Bayesian statistics with R

7. Contrast scientific hypotheses with model selection

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Model selection

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- The proportion of explained variance R² is problematic, because the more variables you have, the bigger R² is.
- Idea: penalize models with too many parameters.

$$AIC = -2\log(L(\hat{ heta}_1,\ldots,\hat{ heta}_K)) + 2K$$

with L the likelihood and K the number of parameters θ_i .

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A measure of goodness-of-fit of the model to the data: the more parameters you have, the smaller the deviance is (or the bigger the likelihood is).

$$AIC = -2 \log(L(\hat{\theta}_1, \dots, \hat{\theta}_K)) + 2K$$

A penalty: twice the number of parameters K

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- Two models are difficult to distinguish if $\Delta AIC < 2$.

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 $DIC = -2 \log L(data \mid \theta) + 2p_D$

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- DIC is intended as a generalisation of AIC, and with little prior information, p_D should be approximately the true number of parameters.
- The model with the smallest DIC is estimated to be the model that would best predict a replicate dataset with same structure as that observed.

DIC in Jags

storks

#> Inference for Bugs model at "code/logistic.txt", fit using jags, #> 2 chains, each with 2000 iterations (first 1000 discarded) #> n.sims = 2000 iterations saved. Running time = 0.059 secs #> mu.vect sd.vect 2.5% 25% 50% 75% 97.5% Rhat n.eff #> a 1.555 0.054 1.449 1.518 1.555 1.593 1.658 1.011 220 #> b.rain -0.160 0.066 -0.283 -0.206 -0.162 -0.116 -0.026 1.004 1300 #> b.temp 0.035 0.062 -0.082 -0.008 0.033 0.075 0.165 1.007 330 #> deviance 204.639 2.446 201.823 202.855 203.982 205.837 210.943 1.006 260 #>

#> For each parameter, n.eff is a crude measure of effective sample size, #> and Rhat is the potential scale reduction factor (at convergence, Rhat=1). #>

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#> DIC info (using the rule: pV = var(deviance)/2)
#> pV = 3.0 and DIC = 207.6
#> DIC is an estimate of expected predictive error (lower deviance is better).
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- Hooten, M.B. and Hobbs, N.T. (2015), A guide to Bayesian model selection for ecologists. Ecological Monographs, 85: 3-28. https://doi.org/10.1890/14-0661.1
- Conn, P.B., Johnson, D.S., Williams, P.J., Melin, S.R. and Hooten, M.B. (2018), A guide to Bayesian model checking for ecologists. Ecol Monogr, 88: 526-542. https://doi.org/10.1002/ecm.1314