

SEXUAL DIMORPHISM IN THE *Sceloporus undulatus* SPECIES COMPLEX

A Thesis

by

DREW EDWIN DITTMER

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

MASTER OF SCIENCE

August 2012

Major Subject: Wildlife and Fisheries Sciences

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Approved by:

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## ABSTRACT

Sexual Dimorphism in the *Sceloporus undulatus* Species Complex.

(August 2012)

Drew Edwin Dittmer, B.S., University of Missouri-Columbia

Co-Chairs of Advisory Committee: Dr. Toby J. Hibbitts  
Dr. Lee A. Fitzgerald

The Fence Lizard (*Sceloporus undulatus* complex) is a wide ranging North American species complex occurring from the eastern seaboard westward through the great plains and central Rocky Mountains and into the American Southwest. A recent phylogeny suggests four species lineages occur within *S. undulatus*. Traits within an interbreeding species that are influenced by sexual selection are under different selection pressures and may evolve independently from the selective forces of habitat.

*Sceloporus* lizards have several characters that are influenced by sexual selection. I investigated sexual size dimorphism and allometric relationships of body size (snout vent length), torso length, rear leg length and three measurements of head size in 12 populations from the four species in the *S. undulatus* complex (N=352) specifically looking for variation among the 4 species. Additionally I investigated the size of signal patches between males and females in three species (N=339 specimens of *S. consobrinus*, *S. cowlesi*, *S. tristichus*) of the *S. undulatus* complex. Sexual confusion, was recently described in a population of the *Sceloporus undulatus* complex occurring in White Sands, New Mexico and the behavior is correlated with variation in badge size

between male and female lizards. To make inferences about sexual confusion at the species level I investigated the presence and absence of signal patches in female lizards, and compare the sizes of signal patches between males and females. My analyses suggest that torso length and head size are significant sources of sexual size dimorphism but the findings differ from earlier published investigations of sexually dimorphic characters in the species complex. I also find support for the *S. undulatus* complex being generally a female larger species complex. However two of the 12 populations I investigated displayed male biased sexual size dimorphism. Analysis of signal patches across three species of the *S. undulatus* complex suggests that sexual dimorphism in signal patch size for *S. cowlesi* and *S. tristichus* may not prevent sexual confusion. While the near total absence of signal patches in female *S. consobrinus* is evidence that sexual confusion is not possible with regards to signal patches.

## DEDICATION

This thesis is dedicated to my grandfather Wilbur Edwin Meinershagen. He always supported my interest in reptiles and amphibians and when I was 11 years old he helped me capture my first *Sceloporus* lizard in New Mexico. He was also the best listener I've ever known.

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I wish to thank all members of the Herpetology Lab at Texas A&M, Dr. Wade A. Ryberg, Daniel J. Leavitt, Michael L. Treglia, Nicole Smolenskey and Christopher M. Schalk. Each member provided valuable feedback during my research, and often provided just the right amount of distraction that kept my graduate school experience fun and engaging.

Finally I wish to thank Dr. Douglas A. Eifler and his wife Dr. Maria Eifler. Doug and Maria have included me on much of their field research since 2003. Doug and Maria encouraged me pursue an advanced education studying reptiles and amphibians and are

two of the most genuine people I have ever known. I would not have made it in this far without their mentorship, guidance, and support.



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## CHAPTER I

### INTRODUCTION

Phenotypic differences between males and females are known as sexual dimorphisms. These differences in most cases cannot be explained by natural selection. Darwin (1871) proposed the theory of sexual selection to explain sexual dimorphism. Three patterns of sexual dimorphism are commonly observed. The first and often easiest to observe is the asymmetry in size between sexes or sexual size dimorphism (SSD). The second is the presence of weapons such as antlers or horns which are often referred to as armaments. The third common dimorphism is the presence of conspicuously colored scales, feathers, fins or hair, referred to as ornaments. All three of these dimorphisms can occur simultaneously. Commonly these dimorphisms are described as signals, which presumably communicate some aspect of individual fitness to (Hibbitts et al. 2007). An important question surrounding these signals is what these signals are advertising and to what audience are the signals being directed.

Signals that are considered armaments are generally used in male-male contests (Andersson 1994). In regards to SSD in species where males are larger than females, size in and of itself can be advantageous in contests for females (Darwin 1871; Ghiselin 1974) and may also function as an armament. The form of male-male competition known as endurance rivalry can be explained in a SSD context as larger males are able to defend larger territories, engage in more territorial disputes and

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therefore breed with more females (Andersson 1994). Armaments and SSD can also be signals that are subject to the selective forces of female choice. Females may choose to breed with larger males or males with the best armaments as males possessing superior traits may be signaling their physical condition, foraging ability, or they may be defending territories that have the best resources for the female's offspring (Andersson 1994)

Assuming females to be the discriminating sex, male-larger SSD is easily explained within the bounds of sexual selection, but female larger SSD is more complex and difficult to explain. Additionally there are no known cases among vertebrate species where females possess armaments that are larger than those of males, or where armaments are absent in males but present in females. Furthermore female larger SSD is the most common pattern of SSD among vertebrates and is rarely explained by sex role reversal, where large females compete for breeding access to males (Andersson 1994; Fairbairn 2007). Large female size is best explained as contributing to an increase in reproductive output. This is not necessarily a sexually selected trait (Andersson 1994) and is most commonly explained by the female fecundity advantage hypothesis, where larger females are able to produce either more or larger offspring (Darwin 1871; Ghiselin 1974; Andersson 1994). However in some cases males may actually select larger females (Côte and Hunte 1989), and in these instances both sexual selection and fecundity selection may be acting in concert. Additionally it has been shown that growing to a larger size takes time and energy and larger females may not have higher reproductive success over their lifetimes (Shine 1988). Further complicating female SSD

is the small male advantageous hypothesis. Small male size may be advantageous in scramble competitions, where small males mature earlier and find females faster. Small male size may also be beneficial in resource poor environments, where fewer resources are required to actively seek females (Andersson 1994).

Sexual Size Dimorphism and armaments are challenging to explain when considered from a female perspective. Armaments play a direct role in settling disputes, compared to ornaments which have been described as being uncostly to produce, and can be manipulated to deceive conspecifics (Whiting et al. 2003). The examples of signal based deception are numerous (Warner 1984; Gross 1991; Sinervo and Lively 1996). Much of the research on ornaments as signals has focused on their function in sexual selection contexts and has normally taken a decidedly male perspective. However, like SSD, signals can also be subject to selective forces other than sexual selection (Ryan et al. 1982), and signals are observed in females of many species. Lizards, in particular have been the focus of studies investigating the function of conspicuous color signals in females (Stuart-Fox and Whiting 2005; Chan et al. 2009). In many lizards conspicuous colors are sexually dichromatic, and females bear the strongest coloration while they are gravid (Vinegar 1972; Cooper 1984; Jones and Lovich 2009). Research has found support for the rejection hypothesis, where gravid females display these signals to courting males in an effort to avoid unnecessary copulation (Vinegar 1972; Cooper 1984; Chan et al. 2009).

While some female lizards have conspicuous colors strongly associated with the gravid condition, other lizards, including the genus *Sceloporus*, have similar conspicuous

signal patches in males and females. Specifically, in the four species of the *Sceloporus undulatus* complex, blue and black signal patches are present on the throat and belly. These signal patches are common in both males and females in at least two of the four species (*S. cowlesi* and *S. tristichus*). A field investigation by Robertson and Rosenblum (2010) investigated signal patches in *S. cowlesi* and described a behavior that is termed “sexual confusion”, which is related to the overlap in signal patch size of males and females (2010). In *S. cowlesi* sexual confusion occurs when resident male confuses another male as a female. There are a couple of studies reporting male lizards interpreting conspecific males as females and courting or even breeding with them (Noble and Bradley 1933; Vinegar 1972), but Robertson and Rosenblum’s 2010 work is the first to quantify and correlate the behavior to the variation in a sexually dichromatic signal.

Prior to 2002, the *Sceloporus undulatus* complex was considered to be one species with as many as ten subspecies occurring from the eastern and gulf coasts of the United States to the Rocky Mountains and the American southwest. Leaché and Reeder (2002) identified four species within the *S. undulatus* complex, (*S. undulatus*, *S. consobrinus*, *S. tristichus* and *S. cowlesi*) and no subspecies are presently recognized (Leaché 2009). Therefore the *S. undulatus* complex is an excellent group to explore SSD patterns at a large geographic scale and to explore the labile nature of sexual dimorphism (Cox et al. 2009). There are very few Broad scale investigations of SSD across geographically wide ranging species or genera (Roitberg 2009), and the *S. undulatus* species complex offers the opportunity to look for patterns of sexual dimorphism of



sexually selected traits that occur in a species with stereotyped territorial defense (Cooper Jr and Burns 1987). *Sceloporus undulatus* species complex has been a model organism for investigations of Sexual Size Dimorphism (SSD) as well as studies addressing evolution, ecology and morphology (Haenel et al. 2003; Pinch and Claussen 2003; Robertson and Rosenblum 2010). The recognition of four species within the *S. undulatus* species complex provides an opportunity to describe variation in SSD within each one of these species and to look for patterns that may have been obscured by considering *S. undulatus* as a single widespread species with multiple subspecies (Vinegar 1975; Fitch 1978).

Herein I describe my approach to investigate sexual dimorphism of body size, sexually selected morphological characters and conspicuously colored signal patches among the four species of the *Sceloporus undulatus* complex. My project has two major objectives that will be addressed in two chapters. First I will describe sexual size dimorphism (SSD) and the sexual dimorphism of selected traits in the *S. undulatus* species complex. Specifically, I will determine if patterns of SSD and the relationship of head size to body size are consistent within the newly described clades of the *S. undulatus* species complex. Also, I will determine if support for the female fecundity advantage and small male advantage hypotheses exists within the species complex. Second, I will quantify the size of signal badges in three species of the *S. undulatus* complex, *S. consobrinus*, *S. cowlesi*, and *S. tristichus*. I will compare my findings to those of Robertson and Rosenblum (2010) and determine if similar overlap in badge size among the three species and between males and females within species provides

evidence of sexual confusion in populations other than that observed by Robertson and Rosenblum (2010).

## CHAPTER II

### VARIATION IN SEXUAL SIZE DIMORPHISM AMONG 4 SPECIES IN THE

#### *Sceloporus undulatus* SPECIES COMPLEX

##### Introduction

Patterns of sexual dimorphism can evolve through natural selection, sexual selection, or fecundity selection. Most commonly SSD evolves due to sexual selection (Darwin 1871). This primarily occurs in two ways: by contest competition between members of the same sex or by mate choice where in most cases females select males based on secondary sexual characters (Andersson 1994). Another common source of SSD is through fecundity selection (Darwin 1871). This occurs when females have secondary sexual characters that develop to accommodate increased fecundity of females such as increased torso length or width enabling a female to carry more or larger eggs. Sexual size dimorphism can also develop through natural selection (Andersson 1994; Wikelski and Trillmich 1997; Székely et al. 2000). For example, males and females of file snakes have extreme differences in adult body size that is best explained by dietary differences between the sexes (Houston and Shine 1993).

In species where males are the larger sex (male-biased SSD) or where males have weapons (e.g. horns, claws, or increased muscle mass), contest competition favors the larger individual or the larger weapon (Darwin 1871; Ghiselin 1974). The winner of such contests contribute their genes for larger body size or larger weapons to the next generation resulting in male-biased SSD (Ghiselin 1974; Lindenfors et al. 2009). Females may also show a preference for males that are larger or that have larger

weaponry, which could lead to male-biased SSD, but this has received little attention due to the difficulty of determining a female's preference and the difficulty of disentangling selection for male-biased SSD via contest competition (Andersson 1994; Olsson et al. 2002; Hibbitts et al. 2007).

While male-biased SSD can generally be explained by sexual selection, across all animals females are usually the larger sex (Andersson 1994; Fairbairn 2007). The dominant explanation for female-biased SSD is the female fecundity-hypothesis, where larger females can produce more and or larger offspring (Darwin 1871). Female-biased SSD due to the female fecundity hypothesis occurs occasionally in a few mammals such as baleen whales, vespertilionid bats and hares (Ralls 1976). In birds like wading birds and some predatory birds, female-biased SSD is fairly common (Andersson 1994). Fish and amphibians are largely female-biased (Andersson 1994; Fairbairn 2007). In reptiles turtles are the group with the most consistent pattern of female-biased SSD (Andersson 1994; Cox et al. 2009). Female-biased SSD is also supported by the small male advantage hypothesis, where smaller males have a selective advantage because they require fewer food resources and are able to mature earlier and can therefore devote more resources towards acquiring mates. Examples of this hypothesis have been found most commonly in fish, butterflies and endoparasites (Ghiselin 1974; Andersson 1994).

Among the vertebrates, reptiles are popular organisms for studies of SSD, and provide a variety of opportunities to study the selective forces operating on male-biased and female-biased SSD. Lizards generally show male-biased SSD, but the alternative is not unusual (Cox et al. 2009) and except for studies of a few species (Pianka and Parker

1975; Zamudio 1998) reasons explaining female-biased SSD are understudied in lizards (Andersson 1994; Cox et al. 2009). Morphological characters associated with contest competition and fecundity selection are most commonly observed to exhibit SSD in lizards (Sugg et al. 1995; Herrel et al. 1999; Herrel et al. 2001; Olsson et al. 2002). A few studies have attempted to describe the variation in SSD across wide spread genera or species (Fitch 1978; Hibbitts et al. 2005; Roitberg 2009), but most studies of SSD in lizards have focused on the mechanisms and processes that influence SSD patterns within localized lizard populations (Shine et al. 1998; Cox 2005; Cox 2007; Kratochvíl and Frynta 2009). In particular, the lizard species *Sceloporus undulatus* has been a model organism for investigations of SSD as well as studies addressing evolution, ecology and morphology (Haenel et al. 2003; Pinch and Claussen 2003; Robertson and Rosenblum 2010). Prior to 2002 *Sceloporus undulatus* was considered one species occurring from the eastern and gulf coasts of the United States to the Rocky Mountains and the American southwest. The use of molecular systematics has helped to identify four species within the *S. undulatus* complex (*S. undulatus*, *S. consobrinus*, *S. tristichus* and *S. cowlesi*) (Leaché and Reeder 2002; Leaché 2009). An improved phylogenetic hypothesis offers the opportunity to describe variation in SSD and to look for patterns that may have once been confounded by the historical taxonomy of the group (Fitch 1978).

My study targets the variation in snout-vent length (SVL), rear leg length, torso length and three measurements of head size to quantify patterns of SSD within the 4 species of the *S. undulatus* species complex. I aim to determine if these species display

female-biased or male-biased SSD, and if traits normally associated with contest competition (head size) show positive allometry in males. For support of the small-male advantage hypothesis I expect females to be the larger sex, and in males I expect isometry or negative allometry of traits associated with female fecundity and contest competition. For support of the female-fecundity advantage hypothesis I expect to see positive allometry of torso length and female-biased SSD in SVL. My analyses identify species-specific patterns of variation in SSD that are likely attributable to different selective forces acting on populations of the 4 species.

#### Materials and Methods

Previous published information on size at sexual maturity of both males and females on the formerly wide ranging *S. undulatus* (Tinkle and Ballinger 1972; Cooper and Vitt 1989) is no longer accurate in light of the recent phylogenetic hypothesis that splits the *S. undulatus* species complex into four species (Leaché and Reeder 2002). . Therefore, I chose to use Q-Q plots to cull small lizards from my samples prior to conducting any additional analyses. A Q-Q- plot can be used for assessing one dimensional data sets to identify outliers and to assess the normality of a data set distribution (Wilk and Gnanadesikan 1968). I checked Q-Q plots with SVL paired against each morphological measure; individual outliers that represented small lizards were culled from our sample.

I measured 6 morphological characters on 179 female and 173 male museum specimens on the *S. undulatus* complex. The following measurements were taken on each lizard using digital calipers: snout-vent length (SVL), head length (HL), head depth

(HD), head width (HW), torso length (Torso), and right rear leg length (RRL). All measurements were natural log-transformed prior to analysis. For each of the four species I chose three populations (n=12 total populations). I chose populations based on maps in Leaché and Reeder (2002). To quantify SSD and to determine male-biased or female-biased SSD I used the size dimorphism index (SDI) described in Lovich and Gibbons (1992)  $((\text{larger sex}/\text{smaller sex})-1)$ . The SDI is preferable to other methods that quantify SSD because the results are symmetrical around 0 and directional (Roitberg 2009).

