

## Model selection

1. Controversial area of statistics
2. Several alternatives - different "schools of thought"
3. Depends on your aim in fitting a model
4. ...and your study system

## 1. Hypothesis testing (t-test, F test)

## 4. Information theoretic (IT)

 approach (Burnham \& Anderson 2002) Specify a priori 10-15 models. Calculate differences in AIC2. No model selection (Bolker 2008) Only remove interactions
3. Classical stepwise selection (AIC, BIC)

## I. Hypothesis testing

- Drop least significant term
- Refit model
- Continue until only significant terms

I suggest never using this approach

- The best-fitting model may include nonsignificant terms
- Referees will (rightly) criticize this approach
- Consider what a P-value actually represents


## 2. Do nothing

- Perfectly valid (and you can't be criticized for the model selection approach you might otherwise use)
- Illustrates which terms in the model have significance and which don't (this could be your main question)
- A priori you selected certain covariates, so why remove them?
-(do remove collinear terms)

3. Classical stepwise selection - Use backward (start with full model and remove terms) or forward (start just with intercept and add terms) selection

- Use Akaike Information Criteria (AIC) to arrive at best-fitting model (also BIC, and for Bayesian models DIC, WAIC)

4. Information theoretic (IT) approach - Formulate (a priori) 10-15 alternative models

- Run all models, then compare using AIC
- Advocated by respected statisticians (Burnham \& Anderson, 2002)
- A very powerful approach
- ....but requires a lot of information/understanding
- Usually the case in fisheries models

| model | fitted model | source |
| :--- | :--- | :--- |
| M01 | temperature + salinity | Heuts (1947) |
| M02 | presence/absence of fish predators | Hoogland et al. (1956) |
| M03 | latitude x longitude | Münzig (1963) |
| M04 | temperature | Wootton (1976) |
| M05 | presence/absence dragonfly larvae | Reimchen (1994) |
| M06 | pH | Giles (1983) |
| M07 | elevation | Raeymaekers et al. (2007) |
| M08 | salinity | Myre \& Klepaker (2009) |
| M09 | presence/absence Schistocephalus solidus | MacColl et al. (2013) |
| M10 | presence/absence Pungitius pungitius | Reimchen et al. (2013) |
| M11 | turbidity + presence/absence of fish predators | Spence et al. (2013) |
| M12 | pH + presence/absence of fish predators | Klepaker et al. (2016) |
| M13 | pH + presence/absence of fish predators + turbidity | Magalhaes et al. (2016) |
| M14 | presence/absence of fish predators + P. pungitius | this study |
| M15 | temperature + standard length + pH | Mor |

## My suggestion

1. Use IT when possible
2. Alternatively, depending on the aims of your study, either

- Perform no selection, or
- Manual backward selection

3. Avoid using hypothesis testing

## How to deal with zero catches?

- Do not ignore zeros - these are critical data!
- Use an appropriate distribution that can accommodate zero observations
- Simulate from your model to ensure the model accommodates the proportion of zeros in the data
- We will do this (Hilsha analysis)


## How many zeros is too many?

- No specific threshold
- Fit model, then simulate from it
- Does the observed number exceed the predicted (by a lot)

What distribution is appropriate for (many) zeros?

- Gaussian (?), Poisson, negative binomial, Bernoulli, binomial
- Model validation: check by simulating from model and compare proportion of zeros in simulated data sets with observed proportion - they should match
- Use 'testZerolnflation' command in 'DHARMa' package
- We will do this (Hilsha analysis)


## Why do we get lots of zeros?

- Unsuitable conditions - no catch
- Suitable conditions - no catch
- Suitable conditions - not catchable
- Suitable conditions - make error

What type<br>of zeros<br>do you<br>have?

## How to handle lots of zeros

- Fit zero-inflated (mixture) models
- Fit zero-adjusted (hurdle) models


## ZIP, ZAP!

- Zero-inflated models differ from zeroadjusted models
- Zero-inflated models - model zeros as counts (some of which are zero)
- Zero-adjusted models explicitly model zeros as a Bernoulli model, and counts (zerotruncated data) using Poisson, NB, Gamma


## Zero-inflated models

- Model data in two parts:
- Binomial part; zeros vs. count (use binomial distribution)
- Zero-truncated data, using Poisson, negative binomial, gamma
- Able to identify which variables result in a catch (binomial part) and if a catch occurs, the size of the catch (zero-truncated part)
- We will use a ZINB model with the Hilsha analysis


## Tweedie distribution

- A family of distributions
- Not widely used
- Easy to implement with the 'glmmTMB' package
- Able to generate a compound PoissonGamma distribution


## Approach

1. Formulate the question Standardise CPUE for Bangladesh hilsha catch

