 0, t) ) \times \exp(-lambda(dg) \times t) ;
\end{verbatim}

\textbf{end}

\% FORCING FUNCTIONS

\% Fuel Temperature
\begin{verbatim}
srcT = rho \times cp \times diff(T, t) - diff(k \times diff(T, x), x) -
    diff(k \times diff(T, y), y) - normalization_const \times
    (energy_per_fission1 \times xsfiss1 \times PHI1 +
    energy_per_fission2 \times xsfiss2 \times PHI2) ;
\end{verbatim}

\% Fast Flux
\begin{verbatim}
srcFLX1 = invvel1 \times diff(PHI1, t) - diff(xsdiff1 \times diff(PHI1, x), x) -
    diff(xsdiff1 \times diff(PHI1, y), y) + xsrem1 \times PHI1 -
    (1-beta_tot) \times (xsnufiss1 \times PHI1 + xsnufiss2 \times PHI2) -
    xsscatt21 \times PHI2 ;
\end{verbatim}

\textbf{if} dgroups > 0

\begin{verbatim}
for \ dg = 1 : dgroups
    srcFLX1 = srcFLX1 - lambda(dg) \times PREC(dg) ;
    srcPREC(dg) = diff(PREC(dg), t) -
    beta(dg) \times (xsnufiss1 \times PHI1 + xsnufiss2 \times PHI2) +
    lambda(dg) \times PREC(dg) ;
end
\end{verbatim}
% Thermal Flux
srcFLX2 = invvel2 * \text{diff}(\text{PHI2},t) - \text{diff}(\text{xsdiff2} * \text{diff}(\text{PHI2},x),x)
- \text{diff}(\text{xsdiff2} * \text{diff}(\text{PHI2},y),y) + \text{xscatt12} * \text{PHI2}
- \text{xsrem2} * \text{PHI2};

\text{Power} = \text{normalization}\_\text{const} * (\text{energy}\_\text{per}\_\text{fission1} * \text{xsfiss1} * \text{PHI1}
+ \text{energy}\_\text{per}\_\text{fission2} * \text{xsfiss2} * \text{PHI2});

% Total power in the domain as a function of time
totPower = \text{int} (\text{int} (\text{Power}, y, 0, \text{LY}), x, 0, \text{LX});

codeT = \text{ccode}(\text{T})
codeFLX1 = \text{ccode}(\text{PHI1})
codeFLX2 = \text{ccode}(\text{PHI2})
\text{for} \ dg = 1 : \text{dgroups}
    \text{fprintf} (\text{`\ncodePREC{%d} = \n
'}, dg);
    \text{disp} (\text{ccode}(\text{PREC}(dg)))
end

codeST = \text{ccode}(\text{srcT})
codeSFLX1 = \text{ccode}(\text{srcFLX1})
codeSFLX2 = \text{ccode}(\text{srcFLX2})
\text{for} \ dg = 1 : \text{dgroups}
    \text{codeSPREC}(dg) = \text{ccode}(\text{srcPREC}(dg));
    \text{fprintf} (\text{`\ncodeSPREC{%d} = \n
'}, dg);
    \text{disp} (\text{codeSPREC}(dg))
end

codePower = \text{ccode}(\text{totPower})

Code Snippet A.2 MMS Script for Coupled Neutronics/Conduction Problem
APPENDIX B
CROSS-SECTION DATA FOR EIGENVALUE PROBLEMS

B.1 2-D Two-group IAEA Benchmark Problem

This problem is based on the benchmark introduced in ANL [115] and contains 4 materials. The cross-sections are obtained from the 3D data by accounting for leakage in z-direction with a $B g_z^2 = 8 \times 10^{-5}$.

The lattice configuration is of the form given below with each assembly having dimensions $= [10, 10]$ cms.

```
3 1 1 1 1 1 1 3 1 1 1 1 1 2 2 4 4
1 1 1 1 1 1 1 1 1 1 1 1 2 2 4 4
1 1 1 1 1 1 1 1 1 1 1 1 2 2 4 4
1 1 1 1 1 1 1 1 1 1 1 1 2 2 4 4
1 1 1 1 1 1 1 1 1 1 1 2 2 2 4 4
1 1 1 1 1 1 1 1 1 1 2 2 2 4 4 4
1 1 1 1 1 1 1 1 1 1 2 2 4 4 4 4
1 1 1 1 1 1 1 1 1 2 2 4 4 4 4 4
3 1 1 1 1 1 1 3 3 2 2 2 2 4 4 0 0
3 1 1 1 1 1 1 3 3 2 2 2 2 4 4 0 0
1 1 1 1 1 1 2 2 2 2 4 4 4 4 0 0
1 1 1 1 1 1 2 2 2 2 4 4 4 4 0 0
1 1 1 1 1 1 2 2 2 2 4 4 4 4 0 0
1 1 1 1 1 1 2 2 2 2 4 4 4 4 0 0
1 1 1 1 1 1 2 2 2 2 4 4 4 4 0 0
1 1 1 1 1 1 2 2 2 2 4 4 4 4 0 0
2 2 2 2 2 2 2 2 2 2 4 4 4 4 0 0 0 0
2 2 2 2 2 2 2 2 2 2 4 4 4 4 0 0 0 0
2 2 2 2 2 2 2 2 2 2 4 4 4 4 0 0 0 0
4 4 4 4 4 4 4 0 0 0 0 0 0 0 0 0 0
4 4 4 4 4 4 4 0 0 0 0 0 0 0 0 0 0
```
B.2 Cross-section Data for 2-D Two-group Homogenous Medium Problem

\[ D_1 = 1.0, D_2 = 0.4 \]
\[ \Sigma_{r,1} = 0.04, \Sigma_{r,2} = 0.08 \]
\[ \nu \Sigma_{f,1} = 0.01, \nu \Sigma_{f,2} = 0.13 \]
\[ \Sigma_{s,1\rightarrow2} = 0.02, \Sigma_{s,2\rightarrow1} = 0.001 \]
VITA

Vijay Subramaniam Mahadevan was born in Chennai, India. After high school, he attended the Regional Engineering College, affiliated with Bharathidasan University, Trichy, India and graduated in May 2002 with a Bachelor of Technology degree in chemical engineering. After working two years at Dell Computers Inc., Bangalore, India as a software developer, he entered the nuclear engineering graduate program at Texas A&M University in August 2004. After graduating with a Master of Science degree in December 2006, he continued to pursue his research interests in numerical methods for coupled multi-physics simulations as a Graduate Research Assistant. In Fall 2008, he received the Givens Fellowship from Argonne National Lab. Apart from the fellowship, he has spent several summers at Idaho National Lab and worked with research scholars who have similar interests.

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