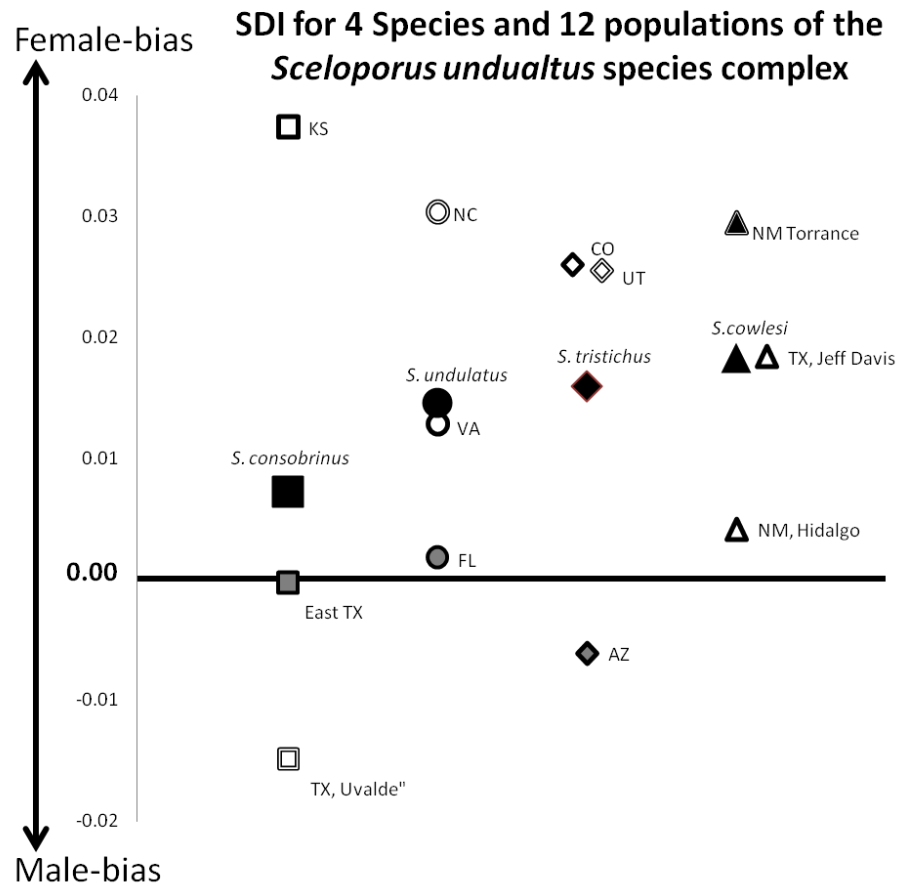
To determine the morphological characters that express the most variation I used Microsoft Excel® with the PopTools add-in (Hood 2010) to perform a principle components analysis (PCA) on the size measurements. The PCA has been used to describe patterns of sexual dimorphism in lizards (Zamudio 1998) and results from PCA informed my choice of which morphological characters to subject to reduced major axis (RMA) regression. It is advisable to use RMA when both the categorical and dependent variable are measured with error (Johnson et al. 2005; Fairbairn 2007). I used RMA to determine variation in allometry across the 4 species and 12 populations. RMA has been used to describe allometry in lizards and other taxa (Johnson et al. 2005; Schulte-Hostedde et al. 2011).

To perform RMA I used RMA v. 1.17 (Bohonak 2004) and entered SVL as the dependent variable and Torso, RLL, HL, HW and HD as the independent variables. The RMA was run for all lizards combined, each of the four species combined, and for each population of lizards.

## Results

*Sexual Dimorphism Index*—Females were the larger sex when analyzing all lizards measured within the *S. undulatus* species complex (Figure 1). Females were also generally larger when analyzing SDI for each species separately. One population of *S. consobrinus* from east Texas showed no noticeable sexual size dimorphism in relation to SVL (SDI=-0.000306). *Sceloporus consobrinus* from Uvalde Co., Texas showed the highest degree of male-biased size dimorphism and *S. tristichus* from Coconino Co., Arizona was the only other population to show male-biased dimorphism. Among the four species, the greatest variation in SDI between populations was for *S. consobrinus* ( $SDI_{\max}-SDI_{\min}=0.0523$ ) followed by *S. tristichus* ( $SDI_{\max}-SDI_{\min}=0.0322$ ). *Sceloporus undulatus* and *S. cowlesi* showed little difference in degree of SDI ( $SDI_{\max}-SDI_{\min}=0.0286$ ; and  $SDI_{\max}-SDI_{\min}=0.0254$  respectively).





**Figure 1.** Sexual size dimorphism index (SDI) for all four species groups and all 12 populations of the *S. undulatus* complex. Points are jittered to more clearly show populations with equivalent SDI values. Black shapes are the results for all populations pooled into their respective species. Individually colored shapes show the results for each population analyzed independently.

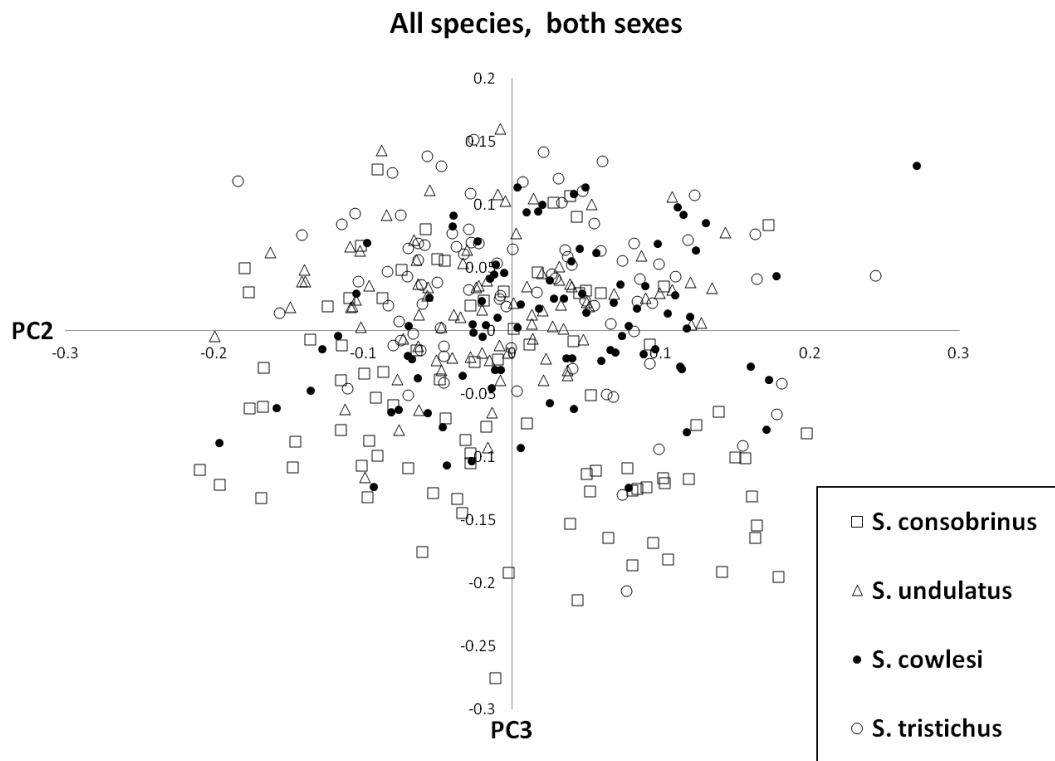
*Principle Component Analysis*--The PCA for all lizards yielded 5 eigenvectors. The first eigenvector (PC1) explained 85.38% of the cumulative variation, and showed strong positive loadings of similar magnitude for all measured variables (Table 1). This can be interpreted as being a measure of overall size across all individuals. Because the SDI is preferable to PCA for interpreting the degree of SSD in regards to body size for

each species I did not use PC1 scores in my analyses. The second (PC2) and third (PC3) eigenvectors explained an additional 12.89% of the variation and were strongly influenced by the measures of Torso, RLL and HD, while PC4 and PC5 were influenced entirely by measures of head size but only explained an additional 3.19% of the variation

**Table 1.** Results of PCA showing eigenvalues, cumulative variances and eigenvectors. Loadings in boldface are considered the most significant for this study

| Eigenvectors for all Variables |        |         |              |                |                |                |               |               |
|--------------------------------|--------|---------|--------------|----------------|----------------|----------------|---------------|---------------|
| PC Scores                      | % Var  | Cum Var | Eigen Values | HL             | HW             | HD             | TORSO         | RLL           |
| PC 1                           | 0.8538 | 0.8538  | 0.0992       | 0.4128         | 0.4940         | 0.4551         | 0.4462        | 0.4236        |
| PC 2                           | 0.0661 | 0.9199  | 0.0077       | -0.1342        | -0.2755        | -0.3957        | <b>0.8651</b> | -0.0342       |
| PC 3                           | 0.0483 | 0.9681  | 0.0056       | 0.1379         | -0.2294        | <b>-0.4358</b> | -0.2181       | <b>0.8312</b> |
| PC 4                           | 0.0200 | 0.9881  | 0.0023       | <b>-0.6717</b> | -0.3192        | <b>0.5710</b>  | 0.0688        | 0.3408        |
| PC 5                           | 0.0120 | 1.0000  | 0.0014       | <b>0.5843</b>  | <b>-0.7250</b> | 0.3470         | 0.0140        | -0.1115       |

When PC2 scores were plotted against PC3 scores, a separation was observed in quadrant IV of the cartesian plane (Figure 2). Most of the individuals in quadrant IV were lizards from the Uvalde population and all of the males measured from Uvalde fall into this quadrant. This quadrant can be explained as lizards with short RLL and long Torso. It is interesting that *S. consobrinus* from Uvalde are one of only two populations



**Figure 2.** Scatter plot of PC2 and PC3 including both sexes and 4 species analyzed. PC2 is primarily a measure of Torso Length, and PC3 is primarily measure of increasing rear leg length and decreasing head depth.

measured to show male-biased SSD. Uvalde is relatively close to the Trans Pecos area of West TX which loosely defines part of the boundary between *S. cowlesi* and *S. consobrinus*. However the PCA showed almost no overlap with the individuals from Uvalde and individuals of *S. cowlesi*. Further, more *S. tristichus* from Coconino Co., AZ showed male-biased SSD but did not show any separation from the other lizards measured.

*Reduced Major Axes Regression (RMA)*--Male lizards showed no positive allometry for any of the 6 morphological measurements when regressed against SVL (Table 2 for summarized results, full RMA results are in Appendix 2). Male *S. consobrinus* had negative allometry for Torso, HD, HL and HW and isometric allometry for RLL. *Sceloporus undulatus* males demonstrated negative allometry only for HD and HW, all other measures were isometric. Male *S. tristichus* show negative allometry only for Torso, and all other measures were isometric. *Sceloporus cowlesi* males demonstrate negative allometry only for HD and HW and all other measures were isometric.

**Table 2.** Summary of reduced major axes regressions showing variation in allometric growth across the *S. undulatus* complex and sample sizes. Allometric growth is expressed as Negative (N), Positive (P), or Isometric (0). The rows corresponding to the species name are a composite value representing the pooled populations

|                        | Torso |        | Rear Leg Length |        | Head Depth |        | Head Length |        | Head Width |        | Sample Size | Sample Size |
|------------------------|-------|--------|-----------------|--------|------------|--------|-------------|--------|------------|--------|-------------|-------------|
|                        | Male  | Female | Male            | Female | Male       | Female | Male        | Female | Male       | Female | Male        | Females     |
| <i>S. consobrinus</i>  | N     | 0      | 0               | 0      | N          | 0      | N           | P      | N          | 0      | 44          | 46          |
| Kansas                 | 0     | 0      | 0               | 0      | N          | 0      | N           | P      | N          | 0      | 15          | 15          |
| Uvalde, TX             | 0     | N      | 0               | 0      | 0          | 0      | 0           | P      | N          | 0      | 15          | 15          |
| East, TX               | 0     | 0      | 0               | 0      | N          | N      | 0           | 0      | 0          | 0      | 15          | 16          |
| <i>S. undulatus</i>    | 0     | 0      | 0               | P      | N          | 0      | 0           | P      | N          | 0      | 42          | 45          |
| Virginia               | N     | 0      | 0               | 0      | 0          | 0      | 0           | P      | 0          | P      | 15          | 15          |
| North Carolina         | 0     | 0      | 0               | 0      | 0          | 0      | 0           | P      | 0          | 0      | 15          | 15          |
| Florida                | 0     | N      | 0               | 0      | 0          | 0      | 0           | 0      | 0          | 0      | 12          | 15          |
| <i>S. tristichus</i>   | N     | 0      | 0               | 0      | 0          | 0      | 0           | 0      | 0          | 0      | 43          | 46          |
| Colorado               | 0     | N      | 0               | 0      | N          | 0      | 0           | 0      | 0          | 0      | 15          | 15          |
| Utah                   | N     | N      | 0               | 0      | 0          | 0      | 0           | 0      | 0          | 0      | 15          | 15          |
| Arizona                | 0     | 0      | 0               | 0      | 0          | 0      | 0           | 0      | 0          | 0      | 13          | 16          |
| <i>S. cowlesi</i>      | 0     | 0      | 0               | P      | N          | 0      | 0           | P      | N          | 0      | 43          | 42          |
| Hidalgo, NM            | 0     | 0      | 0               | 0      | 0          | 0      | 0           | P      | 0          | 0      | 13          | 12          |
| Torrance, NM           | 0     | N      | 0               | 0      | N          | 0      | 0           | 0      | 0          | 0      | 15          | 15          |
| Jeff Davis, Texas      | 0     | 0      | 0               | 0      | N          | 0      | 0           | P      | 0          | 0      | 15          | 15          |
| <b>All Individuals</b> | 0     | 0      | N               | P      | N          | 0      | 0           | 0      | N          | 0      | 173         | 179         |

At the population level males of the *S. undulatus* species complex were mostly isometric. Male *S. undulatus* from Virginia and *S. tristichus* from Utah yield negative allometry for Torso. In the following populations, males show a pattern of negative allometry: *S. undulatus* from Kansas and Uvalde Co., Texas, *S. tristichus* from Colorado, and *S. cowlesi* from Torrance Co., New Mexico and Jeff Davis Co., Texas. Head width was the only other body measurement where male *S. consobrinus* from Kansas and Uvalde Co., TX show negative allometry.

Female lizards yielded much broader variation in allometric relationships. Across all individuals females demonstrate isometry for Torso, HD, HL and HW and positive allometry for RLL. The RMA analysis showed no negative allometry when all females were included together. Female *S. consobrinus* showed positive allometry for HL and isometry for all other measured variables. *S. undulatus* and *S. cowlesi* females showed positive allometry for RLL and HL and otherwise demonstrated isometry. The female *S. tristichus* group is the only female group analyzed to display isometry for all morphological measures.

At the population level female lizards generally showed isometry for all traits. For measures of Torso females from the following populations showed negative allometry: *S. consobrinus* from Uvalde Co., TX, *S. undulatus* from Florida, *S. tristichus* from Colorado and Utah, and *S. cowlesi* from Torrance Co., New Mexico. Female *S. consobrinus* from East Texas showed negative allometry for HD and are the only other population to show negative allometry among all females. For the variable HL females from the following populations have positive allometry: *S. consobrinus* from Kansas and

Uvalde Co., Texas, *S. undulatus* from Virginia and North Carolina, *S. cowlesi* from Hidalgo Co., New Mexico and Jeff Davis Co., Texas. Female *S. undulatus* from Virginia showed positive allometry for HW and were the only other population with a positive allometric relationship.

### Discussion

Despite two populations showing male-biased SSD, the *S. undulatus* species complex in general can be described as having female-biased SSD. This pattern of SSD is explained by two hypotheses that are not mutually exclusive: the female fecundity hypothesis and the small male advantage hypothesis. Considering the latter hypothesis, in female-larger species, selection should favor small males because their small size requires fewer food resources, allowing more time for them to search out potential mates (Blanckenhorn 1999). I found that males were isometric or had negative allometry when comparing SVL to HD, HW, HL, Torso and RLL. I expected this pattern to be present if the female-biased SSD was explained by the small male advantage hypothesis. However the known behavioral patterns of lizards in the *S. undulatus* species complex challenge parts of the small male advantage hypothesis, as these lizards are known to defend territories through the use of push-up displays, (Jones and Lovich 2009; Robertson and Rosenblum 2010) which would normally be associated with male-biased SSD. Female-biased SSD is usually only associated with rare cases of reversed sex roles where females select for males (Andersson 1994). Other signals are presumably used in contests between males in the *S. undulatus* species complex including their throat and belly patches as well as the aforementioned push-up displays. In the dwarf chameleon, also in which females are

larger, the fighting ability of males has been determined to have little to do with body size and more to do with specific traits (casque size and flank patch). These traits are associated with aggressiveness of individual males in male-male contests (Stuart-Fox et al. 2006). Interestingly, although none of the populations I studied showed positive allometry between SVL and head size, it has been found previously in the *S. undulatus* species complex (Cooper and Vitt 1989). This evidence would suggest the small male advantage hypothesis would not be an appropriate explanation for female-biased SSD in all populations, but the results do indicate that populations with morphological characteristics consistent with contest competition are rare. Future research aimed at describing aggressiveness between interacting males in the *S. undulatus* species complex could help elucidate what factors determine a successful male and provide further support to small male advantages and how male territoriality plays a role.

