

# EVALUATING ENM PERFORMANCE WITH NULL MODELS

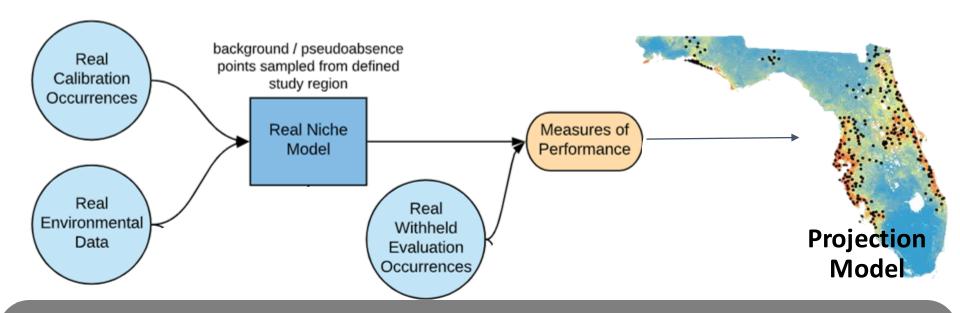
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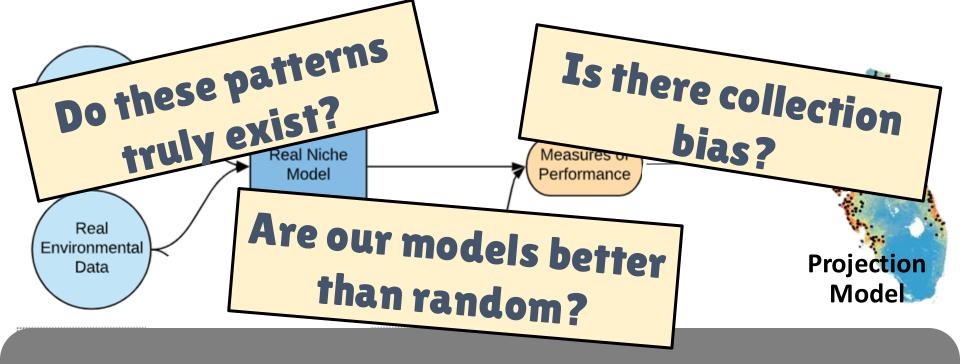




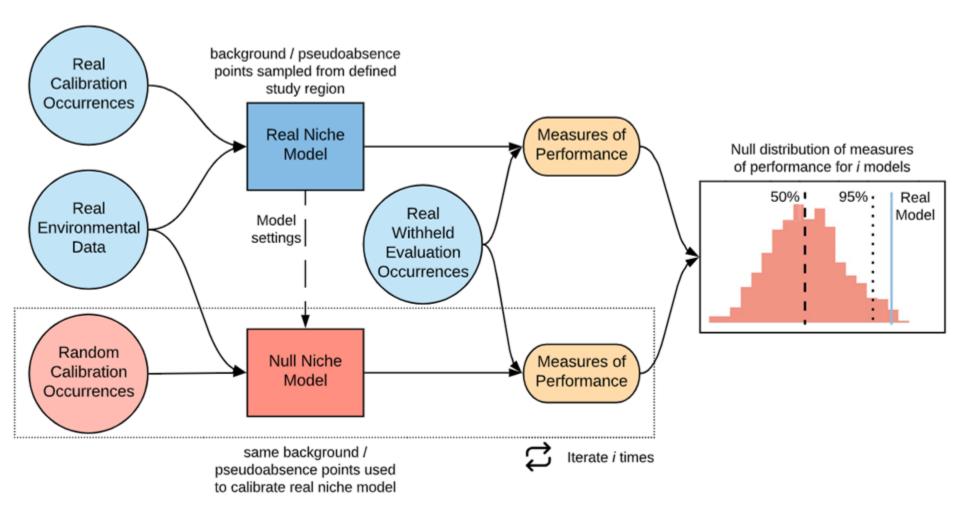




## The Typical ENM Process



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#### Actual vs. predicted habitat suitability values

#### PREDICTED VALUES

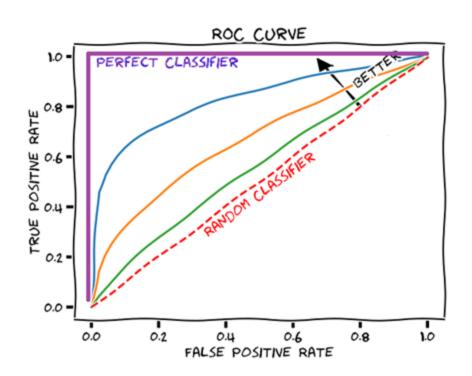
	Positive	Negative
Positive	True Positive (TP)	False Negative (FN)
Negative	False Positive (FP)	True Negative (TN)

**ACTUAL VALUES** 

#### PREDICTED VALUES

		Apple		Strawberry		
<b>ACTUAL VALUES</b>	Apple	<b>Č</b>	ě	Š	Ď	* *
	Strawberry	*	•		<b>ॐ</b>	****

#### **Area Under the Curve (AUC)**



False positive rate (specificity) vs. true positive rate (sensitivity)

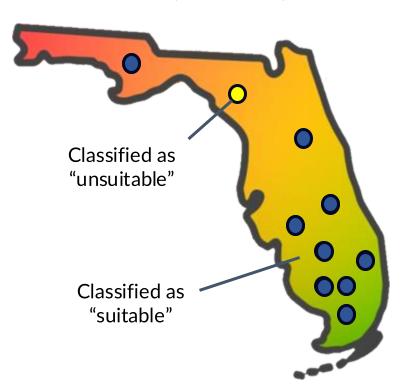
AUC = 1.0 -> perfect model AUC = 0.5 -> model performs no better than random AUC = <0.5 -> a sign of error

