# How cryptic is cryptic diversity? Machine learning approaches to fine scale variation in the morphology of *Emys marmorata*.

Peter D Smits<sup>1</sup>, Kenneth D Angielczyk<sup>2</sup>, James F Parham<sup>3</sup>

<sup>1</sup>Committee on Evolution Biology, University of Chicago, <sup>2</sup>Department of Geology, Field Museum of Natural History, <sup>3</sup>Department of Geological Sciences, California State University – Fullerton

May 9, 2013

# Cryptic diversity

# Emys marmorata



wikimedia

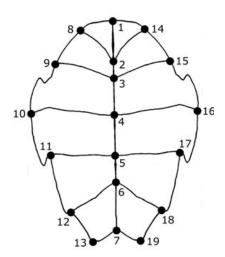
# Morphological hypotheses

## Phylogenetic hypotheses

Number of subspecies and where they occur.

## Methods: morphometrics

- plastral ("belly") shape
- landmarks averaged across bilat axis
- ► total 13 landmarks, 7 on bilat axis, 6 off
- geographic information known/inferred



## Methods: unsupervised learning

Fancy way of saying clustering or density estimation.

Partitioning around mediods (PAM) compared with "gap" statistic.

(dissimilarity based) Evidence accumulation clustering

#### Methods: supervised learning

Fancy way of saying classification and regression.

Here, features (principal components) predict class (subspecific assignment).

Multnomial logistic regression

Random forests

## Methods: model training and selection

Unknown appropriate number of features to "best" predict class. Want to minimize false positive, while maximizing true positive.

Split data set, 75-25, training and testing.

Tuning parameters via grid-search. Uncertainty via 10-fold cross-validation.

Multinomial logistic model selected via min AICc. Random forest model via max AUC ROC.

# Results: mophometrics

# Results: gap clustering

#### Results: d-EAC

Results: multinomial logistic regression

#### Results: random forests

#### Best classification scheme

#### **Future**

### Acknowledgements

- Ben Frable, Dallas Krentzel, Michael Foote
- COLLECTIONS