Although most populations had female-biased SSD I found inconclusive support for the female fecundity hypothesis. No populations showed positive allometry for Torso, a trait associated with increased fecundity, and five populations showed negative allometry. I observe two species (*S. consobrinus* and *S. tristichus*) where males show negative allometry for Torso. This could suggest that at least for some populations an increase in body size in females is adequate to increase fecundity and that males discontinue investment in Torso as they age, similar to the pattern found by Sugg et al (1995). It has been determined that in some horned lizard species body size does increase fecundity. However an increase in the degree of SSD does not yield a fecundity advantage in *P. hernandesi* (Zamudio 1998). I note however that body shape in the



genus *Phrynosoma* evolved largely due to adaptations for dietary specialization on ants (Pianka and Parker 1975).

It is clear that fecundity selection and sexual selection contribute to SSD in many organisms. However, Cox et al. (2003) suggested that sexual selection contributes to only 16% of the variation in evolutionary shifts of SSD in lizards (Cox et al. 2003). Cox's study only looked at instances of male-biased SSD in relation to sexual selection, therefore the contribution of lizards with female-biased SSD and the role of fecundity selection is unclear. Natural selection also contributes to mating success and SSD. Predation has been identified as a contributor or inhibitor to differences in SSD (Ryan et al. 1982; Magnhagen 1991). Research into the selective pressures on the *S. undulatus* species complex could identify other significant influences on body size and morphology (Blanckenhorn 2000) such as greater predation on larger males, or increased survival of females with longer legs. Further comparative field studies that identify both sexual selection forces on each sex and other selective pressures may shed light on the variations of SSD observed in the *S. undulatus* species complex.

In addition to predation hormones, especially testosterone, have been shown to have an effect on variety of SSD patterns in closely related species of *Sceloporus* lizards (Cox 2005). Testosterone has been shown to inhibit male growth in female larger species and promotes male growth in male larger species (Cox 2005). The variation in SDI and the variety of allometric relationships observed in the study re-enforce the description of the *Sceloporus* genus as having directionally labile SSD (Cox et al. 2009). Further

studies examining the effects of testosterone on the *S. undulatus* species complex could identify a hormonal mechanism to the observed variation.

A third explanation for the variation observed in the study is natural selection on ecomorphs. Ecomorphs are well known in the *S. undulatus* species complex and the variety of ecomorphs contributed to the abundant sub-species descriptions prior to the most recent phylogeny (Ferguson et al. 1980; Leaché and Reeder 2002). The PC2 axis described increasing torso lengths as other characters decrease, and on the PC3 axis a gradient of long legged to shallow headed lizards. More simply put I find a gradient of faster lizards to slower lizards across the PC3 axis (Miles et al. 1995; Bonine and Garland Jr 1999), thus the PCA is highlighting the existence of the ecomorphs within this group.

In conclusion *S. undulatus* complex is generally a female-biased species group, in relation to SSD. I find support for the small male advantage hypothesis in that males have negative allometry for measures of head size, and are generally smaller than females. I do not find strong support for the female fecundity advantage hypothesis as no measures of fecundity had positive allometry.

## CHAPTER III

### SEXUAL DIMORPHISM AND PATCH SIZE VARIATION IN THREE LIZARD SPECIES SUGGESTS POTENTIAL FOR SEXUAL CONFUSION

#### Introduction

Many organisms possess traits that convey information about species recognition, fighting ability, or influence female choice of mates. For example the gray tree frog exists as two cryptic species (*Hyla chrysoscelis* and *Hyla versicolor*) that are genetically incompatible and males of each species have different calling rates that serve as auditory signals directed toward potential mates (Littlejohn et al. 1960). Female *H. chrysoscelis* and *H. versicolor* are receptive only to the calls of their respective species (Gerhardt 1982; Gerhardt 1994). Fighting ability can be advertised with weapons or armaments, (i.e. teeth, antlers, horns etc.). Armaments can serve simultaneously to signal fitness to females (Clutton-Brock et al. 1982) and the fighting ability of a male to conspecific males. For instance in male caribou, violent fights between males of matched antler size is a rare occurrence, but light sparring and assessment of mismatched opponents is common (Barrette and Vandal 1990). Signals that generally influence female choice occur in the form of conspicuous colors and ornaments. *Anolis* lizards display a colorful dewlap to potential mates, and experiments that prevented males from extending the dewlap or altered the color of the dewlap reduced the likelihood that a male would be chosen as a mate (Crews 1975; Sigmund 1983).

Compared to armaments, ornaments can be subject to cheating or alternative mating strategies, as ornaments do not require physical force to verify their

effectiveness (Whiting et al. 2003). In many cases cheating the expression of an ornament involves a male mimicking a female, these individuals are considered female mimics or sneaker males (Sinervo and Lively 1996; Wikelski et al. 1996). Female mimicry can be advantageous as it can enable sneaker males to exist in other male territories without engaging in territorial disputes. Thus these sneaker males can obtain mating opportunities that would otherwise be unavailable to them.

The use of ornaments as signal patches, and exploiting these signals in cheating behavior is well studied in male lizards (Hews et al. 1994; Sinervo and Lively 1996; Whiting et al. 2003; Whiting et al. 2009), but only a few studies have investigated the use of signal patches in females. These studies have supported sex role reversal, or the uncommon pattern of females competing with other females for breeding access to males (Andersson 1994; Eens and Pinxten 2000). Particularly in lizards, signal patches and conspicuous female colors are relatively common, and color patches commonly signal to males that the female has already bred and is gravid (Cooper 1984). However, many female lizards have color patches that are similar to those of males of the same species. For most of these patches it is not known what they are signaling or even if they function as a signal or are just ancestral characters without a current known function. In the family Phrynosomatidae, the genus *Sceloporus* has many species that display both male throat and belly patches (also known as badges), and female throat and belly patches. However, among members of the *Sceloporus undulatus* species complex, female lizards often have large ventral patches on the belly and throat. These patches can range from being small and faint to being very similar to the patches of males.

A population of *Sceloporus cowlesi* occurring in White Sands, New Mexico has been shown to overlap in the size of signal patches between males and females and inter and intrasexual interactions may result in sexual confusion (Robertson and Rosenblum 2010). Robertson and Rosenblum (2010) found that male *S. cowlesi* would often attempt to court conspecific males with smaller badges. This is the first time the term sexual confusion has been applied to any species in the *Sceloporus undulatus* species complex, but it is not the first description of courtship between two males. Vinegar (1972) observed several cases of male-male courtship and one case of a female using courtship behavior in an interaction with a con-specific female in *S. tristichus*. While Vinegar (1972) admitted that he was unable to correlate same sex courtship with signal patches or coloration, it is Robertson and Rosenblum who successfully showed that signal patch size is related to sexual confusion in at least one population of *S. cowlesi*.

I chose to investigate patch size in multiple populations across the range of *S. cowlesi* to determine if the observations of Robertson and Rosenblum (2010) were an isolated example of sexual confusion that was occurring in a unique habitat situation. I also chose to do the same analysis with the two closest relatives of *S. cowlesi*, *S. consobrinus* and *S. tristichus* to determine if sexual confusion could potentially be occurring in these taxa as well.

#### Materials and Methods

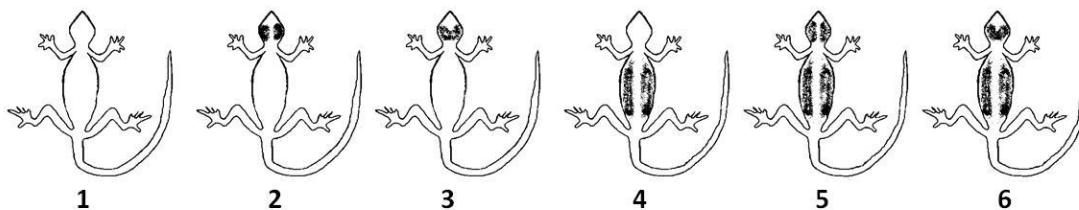
I used a total of 339 museum specimens to quantify sexual dimorphism of signal patches between males and females of the three species; *Sceloporus consobrinus*(*N*), *S. cowlesi*(*N*), and *S. tristichus*(*N*) (Table 3 see Appendix for locality information). No

known morphological characters exist to confidently identify the three species used in this study from one another. Presently the best way to identify museum specimens of these three species is by classifying them according to locality. I used range maps (Powell 2011) and HerpNet (Spencer 2010) to identify specimens from within the described range of each species. I used a SONY Digital Camera (Model Model DSC-H20) mounted on a photo table to photograph the venter all specimens. The camera was positioned 25 centimeters above the specimen and a metric ruler was placed adjacent to every specimen. To measure the variation in the sizes of signal patches between the sexes I used tpsDig2 to outline throat and belly patch size and to measure snout to vent length (SVL), all photos were calibrated to record measurements in millimeters before outlining any patches. The area of signal patches was calculated using the `area.pl` function from the `splancs` package for R, this function calculates the area of non self-intersection polygons (Rowlingson 2012). To ensure only adult lizards were analyzed I used Q-Q plots to examine the distribution of SVL. Individual outliers that represented small lizards were culled from the sample. A Q-Q plot can be used for assessing one dimensional data sets to identify outliers and to assess the normality of a data set distribution (Wilk and Gnanadesikan 1968).

**Table 3** The number of female and male specimens measured from each of the three species .

|               | <i>Sceloporus consobrinus</i> | <i>Sceloporus cowlesi</i> | <i>Sceloporus tristichus</i> |
|---------------|-------------------------------|---------------------------|------------------------------|
| <b>Female</b> | <b>42</b>                     | <b>63</b>                 | <b>64</b>                    |
| <b>Male</b>   | <b>42</b>                     | <b>68</b>                 | <b>60</b>                    |
| <b>Total</b>  | <b>84</b>                     | <b>131</b>                | <b>124</b>                   |

Prior to analyzing the area of signal patches I categorized the type and frequency of patches occurring between males and females of the three species. Throat patches occurred in two forms, as a single colored patch covering the ventral surface of the head and neck, or as two separate patches on the right and left. When present, belly patches always occurred as a right and left patch. The combination of presence and absence of throat and belly patches resulted in 6 categories; (1) no throat or belly patches, (2) two throat patches and no belly patches, (3) one throat patch and no belly patches, (4) no throat patch and two belly patches, (5) two throat patches and two belly patches, (6) one throat patches and two belly patches (Figure 3). A histogram was used to show the number of males and females in each species that are represented in the 6 categories (Figure 4)



**Figure 3** Individual specimens of *S. consobrinus*, *S. cowlesi* and *S. tristichus* were classified according to six categories; (1) no patches (2) two throat patches (3) one throat patch (4) two belly patches and no throat patches (5) two throat patches and two belly patches (6) one throat patch and two belly patches

All statistical analyses were performed using JMP™. The area of the signal patches were summed into a total badge area (TBA) and natural log transformed (LN) (the untransformed measures of TBA and SVL are reported in the Appendix C). To correct for body size in size corrected the LN TBA I divided each value by the corresponding LN SVL, this method has been used to standardize size measures in several studies of reptiles (Lee 1980; Brennan et al. 2008; Doan 2008; Santos and Llorente 2008). This method of size correction is preferable to using residuals from a regression, as the residuals would have produced negative values giving a false sense of variance for lizards that have no measurable TBA. I used a one-way ANOVA to test the null hypothesis that the mean TBA of males and females were equal within a species. I also used a Two One-Sided Tests (TOST) approach to test for a difference between the means of males and females from the ANOVA. The TOST tests that the difference between means is less than or equivalent to a pre-determined value, and is used as assurance that means do not differ by much or are practically equivalent (Ramírez and Ramírez 2009). I employ the TOST and use the minimum male TBA for each respective



species to test the null hypothesis that the difference in mean TBA between males and females is less than the minimum male TBA ( $\mu_{\text{Male TBA}} - \mu_{\text{Female TBA}} \leq \text{Male Minimum TBA}$ ). Rejecting the null hypothesis suggests that sexual dimorphism of badge size is sufficient to prevent sexual confusion in the species. Failure to reject the null hypothesis is evidence that the overlap in TBA of males and females could result in sexual confusion between males and females of a species. This test is intended to illustrate the potential for sexual confusion correlated with variations in badge size as reported by Robertson and Rosenblum (2010) in *Sceloporus cowlesi*

## Results

Females occurred in all six categories of badge and throat patch combinations, while males never occurred in categories one, two or three (Figure 4). Because categories four, five and six have both the largest patches and/or the largest number of patches they have the highest TBA and are the most “male-like” categories. Female *S. tristichus* had more representatives in these male-like categories than female *S. cowlesi* while female *S. consobrinus* were never represented in “male-like” categories. Categories five and six are those that have belly patches and throat patches and *S. tristichus* and *S. cowlesi* males were predominately in these categories while *S. consobrinus* males were relatively evenly distributed across categories four, five, and six. Conversely, male *S. tristichus* had only two individuals in category four and *S. cowlesi* males were not represented in category four.

*Sceloporus consobrinus* males and females were significantly different in respect to TBA ( $p < 0.0001$ , Table 4). Only five of the 42 females had measureable badges. Additionally male *S. consobrinus* always had two belly patches and most had belly patches and throat patches. The TOST also show that *S. consobrinus* are significantly different in relation to TBA (Table 4). The box plots visually illustrate that the difference in TBA for male and female *S. consobrinus* is the greatest for all three species (Figure 5).

**Table 4. ANOVA and TOST of the difference between female and male TBA for *S. consobrinus***

**ANOVA of *S. consobrinus* Male and Female TBA**

| Source | DF | Sum of Squares | Mean Square | F Ratio  | Prob>F |
|--------|----|----------------|-------------|----------|--------|
| Sex    | 1  | 32.9697        | 32.9697     | 835.7350 | <.0001 |
| Error  | 82 | 3.2349         | 0.0394      |          |        |
| Total  | 83 | 36.2046        |             |          |        |

**Means for Oneway ANOVA**

| Sex    | Number | Mean    | Std Error | Lower 95% CI | Upper 95% CI |
|--------|--------|---------|-----------|--------------|--------------|
| Male   | 42     | 0.09549 | 0.03065   | 0.0147       | 0.1763       |
| Female | 42     | 1.34849 | 0.03065   | 1.2677       | 1.4293       |

**TOST for *S. consobrinus* Male and Female Mean TBA**

| Values testing for Practical Equivalence | Test            | t Ratio | p-Value |
|--|-----------------|---------|---------|
| Minimum Male TBA                         | Upper Threshold | 1.8924  | 0.9690  |
| Actual Mean Difference                   | Lower Threshold | 55.9258 | <.0001  |
| Std Error of Difference                  | Max Over Both   |         | 0.9690  |

*Sceloporus cowlesi* males and female were significantly different in respect to TBA ( $p < 0.0001$ , Table 5), while the results of the TOST show that that the difference between male and female TBA is not significantly different from minimum male *S. cowlesi* TBA. Most females measured 0 for TBA, but thirty-five females (greater than half) had positive measures for TBA. Thirteen female *S. cowlesi* had measures for TBA that included large belly patches. The box plots visually show that female *S. cowlesi* show a large range of values for TBA and that some overlap exists in TBA between males and females (Figure 5).

