Software tools for Maximum Likelihood Estimation

Lesson 3 - Controlling the search and assessing uncertainty

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Outline:

- Parameterization
- Fixing parameters
- Bounding parameters
- Review and more detail on asymptotic standard errors
- More than you need to know about the delta method
- Profile likelihood Cls
- Self-test simulations and parametric bootstrap

Parameterization - what we want

- Formal parameters (e.g., what gets sent to nlminb):
 - can legally take values on the entire real number line or are restricted from searching outside bounds (or both)
- Formal parameters estimates have values on similar scales to one another
- Formal parameter estimates are not too highly correlated with one another

Functional invariance and parameter transformation

- Formal (estimated) params transformations of natural params
- Relies on functional invariance of MLE
 - if $NLL(\theta) = NLL(g(\theta))$ then search on θ or on $g(\theta)$
 - Example: estimate log_theta and calculate theta=exp(log_theta) inside the NLL function
 - Another example is natural parameters 0<theta<1 or
 - -1<theta<+1 (e.g., correlations)

Logit transformation (for 0-1 or -1 -+1)

$$b = \operatorname{logit}(a) = \operatorname{log}\left(rac{a}{1-a}
ight)$$
 $a = \operatorname{logit}^{-1}(b) = rac{1}{1+\exp(-b)}$

- $0 \le a \le 1, b$ on real number line, so estimate b then calculate a
- For $-1 \le c \le +1$ from b (estimate b and calculate c)

$$c = 2 \operatorname{logit}^{-1}(b) - 1 = rac{2}{1 + \exp(-b)} - 1$$

For similar quantities

- E.g., catchability for different survey gears, or year or location effects
- One approach: base and offsets
 - Define one level (gear, year, location) as base estimate offsets for others. Formal parameters are the base and offsets
 - Two gears: estimate logq_gill and off_trap, where logq_gill is the log-scale gill net catchability. In your NLL function calculate logq_trap=logq_gill+offset
 - for year-effects, estimate YE_0, offsets for the nyears-1 other years. Then YE[1]=YE_0, and (i>1) YE[i]=YE_0+offset[i-1]
- Alternative (and sometimes better) approach is through mean and deviations from mean

Forcing parameter order

- You want theta[1]< theta[2]< theta[3]
 - Estimate theta[1] and two increments
 - theta[2]=theta[1]+inc[1], theta[3]=theta[2]+inc[2]
- Thought experiment: suppose you are estimating selectivity-at-age and are working with 10 ages so you need sel[i], i=1 to 10.
 - You are willing to assume/define sel[10]=1. You want:
 - 0<=sel[i]<=1
 - sel[i]<=sel[i+1]

My solution

- Estimate "adjust_par" that determine "adjust" constrained between 0 and 1
- Loop backwards from i=9 to i=1, with sel[i]=adjust[i]*sel[i+1]

Getting things set up

```
1 logit=function(x) {
2    log(x/(1-x));
3 }
4
5 inv_logit=function(x) {
6    l/(1+exp(-x));
7 }
8
9 # tr_adjust might be starting values of transformed adjusts
10 tr_adjust=rep(logit(.9),9);
11 tr_adjust;
```

[1] 2.197225 2.197225 2.197225 2.197225 2.197225 2.197225 2.197225 2.197225 [9] 2.197225

Code like this inside NLL function

```
1 adjust = inv_logit(tr_adjust);
2 sel=rep(NA,10);
3 sel[10]=1;
4 for(i in 1:9) {
5   sel[10-i]=adjust[10-i]*sel[10-i+1]
6 }
7 sel
```

```
[1] 0.3874205 0.4304672 0.4782969 0.5314410 0.5904900 0.6561000 0.7290000 [8] 0.8100000 0.9000000 1.0000000
```

Bounds

- You specify these as vectors for lower and upper bounds
- These are passed to nlminb (so not part of RTMB)
- Behavior is different than in admb (bound pars will be failed convergence)
- Code squib

```
#create upper and lower bounds
```

```
lower = c(4,-5,-5,0); upper = c(10,1,5,10);
```

#fit with bounds, here ... are the other arguments we usually use

fit = nlminb(..., upper=upper, lower=lower);

Fixing (not estimating) a parameter

- We fix parameters using maps
- You create a map as a named list where parameters you want to fix are included as factors set to missing values
 E.g., if you wanted to fix log_linf:
 - gmmap = list(log_linf=factor(NA));
- The map is passed as an argument to MakeADFun obj=MakeADFun(..., map=grmap);

Exercises

- Modify the vonB program to bound the parameters (bounds in slide on bounds reasonable choices)
- Modify the vonB program to not estimate log_linf using a map
- See if you can combine setting bounds and fixing a parameter

Uncertainty, basic definitions of expected value, variance, covariance, and correlation

$$\mathrm{E}[X] = \int_{-\infty}^{\infty} x f(x) dx$$

$$\operatorname{Var}(X) = E[(X - E[X])^2]$$

$$Cov(X, Y) = E[(X - E[X])(Y - E[Y])]$$

$$\operatorname{Corr}(X,Y) = \operatorname{Cov}(X,Y)/(\sigma_X\sigma_Y)$$

Confidence Intervals (CIs)

- An interval that might contain the true value of a parameter (or other estimated quantity)
- The confidence level is the probability that under repeated sampling the interval does contain the true value.
- E.g., for a 95% CI, it is expected that 95% of intervals constructed in the same way will include the true value being estimated.