AUC evaluates a model's ability to distinguish between presence/background points

#### **Omission Rate (OR)**

OR = the proportion of known presence points that the model fails to predict as suitable

Example: OR = 0.1

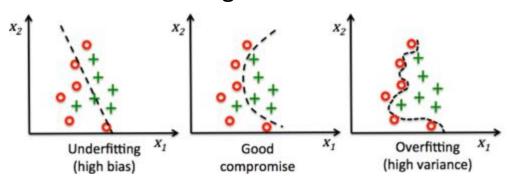


# Feature classes and regularization multiplier

Feature type	Interpretation	Shape
Linear	Continuous variable	_
Quadratic	Square of the variable	$ oxed{ } $
Hinge	As threshold type, but response after the threshold (knot) is linear	

#### Balance complexity and fit

- Feature classes: shape of the expected response curve
- Response curve: the relationship between environmental variables and suitability
- Regularization multiplier: penalty applied to reduce overfitting



### Sample code

Using tools in the ENMeval package

ENMevaluate generates models with different combinations of feature classes and RMs

```
eval <- ENMevaluate(
  occs = Galax_urceolata[, c("longitude", "latitude")],
  envs = vifStack,
  tune.args = list(fc = c("L", "Q"), rm = 1:2),
  partitions = "block",
  n.bg = 10000,
  parallel = FALSE,
  algorithm = 'maxent.jar',
)</pre>
```

## Sample code Using tools in the ENMeval package

Filter the resulting models by selecting for the one with the lowest omission rate and highest area under the curve value

#### How does ENMevaluate generate null models?

- 1 Use the same number of presence points as the empirical model, but sample them randomly from the background
- 2 Fit models with those random points using the same algorithm and environmental predictors
- 3 Evaluate performance using real withheld presence data
- 4 Repeat many times to generate a **null distribution** of performance metrics
- 5 Compare the **empirical model's AUC and omission rate** to the null distribution to assess significance

### Sample code

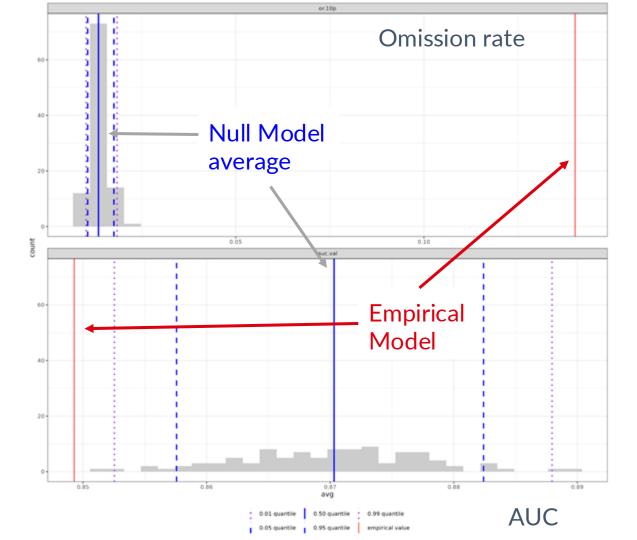
#### Using tools in the ENMeval package

#### Generate null models using parameters from optimal model

```
# Load optimal feature class (fc) and regularization multiplier (rm)
opt.seq <- read.delim("data/05_ENMs/Galax_urceolata_OptModel.txt")</pre>
# Extract parameters
fc <- opt.seq$fc</pre>
rm <- opt.seg$rm
# Run ENMnulls with optimal parameters and 100 iterations
spec.mod.null <- ENMnulls(</pre>
  eval,
  mod.settings = list(fc = fc, rm = rm),
  no.iter = 100
```

#### Output:

Plot shows the median AUC value and OR for the empirical model (red) against the null distribution values (blue)



#### Sample code

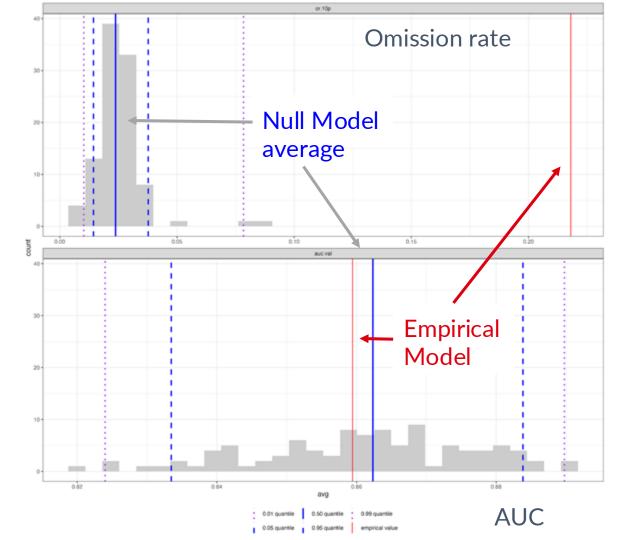
Using tools in the ENMeval package

Generate a new suite of models with a wider range of parameters

```
eval <- ENMevaluate(
  occs = Galax_urceolata[, c("longitude", "latitude")],
  envs = vifStack,
  tune.args = list(fc < c("L", "Q", "H", "LQ", "QH", "LH", "LQH"), rm = 1:5),
  partitions = "block",
  n.bg = 10000,
  parallel = FALSE,
  algorithm = 'maxent.jar',
)</pre>
```

## Output:

New optimal model: fc=H, rm=1



#### Why create null models?

- Determine whether model performance is better than expected by chance.
- Quantify how much better your model is than a null expectation.
- Avoid false confidence by evaluating for bias, spatial autocorrelation and overfitting
- Add rigor to model selection and interpretation