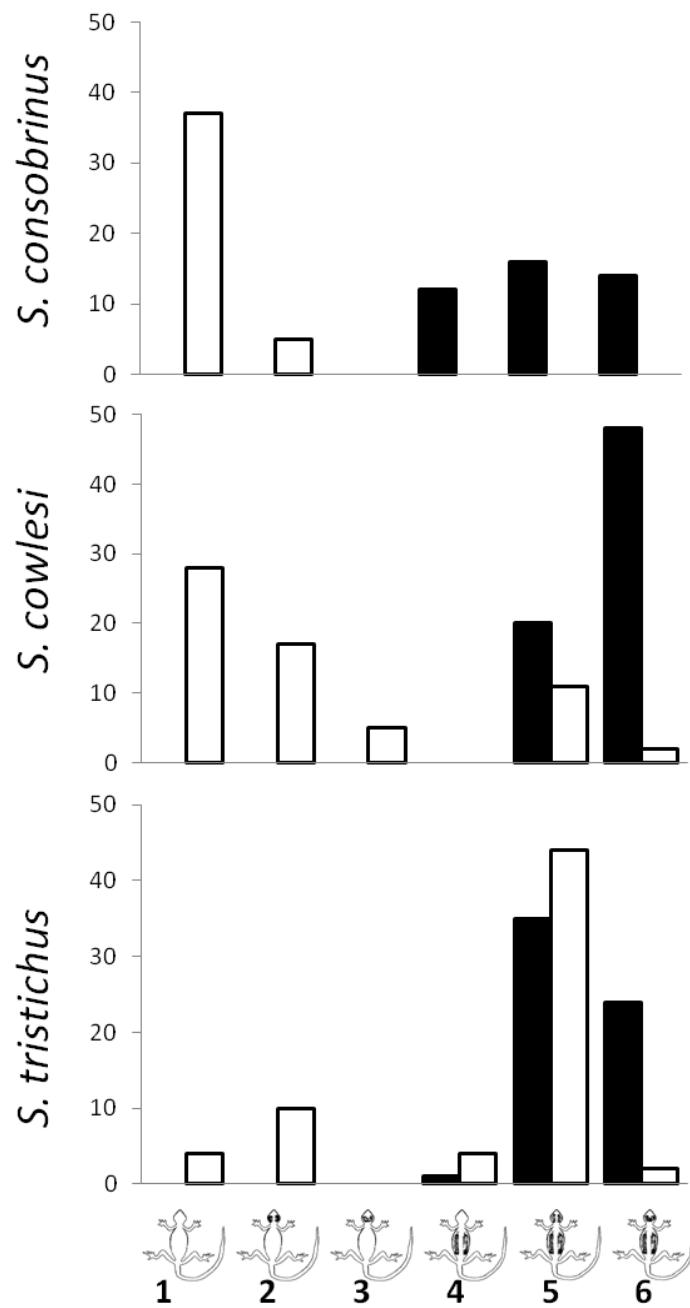
**Table 5. ANOVA and TOST of the difference between female and male TBA for *S. cowlesi***

| <b>ANOVA of <i>S. cowlesi</i> Male and Female TBA</b>      |               |                        |                    |                     |                     |
|--|---------------|------------------------|--------------------|---------------------|---------------------|
| <b>Source</b>  | <b>DF</b>     | <b>Sum of Squares</b>  | <b>Mean Square</b> | <b>F Ratio</b>      | <b>Prob&gt;F</b>    |
| <b>Sex</b>   | 1             | 23.7340                | 23.7340            | 162.7925            | <.0001              |
| <b>Error</b>   | 129           | 18.8073                | 0.1458             |                     |                     |
| <b>Total</b>   | 130           | 42.5412                |                    |                     |                     |
| <b>Means for Oneway ANOVA</b>                              |               |                        |                    |                     |                     |
| <b>Sex</b>   | <b>Number</b> | <b>Mean</b>            | <b>Std Error</b>   | <b>Lower 95% CI</b> | <b>Upper 95% CI</b> |
| Male   | 63            | 0.5706                 | 0.04811            | 0.4448              | 0.6964              |
| Female   | 68            | 1.4225                 | 0.04630            | 1.3014              | 1.5436              |
| <b>TOST for <i>S. cowlesi</i> Male and Female Mean TBA</b> |               |                        |                    |                     |                     |
| <b>Values testing for Practical Equivalence</b>            |               | <b>Test</b>            | <b>t Ratio</b>     | <b>p-Value</b>      |                     |
| <b>Minimum Male TBA</b>                                    | 1.316847      | <b>Upper Threshold</b> | -6.9632            | <.0001              |                     |
| <b>Actual Mean</b>   | 0.8519        | <b>Lower Threshold</b> | 35.4813            | <.0001              |                     |
| <b>Difference</b>  |               | <b>Max Over Both</b>   |                    | <.0001              |                     |
| <b>Std Error of Difference</b>                             | 0.0668        |                        |                    |                     |                     |

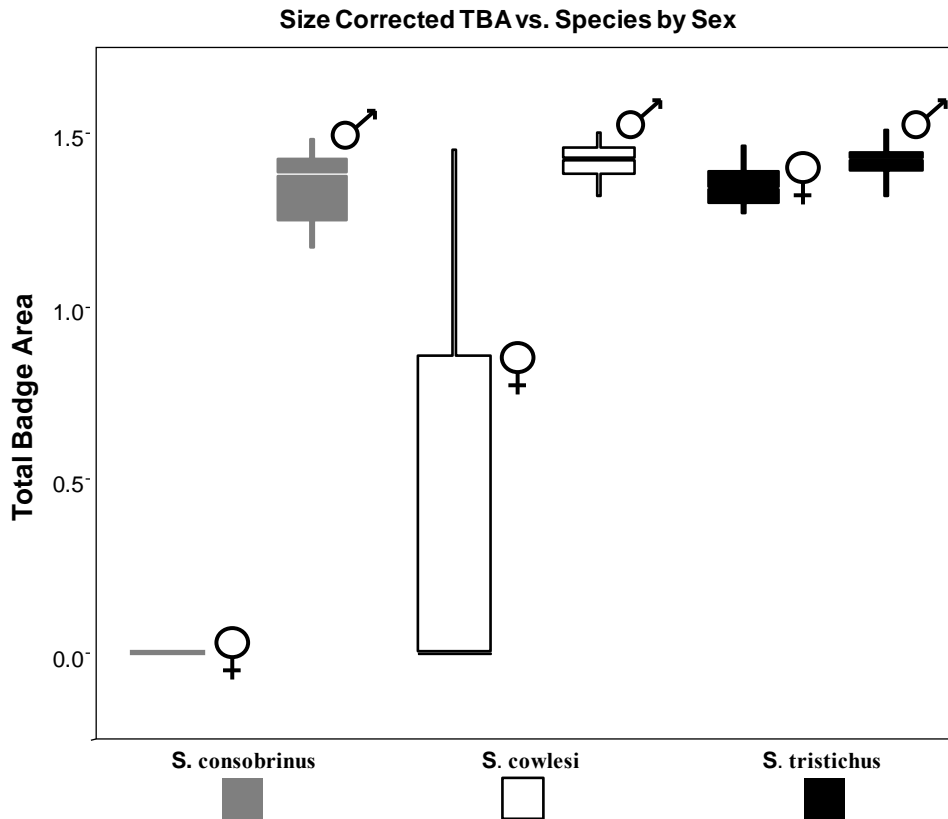
For *S. tristichus* there are significant differences between male and female TBA ( $p < 0.0001$ , Table 6). But the results of the TOST show that the means do not differ by less than the male *S. tristichus* minimum TBA. For female *S. tristichus* only 4 females measured 0 for TBA. Most female *S. tristichus* had belly patches and throat patches and therefore had relatively large measures of TBA. The box plots visually show that the difference in TBA between male and female *S. tristichus* is the smallest of all three species (Figure 5).

**Table 6. ANOVA and TOST of the difference between female and male TBA for *S. tristichus***

| <b>ANOVA of <i>S. tristichus</i> Male and Female TBA</b>      |               |                        |                    |                     |                     |
|---|---------------|------------------------|--------------------|---------------------|---------------------|
| <b>Source</b>   | <b>DF</b>     | <b>Sum of Squares</b>  | <b>Mean Square</b> | <b>F Ratio</b>      | <b>Prob&gt;F</b>    |
| Sex   | 1             | 1.547706               | 1.54771            | 20.9218             | <.0001              |
| Error   | 122           | 9.025032               | 0.07398            |                     |                     |
| <b>Total</b>  | <b>123</b>    | <b>10.572737</b>       |                    |                     |                     |
| <b>Means for Oneway ANOVA</b>                                 |               |                        |                    |                     |                     |
| <b>Sex</b>  | <b>Number</b> | <b>Mean</b>            | <b>Std Error</b>   | <b>Lower 95% CI</b> | <b>Upper 95% CI</b> |
| Male  | 64            | 1.19159                | 0.03400            | 1.1243              | 1.2589              |
| Female  | 60            | 1.41515                | 0.03511            | 1.3456              | 1.4847              |
| <b>TOST for <i>S. tristichus</i> Male and Female Mean TBA</b> |               |                        |                    |                     |                     |
| <b>Values testing for Practical Equivalence</b>               |               | <b>Test</b>            | <b>t Ratio</b>     | <b>p-Value</b>      |                     |
| <b>Minimum Male TBA</b>                                       | 1.315062      | <b>Upper Threshold</b> | -22.3324           | <.0001              |                     |
| <b>Actual Mean Difference</b>                                 | 0.223558      | <b>Lower Threshold</b> | 31.4805            | <.0001              |                     |
| <b>Std Error of Difference</b>                                | 0.048875      | <b>Max Over Both</b>   |                    | <.0001              |                     |



**Figure 4.** Frequency of males and females in the 6 signal patch categories by species.



**Figure 5.** Box plots showing the range of TBA for males and females by species. The box plot for female *S. consobrinus* represents the only 4 females that have measures of TBA greater than 0.

## Discussion

In *S. undulatus* species complex, sex recognition has been determined to be based on the extent of signal patch size (Cooper Jr and Burns 1987). But in lizards with significant overlap in signal patch size, males with smaller badge sizes are interpreted as females (Robertson and Rosenblum 2010). I found that sexual dimorphism in signal patches is significant enough to prevent sexual confusion in *S. consobrinus*. However, in *S. cowlesi* and *S. tristichus* enough overlap in TBA exists (Figure 5) to suggest that

sexual confusion may be occurring. I sampled specimens across a wide geographic distribution therefore the data suggest that sexual confusion in *S. tristichus* and *S. cowlesi* is possible at the species level. Female variation in TBA was the largest in *S. cowlesi* and females in some populations did not have individuals in the “male-like” categories. It could be argued that *S. cowlesi* lives across the widest ranges of habitats, from the novel white and dark sands region into rocky outcrops and deserts. The variation observed in *S. cowlesi* in particular could be the result of many localized adaptations to the wide variety of habitats, this speculation agrees with explanations proposed by Robertson and Rosenblum (Robertson and Rosenblum 2009; Robertson and Rosenblum 2010).

The sexual confusion observed by Robertson and Rosenblum (2010), where a male attempts to court or breed with another male, is not specific only to sceloporine lizards. Males of the species *Ameiva chrysolema* were observed attempting to breed with one another by Noble and Bradley (1933). Attempted male-male courting in other species of *Ameiva* and in *Aspedoscelis sexlineata* have also been observed (Noble and Bradley 1933). In *Ameiva chrysolema* males are generally larger than females, and incidences of sexual confusion were thought to be associated with overlap in the size of males and females. Unlike *Ameiva chrysolema*, *Aspedoscelis sexlineata* has sexually dichromatic green color patches on males, and these patches are reduced or absent in females. Noble and Bradley (1933) observed that these patches are displayed in aggressive male-male interactions, but male lizards courted and bred with females as well as males that failed to display aggressive behavior (Noble and Bradley 1933). These

instances of sexual confusion in *Ameiva* and *Aspedoscelis* are further complicated by the fact that female mimicry is not known in the family Teiidae. These few studies that have documented sexual confusion suggest that the behavior may not be rare. Only Robertson and Rosenblum (2010) offered an explanation for sexual confusion. They posit that sensory drive and ecological speciation in a novel habitat are responsible for sexual confusion in White Sands. It remains to be verified if sexual confusion occurs in other populations of *S. cowlesi* or if it occurs at all in *S. tristichus*. If the evidence from signal badges provided here is shown to result in sexual confusion in more populations and species, then alternative explanations for sexual confusion will be needed.

The investigations of Noble and Bradley (1933) are similar to those of Vinegar (1972) and Robertson and Rosenblum (2010), in that they focus on the interactions of males and the interpretation of sexual signals from the perspective of male lizards. Furthermore data on how signal patches function in *Sceloporus* lizards is biased towards male-male interaction and female-choice experiments. Very few experiments have looked at territoriality in female lizards (Mahrt 1998) or how signal badges function in female-female interactions or how males perceive females with badges larger than their own. Future work targeting interactions between sexes from the perspective of female lizards is necessary to further explain the existence of large female signal patches in *Sceloporus cowlesi* and *Sceloporus tristichus*. Furthermore, investigations focusing on female-female interactions will offer insight into the function of signaling and conspicuous colors in female lizards



## CHAPTER IV

### CONCLUSIONS

My results show that SSD in particular is highly variable at the population level, and some populations display male-larger dimorphism or no SSD which is inconsistent with descriptions of this species complex being female-larger. Furthermore, no measure of head size showed a positive allometric relationship for males of any species or any population. Perhaps more interesting is that head length did have a positive allometric relationship for females of all species except *S. tristichus*. It is difficult to speculate on the positive allometry of head size in females, but future studies could investigate the relationship of throat patches to head size in females. It may be possible that females have larger heads and use throat patch displays more than males in interactions with conspecifics.

Considering the female fecundity advantage hypothesis, when populations are pooled into their respective species, torso length shows isometric allometry with body size. This suggests that larger females do gain a fecundity advantage with an increase in body size. However, growing to a larger body size requires more time and resources, and this has been shown to counter the reproductive advantages of body size (Shine 1988). It should also be pointed out that females from five of the twelve populations I measured actually show negative allometry for torso length relative to body size. This might suggest that there is plasticity in torso length. Future studies that investigate life history and population dynamics with regard to the presence of different morphologies will be beneficial for understanding these variations.

Like the female fecundity advantage hypothesis, the small male advantage hypothesis seems to apply well to some populations of lizards, and poorly to others. Generally speaking females are larger than males in the *S. undulatus* complex, but there are two populations which show male-larger SSD and two more that show little or no SSD. While this clearly supports Cox et al's (2007) description of the labile nature of SSD in the *Sceloporus* genus, it does little to explain why there is a variety of SSD patterns within populations of the same species. It is possible that the small male advantage is supported for populations of the *S. undulatus* complex occurring in resource limited environments, but field studies will be needed to confirm this.

In consideration of the merits of the analyses I used to investigate SSD, my results highlight the value of using RMA regression for describing allometric relationships. RMA is preferable when both the predictor and the predicted variables are measured with error. Moreover, my use of RMA to analyze allometry of head size yielded results different from those previously published for *S. undulatus* that used ANCOVA (Cooper and Vitt 1989). This suggests that previously published results of allometric relationships of head to body size in lizards and allometric relationships of other taxa may benefit from being reanalyzed with RMA.

This investigation contributes to the importance of evaluating descriptive characteristics in species in concert with species trees generated from molecular phylogenetics. Certainly molecular phylogenetics is an excellent tool for elucidating evolutionary relationships of species. However, molecular phylogenetics leaves something to be desired when previously assumed morphological characters no longer

apply to many species. This is especially true for species that have a rich history of morphological descriptions and investigation, similar to the *S. undulatus* complex, as these previous descriptions can be confounded by new phylogenies. The results presented in this thesis illustrate the broad heuristic value of investigating sexual dimorphism and morphological patterns in the context of new molecular phylogenies.

In consideration of sexually dimorphic signals, the results presented in this thesis indicate that signal badges may not serve perfectly parallel functions in all species of the *S. undulatus* complex, particularly for *S. cowlesi* and *S. tristichus*. It has been shown empirically that sexual confusion results from overlap in signal patch size in males and females for a population of *S. cowlesi* occurring at White Sands, New Mexico (Robertson and Rosenblum 2010). However, my results suggest that sexual confusion may be common among *S. cowlesi*, and perhaps even more common among *S. tristichus*. I cannot disagree with Robertson and Rosenblum's (2010) observation of sexual confusion, but my data suggests that other explanations are possible. My data clearly shows that signal badges occur at a high frequency in female *S. cowlesi* and *S. tristichus*. Furthermore the variation in badge size between males and females of *S. cowlesi* and *S. tristichus* occurs in areas that are not novel habitats. Because signal patches occur on the belly and throat (i.e. inconspicuous areas) of *S. undulatus* complex, they are resistant to the forces of natural selection via predation. So the function of signal patches is best suited for investigations of how the signal patches are used in same sex and opposite sex interactions. Females that use similar signals as males may also be engaging in territorial disputes with conspecific females, perhaps to defend the best resources for their

offspring. The possibility that female signal badges function as a rejection signal to resist copulation with males while they are gravid is not supported as there is no evidence that female signal badges vary in size or color intensity with reproductive condition. It should be the priority of future field studies on *S. undulatus* complex to investigate female-female interactions and female-male interactions from the perspective of the female. In this way it will be possible to understand the function of signal patches in females.

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## APPENDIX A

Herein I have provided the locality data and voucher numbers for specimens used in the investigation described in chapter I. We used HerpNet (<http://www.herpNet.org>) to query and choose specimens. Specimens were obtained from the following institutions:

University of Kansas Natural History Museum and Biodiversity Research Center

(KUNHM), Smithsonian National Museum of Natural History, Washington, D.C. (USNM),

Museum of Southwestern Biology, University of New Mexico (MSB), Florida Museum of

Natural History, University of Florida (UF), Monte L. Bean Museum, Brigham Young

University, Provo, UT (BYU). All specimens were sent to the Texas Cooperative Wildlife

Collection, Texas A&M University (TCWC) where we complimented our project with

specimens from the TCWC.