Asymptotic variance covariance matrix and Wald confidence intervals

- The estimated variances for the parameter estimators are the diagonal elements from matrix, SEs are their square roots.
- A Wald test assumes $Z=\left(\hat{\theta}-\theta_0\right)/SE$ comes from a N(0,1) distribution, so one would expect Pr(|Z|>1.96)=0.05
 - ullet One would reject $H_0: heta = heta_0$ at the 0.05 level if |Z| > 1.96
- A Wald CI inverts a Wald test

Wald Confidence Interval

- A $100(1-\alpha)\%$ CI is: $\hat{\theta}\pm\Phi^{-1}(1-\alpha/2)SE$, Φ is the CDF for a N(0,1) distribution, Φ^{-1} is the inverse CDF or quantile function for N(0,1)
- This is based on inverting a Wald test (i.e., find the Z values that just make the Wald test significant)
- ullet Special case 95% CI: $\hat{ heta} \pm 1.96SE$

Delta method

- This is what ADREPORT uses to get variances for derived (non-parameters) quantities.
- $\bullet \ \operatorname{C\^{o}v}[g(\underline{\theta}),h(\underline{\theta})] \cong \sum_{i} \sum_{j} \operatorname{C\^{o}v}[\theta_{i},\theta_{j}] \frac{\partial g(\underline{\theta})}{\partial \theta_{i}} \frac{\partial h(\underline{\theta})}{\partial \theta_{j}}$
 - Formula for variance is just special case of this.
- Main point is you can approximate variance of any function of parameters based on covariances of parameters and derivatives of function with respect to parameters
- If you can get Variances (and hence SEs) you can get Wald
 CIs

Likelihood ratio test

 Likelihood profile CI based on inverting a Likelihood Ratio test, with test statistic:

$$LRT = -2\lograc{\supig(L({ar heta}),{ar heta}\in{ar heta}ig)}{\sup(L({ar heta}),{ar heta}\in{ar heta}ig)} = -2\left[\log(\supig(L({ar heta}),$$

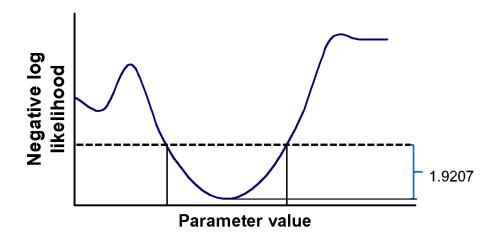
- ullet $LR_r \sim X_
 u^2$ with u df (based on number of restrictions)
- Of interest here is $\underline{\theta} \in \underline{\Theta}_0$ with one parameter fixed, testing if that focal parameter differs from fixed value.
 - ullet In this case $\operatorname{crit} = CDF_{X_1^2}^{-1}(1-lpha)$

Steps in Likelihood ratio test for one fixed parameter

- 1. Fit the full model and model with one focal parameter fixed, and extract likelihood (or log likelihood) from results and calculate the test statistic.
- 2. Find the critical value (it will always be 3.8415 if one parameter is fixed and your test is at the 0.05 level, otherwise use the chi-square quantile function).
- 3. Compare the test statistic with crit and declare significant if crit exceeded.

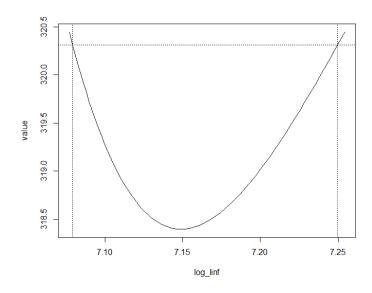
Likelihood profile CI steps

- 1. Fit the unrestricted model.
- 2. Repeatedly fit a restricted version of the model with the focal parameter fixed at a range of finely spaced values.
- 3. Find the upper and lower values (above and below the point estimate from the unrestricted model) that produce a significant likelihood ratio test.



Code to get likelihood profile Cl

```
library(TMB); #load for CI
log_linf_prof = tmbprofile(obj, "log_linf");
confint(log_linf_prof);
plot(log_linf_prof);
```



Coding issue

Once you load the TMB package, calls to MakeADFun() do not work unless they explicitly name the RTMB package as in:

RTMB::MakeADFun(...);

Limitation of likelihood profile Cls

- In TMB for parameters only
- Some have suggested constraining fit with penalty so desired values of derived quantities could be matched. Theory for this is not fully worked out and its not implemented.
- Hence worth exploring simulation approaches (aka parametric bootstrap
 - Simulations of course are for more than CIs

Simulations using RTMB

- To simulate data from your model you:
 - Mark data you want simulated in your NLL function y=OBJ(y);
 - After fitting (or before if you want to simulate from starting values)
 - simy=my_obj\$simulate()\$y
- Common plan: (1) replace data model was fit to with simulated data, (2) refit the model to this copy and save pertinent results. Repeat 1&2 many times and summarize.
- I will demo doing this for one data set and have you attempt to do a full simulation study.