### No free lunch in inference

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### no free lunch

- ➤ Conjecture: Data-driven model tuning can increase the accuracy of a point estimate, but cannot decrease its uncertainty (without further strong assumptions)
- accuracy: e.g. mean squared error
- uncertainty: e.g. width of the confidence interval

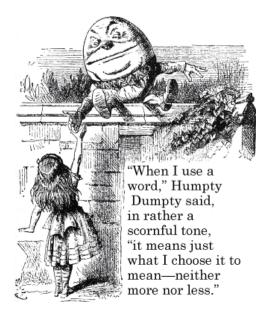
# scope of this talk

- ▶ full-rank (p < n), **non-sparse** problems
- science oriented (ecology/evolution)

# what are we doing when we do statistics?

- exploration
  - look for interesting patterns
  - confirm with followup observations
- prediction
  - best guess at future outcomes under specified conditions
- inference
  - estimate effects of processes and their uncertainty

# terminology



# what do I mean by inference?

- **not** concerned with *formal* causal inference
- evaluation of uncertainty
- what would we expect to see in future data?
  - coefficients: uncertainty around the effect of a change in the predictors
  - predictions: uncertainty around the value observed for specified predictor values
  - p-values: uncertainty around a counterfactual null

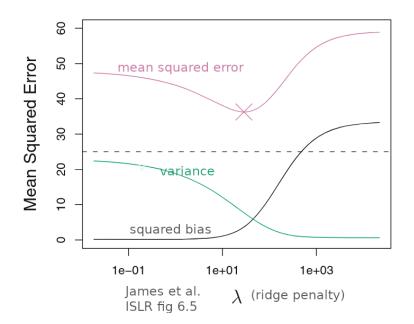
# data-driven model tuning for point estimates

- avoid omitting potentially important predictors
- avoid overfitting
- ➤ ≈ optimize bias-variance tradeoff



Rackham 1918 Wikipedia

# e.g. bias-variance tradeoff in ridge regression



## data-driven tuning

- stepwise/subset regression, ridge/lasso/elastic net, random forests, boosting . . .
- ▶ need to choose *appropriate* model complexity
  - model size (selection) or complexity (shrinkage)
    - estimate out-of-sample error without (explicit) cross-validation (AIC, Cp, BIC, out-of-bag error, ...)
    - or explicit cross-validation
- may need to tune model hyperparameters (e.g. via cross-validation)

## but what about uncertainty?

- statistical learning strongly focused on prediction
- but appropriate decisions require uncertainty quantification!
  - well appreciated in clinical trials
  - underappreciated in modern data science?

# how do we assess uncertainty quantification?

- false positive/type 1 error rate
- **coverage**: does an x% confidence interval include the true value x% of the time?
- (mentioned  $0 \times$  in James et al. (2013),  $1 \times$  in Hastie et al. (2009))

# coverage example (Li et al. 2018)

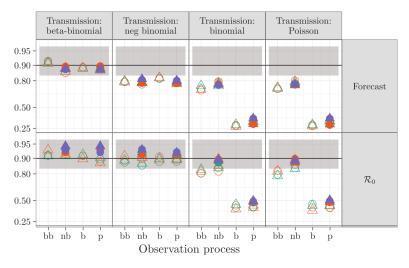


Figure 1: coverage plot for 90% intervals

# why coverage is better than type 1 error (mini-rant)

- type 1 error focuses on rejecting null hypotheses
- NH  $(\beta = 0)$  never(?) true in applied problems outside physics
- ightharpoonup coverage reduces to type-1 error *if*  $\beta = 0$
- type 1 assessment encourages unrealistic simulation setups



#### naive selection methods

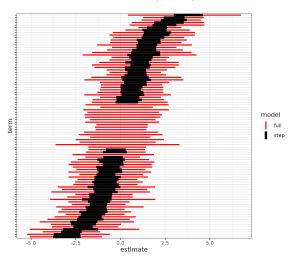
Post-selection inference that *ignores the selection process* is always be overoptimistic

▶ Altman et al. (1989) Any form of data-dependent variable selection is likely to lead to overoptimistic goodness of fit; we expect a worse fit to a new set of data; bootstrapping with stepwise variable selection gave similar individual predictions but larger confidence intervals for estimated survival probabilities.

See also Harrell et al. (1996).

# for example . . .

AIC-stepwise regression, simulated data lm(y ~ .) + step()  $n=100,~p=90,~\beta\sim U(-1,1),~\sigma_r^2=5;~90\%$  Cls



# coverage results (90% Cls)

model	n	n_ok	prop	lwr	upr
full	18000	16284	0.905	0.900	0.909
step	14008	4220	0.301	0.294	0.309

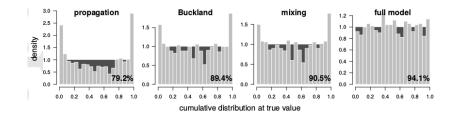
# what about something smarter, e.g. ridge regressoin?

- ▶ Obenchain (1977): properly constructed CI width ≥ full least-squares CI
- ▶ other methods may give OK results (e.g. Crivelli et al. (1995), Vinod (1987), Efron et al. (2020))
  - often involve additional assumptions e.g. on the distribution of parameters
  - bootstrapping methods must incorporate the full tuning process

# multimodel averaging (MMA)

- various methods for constructing MMA CIs (Burnham et al. 2002; Fletcher et al. 2012; Kabaila et al. 2016)
- ► MMA Cls are generally **too narrow** (Turek 2013; Kabaila et al. 2016; Dormann et al. 2018) but cf. Burnham et al. (2002)

# MMA results (Dormann et al. (2018), Figure 5)



## What about post-selection inference?

- Lots of exciting work
- focused on high dimensions, depends on strong assumptions
  - sparsity
  - ightharpoonup coefficient gap (minimum size of smallest  $|\beta|$ )
  - asymptopia
- e.g. Dezeure et al. (2015): When the truth (or the linear approximation of the true model) is nonsparse, the methods are expected to break down . . .
- See Cosma Shalizi's notes at {http://bactra.org/notebooks/post-model-selection-inference.html}

## paying for lunch in other ways

- go Bayesian!
  - Bayesian CIs are well-calibrated by definition, conditional on the model and the priors . . .
    - (Gelman et al. 1995; Cook et al. 2006; Talts et al. 2020)
- pseudo-Bayesian assumptions about effect size distributions

# conclusions: what should you do?

- ► for **inference**:
  - use the full model
  - ▶ a priori model reduction (Harrell 2001)
- ▶ for **prediction**:
  - use CIs from shrinkage estimates with caution
  - use non-neutral, informative Bayesian priors? (Crome et al. 1996)

# there ain't no such thing as a free lunch ...





## what are multifactorial systems?

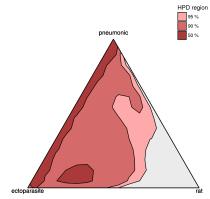
- many processes contribute to pattern
- quantify how each process affects the system, rather than testing whether we can detect its impact
- related:
  - Chamberlin's method of multiple working hypotheses (Raup et al. 1995)
  - tapering effect sizes (Burnham et al. 2002)

### conceptual problem: discretization

- model selection, or evidential statistics (Taper et al. 2016), focus on differentiating discrete hypotheses/models
- submodels are always straw men
- expand models to cover the whole space

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{Estimated contribution of plague transmission modes in Eyam 1665}

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