*Sceloporus undulatus* (complex)

(30) Kansas: Cherokee, Co. (KUNHM: 7817, 2321128771-72, 28775, 28776, 28781, 28784-85, 28788-89, 30084-85, 30093-95, 30097-103, 31092, 75150, 154017, 157817 159826 206700 289052)

(30) Virginia: Henrico, Co. (USNM: 324245, 325231, 325233-44, 325247-48, 325251-52, 325254-56, 325258, 325260-67)

(30) North Carolina: Lenoir, Co. (USNM: 499937-67, 499972)

(30) Colorado: Garfield, Co. (MSB: 85934, 85936, 85938-39, 85941, 85946, 85949, 85956, 85957, 85976, 85985-86, 85987-88, 85991, 85997-99, 86001, 86045, 86047-48, 86048, 86051, 86063, 86066-67, 86069, 86074, 86076, 86084)

(25) New Mexico: Hidalgo, Co. (MSB: 4427-28, 4449, 6168, 7292, 11129, 11153, 12141, 23248, 33813, 41197-98, 48822- 23, 48850, 51156, 51160 51161-63, 51165, 51168-69, 55903, 55915)

(30) New Mexico: Torrance Co. (MSB: 81-82, 7254, 10274, 11550, 18859, 18865-67, 21187-89, 21201-5, 23399, 23525, 26148-50, 26153-54, 26327, 31353, 31915, 42045, 52795, 87871)

(29) Arizona: Coconino, Co. (MSB: 17282-85, 17287-89, 17292-95, 17304-05, 17308-09, 23921-22, 23940, 23957, 23959, 23971, 23997, 35151, 35154-56, 35159-61)

(27) Florida: Alachua, Co. (UF: 461, 1144, 1978, 7790, 14589-90, 14592, 25191, 54449, 54452-53, 54459, 54461, 83731, 83734, 125193, 125258, 125291-92, 146034, 1389-4, 1389-5, 1389-6, 14593-1, 14593-2, 5942-2, 7553-2)

(30) Utah: Emery, Co. (BYU: 2392, 2806, 16296, 16300, 16407, 16464, 16539, 16765, 18032, 18034, 18355-61, 18363-64, 19026, 19054, 19056, 19057, 19096, 19107-08, 19121-22, 19124-25)

(9) Texas: Anderson, Co. (TCWC: 5783, 5784, 5785, 14018, 29246-47, 39132, 54052, 82758)

(5) Texas: Jasper, Co. (TCWC: 18840, 65189, 67838, 79174, 79175)

(30) Texas: Jeff Davis, Co. (TCWC: 13059, 25290, 25918-20, 25922, 25924-27, 35343-47, 36871, 75770, 75810, 75815, 81549, 81556-57, 81584 81656-59, 83169, 83175, 84525)

(4) Texas: Newton, Co. (TCWC: 8855, 8856, 48386, 48387)

(10) Texas: Polk, Co. (TCWC: 125, 5754, 5756-57, 5763-66, 5769, 46511)

(3) Texas: Tyler, Co. (TCWC: 5770, 8849, 14997)



(30) Texas: Uvalde, Co. (TCWC: 48920, 48935, 48951, 48966, 48974, 48979,  
48981, 48986, 48988, 48992, 48995, 48999, 49002, 49008, 49010-11, 49013, 49015, 49017,  
49020, 51202, 51208, 51210-13, 51216, 51218-20)

## APPENDIX B

The full statistical output of reduced major axes (RMA) regression (mean±SD, slope, confidence intervals,  $R^2$ , and allometry).

Full results of the RMA conducted on SVL vs. Torso

| Parameter                     | Mean±SD   | Slope  | CI            | $R^2$ | Allometry |
|-------------------------------|-----------|--------|---------------|-------|-----------|
| All Males                     | 3.36±0.14 | 0.9182 | 0.829-1.008   | 0.585 | Isometric |
| <i>Sceloporus consobrinus</i> | 3.30±0.15 | 0.5792 | 0.4296-0.7288 | 0.298 | Negative  |
| Kansas                        | 3.22±0.15 | 0.9675 | 0.594-1.341   | 0.585 | Isometric |
| Uvalde, TX                    | 3.47±0.09 | 0.7889 | 0.502-1.076   | 0.63  | Isometric |
| East Texas                    | 3.21±0.08 | 1.086  | 0.617-1.555   | 0.481 | Isometric |
| <i>Sceloporus undulatus</i>   | 3.37±0.13 | 0.9313 | 0.777-1.086   | 0.73  | Isometric |
| Virginia                      | 3.45±0.14 | 0.7887 | 0.5828-0.9946 | 0.81  | Negative  |
| North Carolina                | 3.33±0.10 | 1.266  | 0.783-1.749   | 0.595 | Isometric |
| Florida                       | 3.33±0.11 | 0.8441 | 0.544-1.144   | 0.745 | Isometric |
| <i>Sceloporus tristichus</i>  | 3.45±0.14 | 0.7902 | 0.6505-0.9300 | 0.686 | Negative  |
| Colorado                      | 3.37±0.08 | 1.06   | 0.597-1.523   | 0.469 | Isometric |
| Utah                          | 3.38±0.10 | 0.7773 | 0.5766-0.9780 | 0.814 | Negative  |
| Arizona                       | 3.59±0.13 | 0.9463 | 0.505-1.387   | 0.506 | Isometric |
| <i>Sceloporus cowlesi</i>     | 3.31±0.09 | 0.9966 | 0.794-1.199   | 0.584 | Isometric |
| New Mexico, Hidalgo           | 3.33±0.09 | 0.9151 | 0.558-1.272   | 0.655 | Isometric |
| New Mexico, Torrence          | 3.33±0.08 | 0.8052 | 0.422-1.188   | 0.37  | Isometric |
| Jeff Davis, TX                | 3.28±0.10 | 1.001  | 0.659-1.514   | 0.675 | Isometric |
| All Females                   | 3.44±0.17 | 0.9563 | 0.886-1.026   | 0.758 | Isometric |
| <i>Sceloporus consobrinus</i> | 3.33±0.17 | 0.8765 | 0.696-1.057   | 0.542 | Isometric |
| Kansas                        | 3.37±0.16 | 0.993  | 0.731-1.254   | 0.807 | Isometric |
| Uvalde, TX                    | 3.41±0.16 | 0.7554 | 0.5664-0.9443 | 0.826 | Negative  |
| East Texas                    | 3.20±0.12 | 0.9009 | 0.684-1.118   | 0.823 | Isometric |
| <i>Sceloporus undulatus</i>   | 3.45±0.17 | 0.9042 | 0.790-1.018   | 0.833 | Isometric |
| Virginia                      | 3.55±0.18 | 0.9677 | 0.799-1.136   | 0.916 | Isometric |
| North Carolina                | 3.42±0.13 | 1.04   | 0.801-1.279   | 0.853 | Isometric |
| Florida                       | 3.39±0.15 | 0.729  | 0.5166-0.9415 | 0.763 | Negative  |
| <i>Sceloporus tristichus</i>  | 3.54±0.14 | 0.8667 | 0.727-1.006   | 0.72  | Isometric |
| Colorado                      | 3.52±0.11 | 0.5609 | 0.3749-0.7468 | 0.694 | Negative  |
| Utah                          | 3.50±0.10 | 0.7379 | 0.4876-0.9882 | 0.679 | Negative  |
| Arizona                       | 3.59±0.18 | 1.019  | 0.808-1.231   | 0.869 | Isometric |
| <i>Sceloporus cowlesi</i>     | 3.45±0.14 | 0.8828 | 0.748-1.018   | 0.772 | Isometric |

|                      |           |        |               |       |           |
|----------------------|-----------|--------|---------------|-------|-----------|
| New Mexico, Hidalgo  | 3.40±0.14 | 0.8239 | 0.617-1.031   | 0.873 | Isometric |
| New Mexico, Torrence | 3.51±0.15 | 0.6384 | 0.4431-0.8336 | 0.74  | Negative  |
| Jeff Davis, TX       | 3.43±0.10 | 0.9561 | 0.760-1.153   | 0.882 | Isometric |

Full results of the RMA run on SVL vs Head Depth

| Parameter                     | Mean±SD   | Slope  | CI            | R <sup>2</sup> | Allometry |
|-------------------------------|-----------|--------|---------------|----------------|-----------|
| All Males                     | 1.95±0.14 | 0.9052 | 0.8289-0.9815 | 0.689          | Negative  |
| <i>Sceloporus consobrinus</i> | 1.88±0.14 | 0.6384 | 0.5266-0.7502 | 0.677          | Negative  |
| Kansas                        | 1.91±0.15 | 0.747  | 0.6348-0.8593 | 0.937          | Negative  |
| Uvalde, TX                    | 1.79±0.14 | 0.942  | 0.505-1.379   | 0.402          | Isometric |
| East Texas                    | 1.95±0.08 | 0.6056 | 0.4375-0.7736 | 0.785          | Negative  |
| <i>Sceloporus undulatus</i>   | 2.00±0.15 | 0.8061 | 0.6930-0.9192 | 0.807          | Negative  |
| Virginia                      | 2.09±0.13 | 0.8536 | 0.668-1.040   | 0.868          | Isometric |
| North Carolina                | 1.92±0.15 | 0.8566 | 0.614-1.099   | 0.776          | Isometric |
| Florida                       | 1.99±0.12 | 0.7509 | 0.420-1.082   | 0.608          | Isometric |
| <i>Sceloporus tristichus</i>  | 2.01±0.12 | 0.9537 | 0.780-1.127   | 0.668          | Isometric |
| Colorado                      | 1.96±0.11 | 0.7894 | 0.5840-0.9948 | 0.811          | Negative  |
| Utah                          | 2.01±0.09 | 0.801  | 0.502-1.100   | 0.613          | Isometric |
| Arizona                       | 2.08±0.12 | 0.9877 | 0.581-1.394   | 0.615          | Isometric |
| <i>Sceloporus cowlesi</i>     | 1.92±0.13 | 0.6933 | 0.5914-0.7952 | 0.783          | Negative  |
| New Mexico, Hidalgo           | 1.95±0.10 | 0.8867 | 0.672-1.102   | 0.866          | Isometric |
| New Mexico, Torrence          | 1.99±0.11 | 0.6149 | 0.3836-0.8462 | 0.606          | Negative  |
| Jeff Davis, TX                | 1.82±0.13 | 0.758  | 0.5480-0.9681 | 0.786          | Negative  |
| All Females                   | 1.98±0.16 | 1.036  | 0.960-1.113   | 0.752          | Isometric |
| <i>Sceloporus consobrinus</i> | 1.91±0.16 | 0.9261 | 0.834-1.018   | 0.893          | Isometric |
| Kansas                        | 2.04±0.17 | 0.932  | 0.743-1.121   | 0.886          | Isometric |
| Uvalde, TX                    | 1.87±0.10 | 1.207  | 0.874-1.541   | 0.788          | Isometric |
| East Texas                    | 1.84±0.14 | 0.8189 | 0.6846-0.9532 | 0.918          | Negative  |
| <i>Sceloporus undulatus</i>   | 2.02±0.16 | 0.9159 | 0.779-1.053   | 0.765          | Isometric |
| Virginia                      | 2.12±0.16 | 1.076  | 0.892-1.260   | 0.919          | Isometric |
| North Carolina                | 2.02±0.12 | 1.16   | 0.778-1.541   | 0.699          | Isometric |
| Florida                       | 1.90±0.13 | 0.8064 | 0.487-1.125   | 0.564          | Isometric |
| <i>Sceloporus tristichus</i>  | 2.05±0.13 | 0.9327 | 0.768-1.097   | 0.664          | Isometric |
| Colorado                      | 2.06±0.08 | 0.7846 | 0.509-1.060   | 0.656          | Isometric |
| Utah                          | 2.09±0.06 | 1.13   | 0.567-1.692   | 0.308          | Isometric |
| Arizona                       | 2.01±0.19 | 0.9754 | 0.723-1.228   | 0.796          | Isometric |
| <i>Sceloporus cowlesi</i>     | 1.95±0.14 | 0.8827 | 0.746-1.019   | 0.766          | Isometric |
| New Mexico, Hidalgo           | 1.91±0.13 | 0.926  | 0.598-1.254   | 0.747          | Isometric |

|                      |           |        |             |       |           |
|----------------------|-----------|--------|-------------|-------|-----------|
| New Mexico, Torrence | 2.04±0.12 | 0.8359 | 0.614-1.058 | 0.804 | Isometric |
| Jeff Davis, TX       | 1.86±0.11 | 0.8969 | 0.521-1.273 | 0.51  | Isometric |

Full results of the RMA run on SVL vs Rear Leg Length

| Parameter                     | Mean±SD   | Slope  | CI            | R <sup>2</sup> | Allometry |
|-------------------------------|-----------|--------|---------------|----------------|-----------|
| All Males                     | 3.61±0.15 | 0.9042 | 0.8372-0.9712 | 0.759          | Negative  |
| <i>Sceloporus consobrinus</i> | 3.48±0.13 | 0.8473 | 0.659-1.035   | 0.481          | Isometric |
| Kansas                        | 3.55±0.14 | 0.9283 | 0.635-1.222   | 0.721          | Isometric |
| Uvalde, TX                    | 3.47±0.12 | 0.8377 | 0.408-1.267   | 0.267          | Isometric |
| East Texas                    | 3.41±0.11 | 0.8266 | 0.501-1.153   | 0.567          | Isometric |
| <i>Sceloporus undulatus</i>   | 3.65±0.14 | 1.047  | 0.930-1.164   | 0.877          | Isometric |
| Virginia                      | 3.68±0.16 | 1.121  | 0.851-1.391   | 0.838          | Isometric |
| North Carolina                | 3.68±0.13 | 1.138  | 0.977-1.300   | 0.944          | Isometric |
| Florida                       | 3.60±0.13 | 0.9429 | 0.601-1.285   | 0.735          | Isometric |
| <i>Sceloporus tristichus</i>  | 3.72±0.13 | 0.9839 | 0.808-1.159   | 0.68           | Isometric |
| Colorado                      | 3.70±0.07 | 1.368  | 0.871-1.864   | 0.633          | Isometric |
| Utah                          | 3.75±0.12 | 0.8149 | 0.388-1.242   | 0.236          | Isometric |
| Arizona                       | 3.70±0.18 | 0.9004 | 0.643-1.158   | 0.815          | Isometric |
| <i>Sceloporus cowlesi</i>     | 3.60±0.09 | 1.184  | 0.916-1.453   | 0.483          | Isometric |
| New Mexico, Hidalgo           | 3.56±0.10 | 1.026  | 0.557-1.495   | 0.526          | Isometric |
| New Mexico, Torrence          | 3.64±0.06 | 0.967  | 0.474-1.460   | 0.276          | Isometric |
| Jeff Davis, TX                | 3.58±0.09 | 1.366  | 0.951-1.781   | 0.743          | Isometric |
| All Females                   | 3.60±0.15 | 1.079  | 1.010-1.148   | 0.815          | Positive  |
| <i>Sceloporus consobrinus</i> | 3.48±0.13 | 1.149  | 0.966-1.333   | 0.725          | Isometric |
| Kansas                        | 3.42±0.11 | 1.163  | 0.867-1.459   | 0.819          | Isometric |
| Uvalde, TX                    | 3.58±0.14 | 1.069  | 0.671-1.467   | 0.614          | Isometric |
| East Texas                    | 3.45±0.11 | 0.9715 | 0.638-1.305   | 0.641          | Isometric |
| <i>Sceloporus undulatus</i>   | 3.66±0.13 | 1.108  | 1.010-1.206   | 0.918          | Positive  |
| Virginia                      | 3.66±0.17 | 1.121  | 0.851-1.391   | 0.838          | Isometric |
| North Carolina                | 3.67±0.12 | 1.062  | 0.886-1.237   | 0.924          | Isometric |
| Florida                       | 3.64±0.09 | 0.9898 | 0.733-1.246   | 0.813          | Isometric |
| <i>Sceloporus tristichus</i>  | 3.72±0.11 | 0.9962 | 0.811-1.181   | 0.628          | Isometric |
| Colorado                      | 3.67±0.07 | 1.324  | 0.559-2.089   | 0.0701         | Isometric |
| Utah                          | 3.70±0.09 | 1.373  | 0.634-2.112   | 0.193          | Isometric |
| Arizona                       | 3.78±0.14 | 0.9812 | 0.754-1.209   | 0.836          | Isometric |
| <i>Sceloporus cowlesi</i>     | 3.58±0.08 | 1.406  | 1.168-1.644   | 0.72           | Positive  |
| New Mexico, Hidalgo           | 3.59±0.10 | 1.247  | 0.789-1.705   | 0.728          | Isometric |
| New Mexico, Torrence          | 3.58±0.07 | 1.205  | 0.858-1.553   | 0.769          | Isometric |

|                |           |       |             |       |           |
|----------------|-----------|-------|-------------|-------|-----------|
| Jeff Davis, TX | 3.56±0.08 | 1.275 | 0.856-1.694 | 0.699 | Isometric |
|----------------|-----------|-------|-------------|-------|-----------|

Full results of the RMA run on SVL vs. Head Length

| Parameter                     | Mean±SD   | Slope  | CI            | R <sup>2</sup> | Allometry |
|-------------------------------|-----------|--------|---------------|----------------|-----------|
| All Males                     | 2.69±0.13 | 1.028  | 0.968-1.088   | 0.851          | Isometric |
| <i>Sceloporus consobrinus</i> | 2.61±0.10 | 0.8601 | 0.7582-0.9626 | 0.79           | Negative  |
| Kansas                        | 2.59±0.13 | 0.868  | 0.7557-0.9919 | 0.95           | Negative  |
| Uvalde, TX                    | 2.58±0.10 | 0.8746 | 0.690-1.041   | 0.867          | Isometric |
| East Texas                    | 2.65±0.08 | 0.9671 | 0.673-1.828   | 0.45           | Isometric |
| <i>Sceloporus undulatus</i>   | 2.72±0.12 | 1.035  | 0.910-1.161   | 0.856          | Isometric |
| Virginia                      | 2.77±0.10 | 1.072  | 0.872-1.195   | 0.922          | Isometric |
| North Carolina                | 2.67±0.14 | 0.9232 | 0.665-1.313   | 0.783          | Isometric |
| Florida                       | 2.71±0.08 | 1.096  | 0.862-1.330   | 0.908          | Isometric |
| <i>Sceloporus tristichus</i>  | 2.80±0.11 | 0.9599 | 0.852-1.082   | 0.875          | Isometric |
| Colorado                      | 2.76±0.09 | 0.9409 | 0.732-1.179   | 0.787          | Isometric |
| Utah                          | 2.75±0.09 | 0.8088 | 0.563-1.297   | 0.707          | Isometric |
| Arizona                       | 2.89±0.11 | 1.08   | 0.761-1.203   | 0.939          | Isometric |
| <i>Sceloporus cowlesi</i>     | 2.64±0.09 | 1.028  | 0.848-1.182   | 0.666          | Isometric |
| New Mexico, Hidalgo           | 2.61±0.09 | 0.9137 | 0.713-1.123   | 0.618          | Isometric |
| New Mexico, Torrence          | 2.69±0.06 | 1.069  | 0.728-1.550   | 0.526          | Isometric |
| Jeff Davis, TX                | 2.62±0.09 | 1.029  | 0.735-1.200   | 0.869          | Isometric |
| All Females                   | 2.72±0.14 | 1.035  | 0.930-1.153   | 0.751          | Isometric |
| <i>Sceloporus consobrinus</i> | 2.63±0.12 | 1.206  | 1.106-1.333   | 0.865          | Positive  |
| Kansas                        | 2.70±0.12 | 1.194  | 1.014-1.464   | 0.834          | Positive  |
| Uvalde, TX                    | 2.61±0.10 | 1.194  | 1.013-1.522   | 0.834          | Positive  |
| East Texas                    | 2.58±0.11 | 0.9843 | 0.830-1.121   | 0.863          | Isometric |
| <i>Sceloporus undulatus</i>   | 2.73±0.12 | 1.213  | 1.112-1.343   | 0.903          | Positive  |
| Virginia                      | 2.77±0.15 | 1.186  | 1.049-1.424   | 0.924          | Positive  |
| North Carolina                | 2.75±0.11 | 1.223  | 1.086-1.494   | 0.934          | Positive  |
| Florida                       | 2.69±0.10 | 1.08   | 0.899-1.317   | 0.822          | Isometric |
| <i>Sceloporus tristichus</i>  | 2.84±0.12 | 0.9932 | 0.895-1.120   | 0.824          | Isometric |
| Colorado                      | 2.85±0.07 | 0.8901 | 0.506-1.269   | 0.444          | Isometric |
| Utah                          | 2.84±0.06 | 1.107  | 0.677-1.768   | 0.394          | Isometric |
| Arizona                       | 2.82±0.19 | 0.9961 | 0.860-1.069   | 0.946          | Isometric |
| <i>Sceloporus cowlesi</i>     | 2.66±0.10 | 1.204  | 1.055-1.415   | 0.762          | Positive  |
| New Mexico, Hidalgo           | 2.59±0.07 | 1.679  | 1.166-2.452   | 0.805          | Positive  |
| New Mexico, Torrence          | 2.75±0.09 | 1.036  | 0.721-1.333   | 0.851          | Isometric |
| Jeff Davis, TX                | 2.63±0.06 | 1.503  | 1.103-1.919   | 0.44           | Positive  |

## Full Results of the RMA run on SVL vs Head Width

| Parameter                     | Mean±SD   | Slope  | CI            | R <sup>2</sup> | Allometry |
|-------------------------------|-----------|--------|---------------|----------------|-----------|
| All Males                     | 2.25±0.15 | 0.8644 | 0.8046-0.9241 | 0.79           | Negative  |
| <i>Sceloporus consobrinus</i> | 2.17±0.15 | 0.5923 | 0.4751-0.7096 | 0.588          | Negative  |
| Kansas                        | 2.19±0.14 | 0.7993 | 0.6355-0.9632 | 0.883          | Negative  |
| Uvalde, TX                    | 2.06±0.13 | 0.7433 | 0.390-1.097   | 0.37           | Isometric |
| East Texas                    | 2.30±0.10 | 0.6199 | 0.4637-0.7913 | 0.823          | Negative  |
| <i>Sceloporus undulatus</i>   | 2.29±0.13 | 0.8987 | 0.7976-0.9999 | 0.876          | Negative  |
| Virginia                      | 2.38±0.12 | 0.901  | 0.750-1.052   | 0.922          | Isometric |
| North Carolina                | 2.21±0.13 | 1.017  | 0.774-1.261   | 0.841          | Isometric |
| Florida                       | 2.28±0.10 | 0.9524 | 0.669-1.236   | 0.822          | Isometric |
| <i>Sceloporus tristichus</i>  | 2.40±0.12 | 0.928  | 0.838-1.018   | 0.906          | Isometric |
| Colorado                      | 2.32±0.10 | 0.8765 | 0.682-1.071   | 0.862          | Isometric |
| Utah                          | 2.34±0.09 | 0.8244 | 0.602-1.047   | 0.797          | Isometric |
| Arizona                       | 2.46±0.13 | 0.9782 | 0.852-1.105   | 0.962          | Isometric |
| <i>Sceloporus cowlesi</i>     | 2.21±0.12 | 0.7834 | 0.6665-0.9002 | 0.777          | Negative  |
| New Mexico, Hidalgo           | 2.21±0.11 | 0.7925 | 0.525-1.060   | 0.74           | Isometric |
| New Mexico, Torrence          | 2.28±0.08 | 0.8302 | 0.502-1.158   | 0.565          | Isometric |
| Jeff Davis, TX                | 2.12±0.10 | 0.9259 | 0.693-1.158   | 0.824          | Isometric |
| All Females                   | 2.28±0.17 | 1.015  | 0.943-1.086   | 0.773          | Isometric |
| <i>Sceloporus consobrinus</i> | 2.19±0.16 | 1.046  | 0.874-1.217   | 0.771          | Isometric |
| Kansas                        | 2.32±0.15 | 1.085  | 0.865-1.305   | 0.885          | Isometric |
| Uvalde, TX                    | 2.16±0.11 | 1.441  | 0.719-2.164   | 0.3            | Isometric |
| East Texas                    | 2.10±0.12 | 0.8988 | 0.716-1.082   | 0.874          | Isometric |
| <i>Sceloporus undulatus</i>   | 2.31±0.16 | 0.9565 | 0.814-1.099   | 0.765          | Isometric |
| Virginia                      | 2.43±0.13 | 1.304  | 1.040-1.569   | 0.886          | Positive  |
| North Carolina                | 2.33±0.13 | 1.086  | 0.843-1.329   | 0.86           | Isometric |
| Florida                       | 2.19±0.11 | 0.9702 | 0.597-1.344   | 0.587          | Isometric |
| <i>Sceloporus tristichus</i>  | 2.40±0.12 | 0.97   | 0.835-1.104   | 0.793          | Isometric |
| Colorado                      | 2.43±0.07 | 0.8482 | 0.481-1.215   | 0.479          | Isometric |
| Utah                          | 2.4±0.068 | 1.057  | 0.548-1.567   | 0.354          | Isometric |
| Arizona                       | 2.38±0.19 | 0.9962 | 0.849-1.143   | 0.934          | Isometric |
| <i>Sceloporus cowlesi</i>     | 2.24±0.15 | 0.8964 | 0.785-1.008   | 0.849          | Isometric |
| New Mexico, Hidalgo           | 2.17±0.13 | 0.9052 | 0.571-1.239   | 0.726          | Isometric |
| New Mexico, Torrence          | 2.31±0.12 | 0.8063 | 0.6531-0.9595 | 0.899          | Isometric |
| Jeff Davis, TX                | 2.14±0.10 | 0.9906 | 0.700-1.282   | 0.76           | Isometric |

## APPENDIX C

Herein I provide locality data, voucher numbers and raw measurements for specimens used in chapter III of this investigation. I used HerpNet (<http://www.herpNet.org>) to query and choose specimens. Specimens were obtained from the Museum of Southwestern Biology, University of New Mexico (MSB), and the Texas Cooperative Wildlife Collection, Texas A&M University (TCWC). Herein I provide two tables for each sex. Each table includes the state and county of collection, the collection identification number, the raw untransformed Total Badge Area (TBA) and raw untransformed snout to vent length (SVL).

**Collections Data and Raw Measurements for all Female Specimens**

| <b>Collection ID</b> | <b>State</b> | <b>County</b> | <b>Species</b>        | <b>TBA</b> | <b>SVL</b> |
|----------------------|--------------|---------------|-----------------------|------------|------------|
| MSB 56565            | NM           | Lea           | <i>S. consobrinus</i> | 0          | 61.8271    |
| MSB 56778            | NM           | Lea           | <i>S. consobrinus</i> | 0          | 62.3017    |
| MSB 59874            | NM           | Lea           | <i>S. consobrinus</i> | 0          | 63.3444    |
| MSB 60255            | NM           | Lea           | <i>S. consobrinus</i> | 0          | 56.1982    |
| MSB 60417            | NM           | Lea           | <i>S. consobrinus</i> | 0          | 49.8486    |
| MSB 60509            | NM           | Lea           | <i>S. consobrinus</i> | 0          | 49.8486    |
| MSB 60531            | NM           | Lea           | <i>S. consobrinus</i> | 0          | 48.9275    |
| MSB 60533            | NM           | Lea           | <i>S. consobrinus</i> | 0          | 58.8965    |
| MSB 60929            | NM           | Lea           | <i>S. consobrinus</i> | 0          | 55.0320    |
| MSB 57635            | NM           | Roosevelt     | <i>S. consobrinus</i> | 0          | 51.8152    |
| MSB 59893            | NM           | Roosevelt     | <i>S. consobrinus</i> | 0          | 59.9459    |
| MSB 59894            | NM           | Roosevelt     | <i>S. consobrinus</i> | 0          | 54.8725    |
| MSB 59924            | NM           | Roosevelt     | <i>S. consobrinus</i> | 0          | 52.3520    |
| MSB 59927            | NM           | Roosevelt     | <i>S. consobrinus</i> | 0          | 50.4669    |
| MSB 59928            | NM           | Roosevelt     | <i>S. consobrinus</i> | 0          | 55.5981    |
| MSB 59936            | NM           | Roosevelt     | <i>S. consobrinus</i> | 0          | 56.2699    |
| MSB 59940            | NM           | Roosevelt     | <i>S. consobrinus</i> | 0          | 64.4547    |
| TCWC 48966           | TX           | Uvalde        | <i>S. consobrinus</i> | 0          | 49.6148    |
| TCWC 48969           | TX           | Uvalde        | <i>S. consobrinus</i> | 0          | 49.6107    |
| TCWC 79167           | TX           | Bandera       | <i>S. consobrinus</i> | 0          | 70.0019    |

|      |       |    |          |                       |         |         |
|------|-------|----|----------|-----------------------|---------|---------|
| TCWC | 80431 | TX | Kerr     | <i>S. consobrinus</i> | 27.1741 | 72.2300 |
| TCWC | 48920 | TX | Uvalde   | <i>S. consobrinus</i> | 18.4908 | 61.3404 |
| TCWC | 48937 | TX | Uvalde   | <i>S. consobrinus</i> | 49.5254 | 64.6283 |
| TCWC | 48998 | TX | Uvalde   | <i>S. consobrinus</i> | 21.5026 | 56.9294 |
| MSB  | 22906 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 65.2921 |
| MSB  | 22954 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 69.7794 |
| MSB  | 26158 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 78.8293 |
| MSB  | 31158 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 68.1793 |
| MSB  | 31160 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 73.1154 |
| MSB  | 33641 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 68.9705 |
| MSB  | 38476 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 57.2676 |
| MSB  | 38497 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 59.2604 |
| MSB  | 38505 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 61.2265 |
| MSB  | 43648 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 69.2264 |
| MSB  | 46054 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 70.1751 |
| MSB  | 48589 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 73.6866 |
| MSB  | 48590 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 69.1457 |
| MSB  | 61900 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 60.3801 |
| MSB  | 71562 | NM | Eddy     | <i>S. consobrinus</i> | 0       | 73.8084 |
| TCWC | 46514 | TX | Kinney   | <i>S. consobrinus</i> | 0       | 64.2139 |
| TCWC | 46515 | TX | Kinney   | <i>S. consobrinus</i> | 0       | 54.2078 |
| TCWC | 54071 | TX | Kinney   | <i>S. consobrinus</i> | 26.1110 | 62.5240 |
| MSB  | 60365 | NM | Otero    | <i>S. cowlesi</i>     | 0       | 61.9898 |
| MSB  | 23399 | NM | Torrance | <i>S. cowlesi</i>     | 0       | 65.4595 |
| MSB  | 26046 | NM | Valencia | <i>S. cowlesi</i>     | 0       | 69.3213 |
| MSB  | 21113 | NM | Torrance | <i>S. cowlesi</i>     | 56.3167 | 56.9441 |
| MSB  | 11550 | NM | Torrance | <i>S. cowlesi</i>     | 52.8420 | 69.7025 |
| MSB  | 21105 | NM | Torrance | <i>S. cowlesi</i>     | 69.1051 | 73.0519 |
| MSB  | 21117 | NM | Torrance | <i>S. cowlesi</i>     | 36.0364 | 69.4939 |
| MSB  | 26148 | NM | Torrance | <i>S. cowlesi</i>     | 35.5245 | 71.2026 |
| MSB  | 6335  | NM | Otero    | <i>S. cowlesi</i>     | 37.8542 | 64.5813 |
| MSB  | 16282 | NM | Otero    | <i>S. cowlesi</i>     | 26.0071 | 55.7215 |
| MSB  | 48031 | NM | Otero    | <i>S. cowlesi</i>     | 35.6530 | 69.2035 |
| MSB  | 60366 | NM | Otero    | <i>S. cowlesi</i>     | 30.9491 | 60.5681 |
| MSB  | 60367 | NM | Otero    | <i>S. cowlesi</i>     | 30.6146 | 69.5198 |
| MSB  | 60368 | NM | Otero    | <i>S. cowlesi</i>     | 23.5764 | 64.1262 |
| MSB  | 60369 | NM | Otero    | <i>S. cowlesi</i>     | 26.0451 | 72.2332 |
| MSB  | 26150 | NM | Torrance | <i>S. cowlesi</i>     | 39.5927 | 78.1202 |
| MSB  | 21114 | NM | Torrance | <i>S. cowlesi</i>     | 56.7220 | 66.9330 |
| MSB  | 21123 | NM | Torrance | <i>S. cowlesi</i>     | 37.2090 | 59.7812 |



|      |       |    |            |                   |          |         |
|------|-------|----|------------|-------------------|----------|---------|
| MSB  | 21203 | NM | Torrance   | <i>S. cowlesi</i> | 37.8698  | 70.2739 |
| MSB  | 7249  | NM | Valencia   | <i>S. cowlesi</i> | 28.3683  | 65.9678 |
| MSB  | 7251  | NM | Valencia   | <i>S. cowlesi</i> | 13.6073  | 62.8052 |
| MSB  | 21115 | NM | Torrance   | <i>S. cowlesi</i> | 499.5412 | 72.4474 |
| MSB  | 26149 | NM | Torrance   | <i>S. cowlesi</i> | 297.2064 | 68.9566 |
| MSB  | 6334  | NM | Otero      | <i>S. cowlesi</i> | 227.3149 | 68.6022 |
| MSB  | 18865 | NM | Torrance   | <i>S. cowlesi</i> | 244.7470 | 61.0278 |
| MSB  | 21108 | NM | Torrance   | <i>S. cowlesi</i> | 160.9683 | 50.9515 |
| MSB  | 26152 | NM | Torrance   | <i>S. cowlesi</i> | 328.1760 | 65.7085 |
| MSB  | 26327 | NM | Torrance   | <i>S. cowlesi</i> | 311.8219 | 69.4237 |
| MSB  | 26154 | NM | Torrance   | <i>S. cowlesi</i> | 368.5055 | 70.2960 |
| MSB  | 6814  | NM | Valencia   | <i>S. cowlesi</i> | 207.6157 | 57.9157 |
| MSB  | 7092  | NM | Valencia   | <i>S. cowlesi</i> | 262.2492 | 65.7343 |
| MSB  | 21199 | NM | Valencia   | <i>S. cowlesi</i> | 282.4295 | 66.9102 |
| MSB  | 21198 | NM | Valencia   | <i>S. cowlesi</i> | 297.6377 | 64.4957 |
| MSB  | 21191 | NM | Valencia   | <i>S. cowlesi</i> | 181.5809 | 60.7111 |
| TCWC | 51988 | TX | Terrell    | <i>S. cowlesi</i> | 0        | 69.3129 |
| TCWC | 72507 | TX | Terrell    | <i>S. cowlesi</i> | 0        | 59.2620 |
| TCWC | 60553 | TX | Val Verde  | <i>S. cowlesi</i> | 0        | 58.8965 |
| TCWC | 60554 | TX | Val Verde  | <i>S. cowlesi</i> | 0        | 59.3544 |
| TCWC | 64702 | TX | Val Verde  | <i>S. cowlesi</i> | 0        | 61.7394 |
| TCWC | 52205 | TX | Edwards    | <i>S. cowlesi</i> | 0        | 64.9511 |
| TCWC | 67772 | TX | Edwards    | <i>S. cowlesi</i> | 0        | 62.6272 |
| TCWC | 7197  | TX | Edwards    | <i>S. cowlesi</i> | 0        | 61.6159 |
| TCWC | 16015 | TX | Brewster   | <i>S. cowlesi</i> | 0        | 62.6417 |
| TCWC | 16017 | TX | Brewster   | <i>S. cowlesi</i> | 0        | 61.3761 |
| TCWC | 16018 | TX | Brewster   | <i>S. cowlesi</i> | 0        | 54.6724 |
| TCWC | 16020 | TX | Brewster   | <i>S. cowlesi</i> | 0        | 61.7134 |
| TCWC | 16026 | TX | Brewster   | <i>S. cowlesi</i> | 0        | 57.9907 |
| TCWC | 25918 | TX | Jeff Davis | <i>S. cowlesi</i> | 0        | 68.9094 |
| TCWC | 25922 | TX | Jeff Davis | <i>S. cowlesi</i> | 0        | 68.8879 |
| TCWC | 25924 | TX | Jeff Davis | <i>S. cowlesi</i> | 0        | 56.8099 |
| TCWC | 25925 | TX | Jeff Davis | <i>S. cowlesi</i> | 0        | 59.1401 |
| TCWC | 25926 | TX | Jeff Davis | <i>S. cowlesi</i> | 0        | 66.7151 |
| TCWC | 26125 | TX | Jeff Davis | <i>S. cowlesi</i> | 0        | 62.6508 |
| TCWC | 35347 | TX | Jeff Davis | <i>S. cowlesi</i> | 0        | 60.7689 |
| TCWC | 71586 | TX | Jeff Davis | <i>S. cowlesi</i> | 0        | 63.4584 |
| TCWC | 81556 | TX | Jeff Davis | <i>S. cowlesi</i> | 0        | 67.7033 |
| TCWC | 81584 | TX | Jeff Davis | <i>S. cowlesi</i> | 0        | 56.6808 |
| TCWC | 81656 | TX | Jeff Davis | <i>S. cowlesi</i> | 0        | 59.1131 |

|      |       |    |            |                      |          |         |
|------|-------|----|------------|----------------------|----------|---------|
| TCWC | 84525 | TX | Jeff Davis | <i>S. cowlesi</i>    | 0        | 58.4365 |
| TCWC | 16021 | TX | Brewster   | <i>S. cowlesi</i>    | 16.3478  | 58.7332 |
| TCWC | 52209 | TX | Edwards    | <i>S. cowlesi</i>    | 22.4842  | 69.4231 |
| TCWC | 20719 | TX | Val Verde  | <i>S. cowlesi</i>    | 32.3264  | 63.6716 |
| TCWC | 54496 | TX | Val Verde  | <i>S. cowlesi</i>    | 27.1324  | 67.0899 |
| MSB  | 82489 | CO | Montezuma  | <i>S. tristichus</i> | 393.4121 | 78.3175 |
| MSB  | 82490 | CO | Montezuma  | <i>S. tristichus</i> | 281.2941 | 68.7824 |
| MSB  | 82493 | CO | Montezuma  | <i>S. tristichus</i> | 341.9425 | 75.7534 |
| MSB  | 83014 | CO | Montezuma  | <i>S. tristichus</i> | 272.5066 | 67.3408 |
| MSB  | 83006 | CO | Montezuma  | <i>S. tristichus</i> | 317.2331 | 77.6375 |
| MSB  | 83007 | CO | Montezuma  | <i>S. tristichus</i> | 421.5562 | 75.6908 |
| MSB  | 83006 | CO | Montezuma  | <i>S. tristichus</i> | 317.2331 | 77.6375 |
| MSB  | 83011 | CO | Montezuma  | <i>S. tristichus</i> | 347.0950 | 80.3956 |
| MSB  | 87782 | CO | Montezuma  | <i>S. tristichus</i> | 466.9852 | 78.5728 |
| MSB  | 87882 | CO | Montezuma  | <i>S. tristichus</i> | 268.2195 | 74.9558 |
| MSB  | 14579 | NM | Bernalillo | <i>S. tristichus</i> | 0        | 63.1614 |
| MSB  | 14675 | NM | Bernalillo | <i>S. tristichus</i> | 0        | 61.2351 |
| MSB  | 16890 | NM | Bernalillo | <i>S. tristichus</i> | 0        | 64.2908 |
| MSB  | 17264 | NM | Sandoval   | <i>S. tristichus</i> | 0        | 54.3321 |
| MSB  | 13163 | NM | Bernalillo | <i>S. tristichus</i> | 47.1971  | 77.0294 |
| MSB  | 16574 | NM | Bernalillo | <i>S. tristichus</i> | 17.6055  | 55.9998 |
| MSB  | 16576 | NM | Bernalillo | <i>S. tristichus</i> | 21.2051  | 66.2791 |
| MSB  | 16995 | NM | Bernalillo | <i>S. tristichus</i> | 42.2754  | 70.3302 |
| MSB  | 21190 | NM | Bernalillo | <i>S. tristichus</i> | 35.2024  | 72.6971 |
| MSB  | 25974 | NM | Bernalillo | <i>S. tristichus</i> | 26.3806  | 73.6552 |
| MSB  | 16888 | NM | Sandoval   | <i>S. tristichus</i> | 21.4483  | 54.9722 |
| MSB  | 17315 | NM | Sandoval   | <i>S. tristichus</i> | 18.7502  | 60.7277 |
| MSB  | 17316 | NM | Sandoval   | <i>S. tristichus</i> | 20.5248  | 70.8660 |
| MSB  | 76389 | NM | Mickinley  | <i>S. tristichus</i> | 36.5211  | 67.8412 |
| MSB  | 16477 | NM | Bernalillo | <i>S. tristichus</i> | 484.1792 | 74.1342 |
| MSB  | 25837 | NM | Bernalillo | <i>S. tristichus</i> | 274.7938 | 69.1345 |
| MSB  | 15972 | NM | Bernalillo | <i>S. tristichus</i> | 193.4820 | 57.1252 |
| MSB  | 16612 | NM | Bernalillo | <i>S. tristichus</i> | 325.7322 | 70.5585 |
| MSB  | 21208 | NM | Bernalillo | <i>S. tristichus</i> | 360.9930 | 66.2045 |
| MSB  | 23522 | NM | Bernalillo | <i>S. tristichus</i> | 407.4995 | 69.3471 |
| MSB  | 76285 | NM | Cibola     | <i>S. tristichus</i> | 311.7578 | 71.7635 |
| MSB  | 76385 | NM | Cibola     | <i>S. tristichus</i> | 297.8101 | 72.4612 |
| MSB  | 87840 | NM | Cibola     | <i>S. tristichus</i> | 352.3958 | 70.5926 |
| MSB  | 3212  | NM | Mickinley  | <i>S. tristichus</i> | 756.4573 | 95.1929 |
| MSB  | 3217  | NM | Mickinley  | <i>S. tristichus</i> | 476.4050 | 85.8810 |

|     |       |    |           |                      |          |         |
|-----|-------|----|-----------|----------------------|----------|---------|
| MSB | 3218  | NM | Mickinley | <i>S. tristichus</i> | 306.5322 | 75.1365 |
| MSB | 11960 | NM | Mickinley | <i>S. tristichus</i> | 445.1829 | 72.1382 |
| MSB | 76376 | NM | Mickinley | <i>S. tristichus</i> | 163.3097 | 55.6301 |
| MSB | 76387 | NM | Mickinley | <i>S. tristichus</i> | 280.1656 | 68.3112 |
| MSB | 76410 | NM | Mickinley | <i>S. tristichus</i> | 243.8820 | 62.0186 |
| MSB | 77934 | NM | Mickinley | <i>S. tristichus</i> | 377.0175 | 73.0797 |
| MSB | 3181  | NM | San Juan  | <i>S. tristichus</i> | 351.5229 | 77.4198 |
| MSB | 3183  | NM | San Juan  | <i>S. tristichus</i> | 553.4925 | 90.2097 |
| MSB | 3193  | NM | San Juan  | <i>S. tristichus</i> | 294.9486 | 71.3265 |
| MSB | 3194  | NM | San Juan  | <i>S. tristichus</i> | 294.9486 | 71.3265 |
| MSB | 3220  | NM | San Juan  | <i>S. tristichus</i> | 325.1569 | 76.3891 |
| MSB | 21782 | NM | San Juan  | <i>S. tristichus</i> | 387.3275 | 69.9082 |
| MSB | 21785 | NM | San Juan  | <i>S. tristichus</i> | 545.0869 | 74.3095 |
| MSB | 40658 | NM | San Juan  | <i>S. tristichus</i> | 262.7891 | 71.8935 |
| MSB | 40659 | NM | San Juan  | <i>S. tristichus</i> | 288.4467 | 67.9031 |
| MSB | 40668 | NM | San Juan  | <i>S. tristichus</i> | 376.6657 | 79.3108 |
| MSB | 48915 | NM | San Juan  | <i>S. tristichus</i> | 345.5227 | 72.7250 |
| MSB | 48917 | NM | San Juan  | <i>S. tristichus</i> | 369.8052 | 80.5886 |
| MSB | 49987 | NM | San Juan  | <i>S. tristichus</i> | 382.3376 | 72.8808 |
| MSB | 60304 | NM | San Juan  | <i>S. tristichus</i> | 370.3662 | 73.0537 |
| MSB | 63176 | NM | San Juan  | <i>S. tristichus</i> | 306.1003 | 77.3014 |
| MSB | 63177 | NM | San Juan  | <i>S. tristichus</i> | 416.2134 | 77.8749 |
| MSB | 63178 | NM | San Juan  | <i>S. tristichus</i> | 433.9067 | 79.7339 |
| MSB | 16476 | NM | Sandoval  | <i>S. tristichus</i> | 556.0932 | 76.1882 |
| MSB | 17255 | NM | Sandoval  | <i>S. tristichus</i> | 257.3591 | 58.5170 |
| MSB | 26147 | NM | Sandoval  | <i>S. tristichus</i> | 513.6601 | 76.2503 |
| MSB | 73520 | NM | Sandoval  | <i>S. tristichus</i> | 321.0734 | 68.1550 |
| MSB | 73521 | NM | Sandoval  | <i>S. tristichus</i> | 432.6698 | 73.8875 |
| MSB | 73522 | NM | Sandoval  | <i>S. tristichus</i> | 477.7920 | 80.9468 |

**Collections Data and Raw Measurements for all Male Specimens**

| <b>Collection ID</b> | <b>State</b> | <b>County</b> | <b>Species</b> | <b>TBA</b>            | <b>SVL</b> |         |
|----------------------|--------------|---------------|----------------|-----------------------|------------|---------|
| MSB                  | 55935        | NM            | Lea            | <i>S. consobrinus</i> | 181.2423   | 55.7758 |
| MSB                  | 56567        | NM            | Lea            | <i>S. consobrinus</i> | 88.1075    | 46.2609 |
| MSB                  | 60415        | NM            | Lea            | <i>S. consobrinus</i> | 93.1774    | 44.4544 |
| MSB                  | 60490        | NM            | Lea            | <i>S. consobrinus</i> | 182.5660   | 54.8789 |
| MSB                  | 60500        | NM            | Lea            | <i>S. consobrinus</i> | 103.6669   | 48.5572 |
| MSB                  | 60519        | NM            | Lea            | <i>S. consobrinus</i> | 108.5186   | 52.3983 |
| MSB                  | 60928        | NM            | Lea            | <i>S. consobrinus</i> | 135.1659   | 50.2885 |
| MSB                  | 74876        | NM            | Lea            | <i>S. consobrinus</i> | 98.3415    | 45.9313 |
| MSB                  | 56755        | NM            | Roosevelt      | <i>S. consobrinus</i> | 120.4711   | 47.2582 |

|      |       |    |            |                       |          |         |
|------|-------|----|------------|-----------------------|----------|---------|
| MSB  | 56757 | NM | Roosevelt  | <i>S. consobrinus</i> | 106.0882 | 51.1712 |
| MSB  | 59896 | NM | Roosevelt  | <i>S. consobrinus</i> | 156.1133 | 54.7106 |
| MSB  | 57622 | NM | Roosevelt  | <i>S. consobrinus</i> | 148.0697 | 51.9604 |
| MSB  | 16805 | NM | Bernalillo | <i>S. consobrinus</i> | 243.4671 | 52.8744 |
| MSB  | 22948 | NM | Eddy       | <i>S. consobrinus</i> | 454.0167 | 63.8612 |
| MSB  | 38492 | NM | Eddy       | <i>S. consobrinus</i> | 347.5925 | 62.1405 |
| MSB  | 52122 | NM | Eddy       | <i>S. consobrinus</i> | 361.0383 | 64.1824 |
| MSB  | 22951 | NM | Eddy       | <i>S. consobrinus</i> | 405.1210 | 67.4908 |
| MSB  | 22953 | NM | Eddy       | <i>S. consobrinus</i> | 276.6490 | 61.8638 |
| MSB  | 38496 | NM | Eddy       | <i>S. consobrinus</i> | 399.3252 | 66.0095 |
| MSB  | 39040 | NM | Eddy       | <i>S. consobrinus</i> | 348.8635 | 61.4382 |
| MSB  | 39042 | NM | Eddy       | <i>S. consobrinus</i> | 381.1943 | 62.1360 |
| MSB  | 48543 | NM | Eddy       | <i>S. consobrinus</i> | 403.0482 | 60.9830 |
| MSB  | 48730 | NM | Eddy       | <i>S. consobrinus</i> | 540.0926 | 70.0109 |
| MSB  | 60574 | NM | Eddy       | <i>S. consobrinus</i> | 261.7445 | 54.0011 |
| MSB  | 66893 | NM | Eddy       | <i>S. consobrinus</i> | 352.3885 | 65.8394 |
| MSB  | 57617 | NM | Roosevelt  | <i>S. consobrinus</i> | 122.9895 | 49.0768 |
| MSB  | 57624 | NM | Roosevelt  | <i>S. consobrinus</i> | 118.2200 | 48.9209 |
| MSB  | 57634 | NM | Roosevelt  | <i>S. consobrinus</i> | 119.6035 | 47.9229 |
| MSB  | 12253 | NM | Sandoval   | <i>S. consobrinus</i> | 284.2871 | 62.9511 |
| MSB  | 25916 | NM | Sandoval   | <i>S. consobrinus</i> | 368.8750 | 65.1447 |
| TCWC | 18839 | TX | Kerr       | <i>S. consobrinus</i> | 408.2949 | 61.2253 |
| TCWC | 4464  | TX | Kerr       | <i>S. consobrinus</i> | 227.6859 | 54.7788 |
| TCWC | 4465  | TX | Kerr       | <i>S. consobrinus</i> | 249.1970 | 51.5361 |
| TCWC | 4466  | TX | Kerr       | <i>S. consobrinus</i> | 340.9474 | 60.6942 |
| TCWC | 48910 | TX | Uvalde     | <i>S. consobrinus</i> | 317.7559 | 55.8610 |
| TCWC | 48914 | TX | Uvalde     | <i>S. consobrinus</i> | 298.6615 | 53.1151 |
| TCWC | 48927 | TX | Uvalde     | <i>S. consobrinus</i> | 237.1561 | 53.0383 |
| TCWC | 51200 | TX | Uvalde     | <i>S. consobrinus</i> | 224.1894 | 50.4316 |
| TCWC | 51211 | TX | Uvalde     | <i>S. consobrinus</i> | 259.4330 | 58.8137 |
| TCWC | 51216 | TX | Uvalde     | <i>S. consobrinus</i> | 325.9707 | 55.1616 |
| TCWC | 51986 | TX | Kerr       | <i>S. consobrinus</i> | 307.6230 | 56.4987 |
| TCWC | 83925 | TX | Kinney     | <i>S. consobrinus</i> | 231.3822 | 53.2575 |
| MSB  | 66903 | NM | Otero      | <i>S. cowlesi</i>     | 295.2529 | 60.1226 |
| MSB  | 7254  | NM | Torrance   | <i>S. cowlesi</i>     | 304.9273 | 58.5159 |
| MSB  | 10273 | NM | Torrance   | <i>S. cowlesi</i>     | 281.5453 | 55.6281 |
| MSB  | 10274 | NM | Torrance   | <i>S. cowlesi</i>     | 387.3349 | 60.0363 |
| MSB  | 10275 | NM | Torrance   | <i>S. cowlesi</i>     | 433.9721 | 63.9081 |
| MSB  | 18856 | NM | Torrance   | <i>S. cowlesi</i>     | 363.6948 | 56.5156 |
| MSB  | 18866 | NM | Torrance   | <i>S. cowlesi</i>     | 306.2224 | 55.0405 |

|      |       |    |          |                   |          |         |
|------|-------|----|----------|-------------------|----------|---------|
| MSB  | 18897 | NM | Torrance | <i>S. cowlesi</i> | 510.4993 | 66.3333 |
| MSB  | 21104 | NM | Torrance | <i>S. cowlesi</i> | 428.0742 | 62.1130 |
| MSB  | 21106 | NM | Torrance | <i>S. cowlesi</i> | 337.0792 | 60.6567 |
| MSB  | 21111 | NM | Torrance | <i>S. cowlesi</i> | 234.8987 | 50.5936 |
| MSB  | 21112 | NM | Torrance | <i>S. cowlesi</i> | 511.5112 | 69.2392 |
| MSB  | 21116 | NM | Torrance | <i>S. cowlesi</i> | 243.5987 | 54.6450 |
| MSB  | 21118 | NM | Torrance | <i>S. cowlesi</i> | 500.4126 | 70.6780 |
| MSB  | 21120 | NM | Torrance | <i>S. cowlesi</i> | 327.4966 | 57.2691 |
| MSB  | 21121 | NM | Torrance | <i>S. cowlesi</i> | 447.4290 | 64.1158 |
| MSB  | 21124 | NM | Torrance | <i>S. cowlesi</i> | 369.3464 | 64.6491 |
| MSB  | 21201 | NM | Torrance | <i>S. cowlesi</i> | 362.8763 | 61.4913 |
| MSB  | 21202 | NM | Torrance | <i>S. cowlesi</i> | 494.5778 | 62.6123 |
| MSB  | 21204 | NM | Torrance | <i>S. cowlesi</i> | 370.8063 | 55.3519 |
| MSB  | 21205 | NM | Torrance | <i>S. cowlesi</i> | 500.6512 | 65.7948 |
| MSB  | 26153 | NM | Torrance | <i>S. cowlesi</i> | 434.1302 | 65.1019 |
| MSB  | 7256  | NM | Valencia | <i>S. cowlesi</i> | 274.4211 | 56.8248 |
| MSB  | 21149 | NM | Valencia | <i>S. cowlesi</i> | 347.8514 | 65.4772 |
| MSB  | 21150 | NM | Valencia | <i>S. cowlesi</i> | 284.8648 | 61.8063 |
| MSB  | 25805 | NM | Valencia | <i>S. cowlesi</i> | 387.8761 | 65.8234 |
| MSB  | 72549 | NM | Valencia | <i>S. cowlesi</i> | 318.8615 | 61.8655 |
| MSB  | 72551 | NM | Valencia | <i>S. cowlesi</i> | 270.4430 | 59.5551 |
| MSB  | 72553 | NM | Valencia | <i>S. cowlesi</i> | 359.3334 | 57.8202 |
| MSB  | 60364 | NM | Otero    | <i>S. cowlesi</i> | 390.9751 | 66.4605 |
| MSB  | 21928 | NM | Torrance | <i>S. cowlesi</i> | 195.0064 | 46.0954 |
| MSB  | 26317 | NM | Torrance | <i>S. cowlesi</i> | 158.6268 | 47.0995 |
| MSB  | 7089  | NM | Valencia | <i>S. cowlesi</i> | 409.1367 | 65.1113 |
| MSB  | 7091  | NM | Valencia | <i>S. cowlesi</i> | 309.0205 | 59.0148 |
| MSB  | 21195 | NM | Valencia | <i>S. cowlesi</i> | 225.2664 | 52.1502 |
| MSB  | 21193 | NM | Valencia | <i>S. cowlesi</i> | 415.1511 | 67.7589 |
| MSB  | 25919 | NM | Valencia | <i>S. cowlesi</i> | 350.8069 | 70.4360 |
| MSB  | 25920 | NM | Valencia | <i>S. cowlesi</i> | 350.8069 | 70.4360 |
| MSB  | 21197 | NM | Valencia | <i>S. cowlesi</i> | 294.1080 | 58.9256 |
| TCWC | 16016 | TX | Brewster | <i>S. cowlesi</i> | 377.7102 | 58.9717 |
| TCWC | 16019 | TX | Brewster | <i>S. cowlesi</i> | 315.6495 | 53.9358 |
| TCWC | 16022 | TX | Brewster | <i>S. cowlesi</i> | 322.1553 | 58.2118 |
| TCWC | 16023 | TX | Brewster | <i>S. cowlesi</i> | 456.3237 | 58.9042 |
| TCWC | 52206 | TX | Edwards  | <i>S. cowlesi</i> | 331.4372 | 53.6639 |
| TCWC | 52207 | TX | Edwards  | <i>S. cowlesi</i> | 241.3049 | 52.4845 |
| TCWC | 52208 | TX | Edwards  | <i>S. cowlesi</i> | 389.7406 | 58.1135 |
| TCWC | 52210 | TX | Edwards  | <i>S. cowlesi</i> | 263.5506 | 51.3696 |

|      |       |    |            |                      |          |         |
|------|-------|----|------------|----------------------|----------|---------|
| TCWC | 52212 | TX | Edwards    | <i>S. cowlesi</i>    | 284.4878 | 55.9254 |
| TCWC | 35345 | TX | Jeff Davis | <i>S. cowlesi</i>    | 415.1047 | 58.7171 |
| TCWC | 35346 | TX | Jeff Davis | <i>S. cowlesi</i>    | 342.7814 | 54.3209 |
| TCWC | 39871 | TX | Jeff Davis | <i>S. cowlesi</i>    | 176.6597 | 45.6918 |
| TCWC | 81657 | TX | Jeff Davis | <i>S. cowlesi</i>    | 368.2654 | 58.0291 |
| TCWC | 81658 | TX | Jeff Davis | <i>S. cowlesi</i>    | 327.2380 | 55.7157 |
| TCWC | 81659 | TX | Jeff Davis | <i>S. cowlesi</i>    | 436.3496 | 63.5442 |
| TCWC | 39994 | TX | Terrell    | <i>S. cowlesi</i>    | 238.5597 | 53.8550 |
| TCWC | 80230 | TX | Terrell    | <i>S. cowlesi</i>    | 372.5834 | 58.8439 |
| TCWC | 38873 | TX | Val Verde  | <i>S. cowlesi</i>    | 434.7058 | 62.0791 |
| TCWC | 64701 | TX | Val Verde  | <i>S. cowlesi</i>    | 451.5070 | 59.1603 |
| TCWC | 16024 | TX | Brewster   | <i>S. cowlesi</i>    | 361.0415 | 59.2048 |
| TCWC | 16025 | TX | Brewster   | <i>S. cowlesi</i>    | 214.2101 | 58.6182 |
| TCWC | 52211 | TX | Edwards    | <i>S. cowlesi</i>    | 332.8390 | 56.6714 |
| TCWC | 25919 | TX | Jeff Davis | <i>S. cowlesi</i>    | 456.5617 | 69.9480 |
| TCWC | 25920 | TX | Jeff Davis | <i>S. cowlesi</i>    | 350.8069 | 70.4360 |
| TCWC | 35344 | TX | Jeff Davis | <i>S. cowlesi</i>    | 175.9212 | 46.2102 |
| TCWC | 38729 | TX | Jeff Davis | <i>S. cowlesi</i>    | 175.9856 | 47.8658 |
| TCWC | 71820 | TX | Terrell    | <i>S. cowlesi</i>    | 270.3330 | 57.7429 |
| TCWC | 64185 | TX | Val Verde  | <i>S. cowlesi</i>    | 248.3585 | 54.7962 |
| TCWC | 64700 | TX | Val Verde  | <i>S. cowlesi</i>    | 217.8578 | 50.8186 |
| MSB  | 82491 | CO | Montezuma  | <i>S. tristichus</i> | 470.5954 | 75.3390 |
| MSB  | 82492 | CO | Montezuma  | <i>S. tristichus</i> | 604.6043 | 80.0847 |
| MSB  | 21946 | NM | MicKinley  | <i>S. tristichus</i> | 181.7984 | 52.4868 |
| MSB  | 13064 | NM | Bernalillo | <i>S. tristichus</i> | 476.1021 | 65.5888 |
| MSB  | 13162 | NM | Bernalillo | <i>S. tristichus</i> | 633.1817 | 72.6647 |
| MSB  | 13165 | NM | Bernalillo | <i>S. tristichus</i> | 406.4148 | 63.3191 |
| MSB  | 13167 | NM | Bernalillo | <i>S. tristichus</i> | 490.9825 | 71.1721 |
| MSB  | 14674 | NM | Bernalillo | <i>S. tristichus</i> | 377.9645 | 63.7296 |
| MSB  | 14794 | NM | Bernalillo | <i>S. tristichus</i> | 291.0760 | 58.2575 |
| MSB  | 15122 | NM | Bernalillo | <i>S. tristichus</i> | 408.6303 | 67.5817 |
| MSB  | 16575 | NM | Bernalillo | <i>S. tristichus</i> | 230.7800 | 56.3099 |
| MSB  | 16889 | NM | Bernalillo | <i>S. tristichus</i> | 388.7497 | 64.4724 |
| MSB  | 21211 | NM | Bernalillo | <i>S. tristichus</i> | 353.9186 | 61.4328 |
| MSB  | 25945 | NM | Bernalillo | <i>S. tristichus</i> | 382.2088 | 63.4877 |
| MSB  | 25946 | NM | Bernalillo | <i>S. tristichus</i> | 396.8369 | 65.5932 |
| MSB  | 52810 | NM | Cibola     | <i>S. tristichus</i> | 425.6486 | 68.8039 |
| MSB  | 73225 | NM | Cibola     | <i>S. tristichus</i> | 355.5891 | 65.3661 |
| MSB  | 11959 | NM | MicKinley  | <i>S. tristichus</i> | 392.4387 | 62.5325 |
| MSB  | 76407 | NM | MicKinley  | <i>S. tristichus</i> | 417.7706 | 64.5815 |

|     |       |    |            |                      |          |         |
|-----|-------|----|------------|----------------------|----------|---------|
| MSB | 12252 | NM | Sandoval   | <i>S. tristichus</i> | 398.3997 | 62.0818 |
| MSB | 12399 | NM | Sandoval   | <i>S. tristichus</i> | 442.7115 | 68.8376 |
| MSB | 14336 | NM | Sandoval   | <i>S. tristichus</i> | 402.5499 | 69.5504 |
| MSB | 16805 | NM | Sandoval   | <i>S. tristichus</i> | 243.4671 | 52.8744 |
| MSB | 17267 | NM | Sandoval   | <i>S. tristichus</i> | 439.1107 | 69.5592 |
| MSB | 17312 | NM | Sandoval   | <i>S. tristichus</i> | 234.2556 | 60.7991 |
| MSB | 17313 | NM | Sandoval   | <i>S. tristichus</i> | 286.5366 | 58.8229 |
| MSB | 17314 | NM | Sandoval   | <i>S. tristichus</i> | 277.8326 | 59.5461 |
| MSB | 14793 | NM | Bernalillo | <i>S. tristichus</i> | 411.9320 | 65.1778 |
| MSB | 47211 | NM | Cibola     | <i>S. tristichus</i> | 372.3691 | 60.0517 |
| MSB | 54838 | NM | Cibola     | <i>S. tristichus</i> | 395.2521 | 62.6585 |
| MSB | 61191 | NM | Cibola     | <i>S. tristichus</i> | 257.5200 | 57.3948 |
| MSB | 61192 | NM | Cibola     | <i>S. tristichus</i> | 260.3574 | 62.7159 |
| MSB | 76384 | NM | Cibola     | <i>S. tristichus</i> | 313.1257 | 57.6857 |
| MSB | 76419 | NM | Cibola     | <i>S. tristichus</i> | 290.0902 | 62.8790 |
| MSB | 3293  | NM | MicKinley  | <i>S. tristichus</i> | 351.5956 | 64.2600 |
| MSB | 11951 | NM | MicKinley  | <i>S. tristichus</i> | 293.5277 | 55.2260 |
| MSB | 11956 | NM | MicKinley  | <i>S. tristichus</i> | 241.9808 | 54.4328 |
| MSB | 11957 | NM | MicKinley  | <i>S. tristichus</i> | 270.9618 | 56.2316 |
| MSB | 11961 | NM | MicKinley  | <i>S. tristichus</i> | 300.8999 | 57.1684 |
| MSB | 11962 | NM | MicKinley  | <i>S. tristichus</i> | 376.9084 | 61.9648 |
| MSB | 11963 | NM | MicKinley  | <i>S. tristichus</i> | 377.2648 | 60.1912 |
| MSB | 11964 | NM | MicKinley  | <i>S. tristichus</i> | 367.6270 | 60.7053 |
| MSB | 33238 | NM | MicKinley  | <i>S. tristichus</i> | 396.9748 | 60.5939 |
| MSB | 76386 | NM | MicKinley  | <i>S. tristichus</i> | 290.4570 | 56.7679 |
| MSB | 76388 | NM | MicKinley  | <i>S. tristichus</i> | 265.5109 | 58.4221 |
| MSB | 21762 | NM | San Juan   | <i>S. tristichus</i> | 385.0005 | 71.0418 |
| MSB | 21771 | NM | San Juan   | <i>S. tristichus</i> | 329.0188 | 66.1486 |
| MSB | 21776 | NM | San Juan   | <i>S. tristichus</i> | 419.2225 | 72.4818 |
| MSB | 21779 | NM | San Juan   | <i>S. tristichus</i> | 250.0572 | 57.0638 |
| MSB | 21780 | NM | San Juan   | <i>S. tristichus</i> | 354.5142 | 64.6650 |
| MSB | 21781 | NM | San Juan   | <i>S. tristichus</i> | 587.6860 | 78.1571 |
| MSB | 21791 | NM | San Juan   | <i>S. tristichus</i> | 491.5514 | 74.4960 |
| MSB | 21831 | NM | San Juan   | <i>S. tristichus</i> | 344.7640 | 66.7723 |
| MSB | 48037 | NM | San Juan   | <i>S. tristichus</i> | 430.2068 | 78.3072 |
| MSB | 48038 | NM | San Juan   | <i>S. tristichus</i> | 582.8336 | 81.5739 |
| MSB | 48039 | NM | San Juan   | <i>S. tristichus</i> | 467.4871 | 76.1718 |
| MSB | 58276 | NM | San Juan   | <i>S. tristichus</i> | 316.5791 | 67.7555 |
| MSB | 87791 | NM | San Juan   | <i>S. tristichus</i> | 258.3965 | 62.4324 |
| MSB | 17047 | NM | Sandoval   | <i>S. tristichus</i> | 358.3306 | 59.9697 |

MSB 73519 NM Sandoval *S. tristichus* 476.6398 65.0517



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