F: TMB and MCMC

Fish 559; Day 3: 09h30-10h30



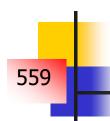
Basics of a Bayesian Assessment

Bayes Theorem:

$$p(\underline{\theta} \mid D) = \frac{L(D \mid \underline{\theta}) p(\underline{\theta})}{\int L(D \mid \underline{\theta}) p(\underline{\theta}) d\underline{\theta}}$$



- We need to specify the prior distribution, $p(\underline{\theta})$, in order to apply Bayes theorem.
- The objective is to represent the posterior by means of a (large) number of vectors of parameters.



Objectives for MCMC

- MCMC is a way to construct a (joint) Bayesian posterior for the parameters of a population dynamics model.
- The basic idea is to set up a (long) "chain" that starts at a pre-specified vector and that then traverses the posterior distribution.
- The sample from the posterior distribution is then every n'th element in the chain (ignoring the first "few" elements of the chain).

559

Overview of the MCMC algorithm

- Select an "initial" parameter vector (often the mode of the posterior) and compute its posterior density (the product of the likelihood and the prior).
- Generate a "new" parameter vector based on the current one (using a "jump" function) and compute its posterior density.
- Replace the current parameter vector by the new parameter vector with probability equal to the ratio of the new to the current posterior density.
- 4. Output the current parameter vector.
- 5. Repeat steps 2)- 4) many times!

Steps 2-4 are referred to as a cycle.



A Little More Detail

- A) Set *i* to 0.
- B) Set y to x.
- C) Apply the "jump function" to y to generate a new vector z (may only differ in one element from y) and compute f(z).
- D) If f(z) > f(y), set i=i+1, $x_i=z$, go to B).
- E) Generate $w \sim U[0,1]$. If w < f(z)/f(y), set i=i+1, $x_i=z$, go to B).
- F) Set i=i+1, $x_i=y$ and go to B).



The Jump Function

- The "jump" function should be chosen to optimize performance (but is usually selected for computational convenience).
- It should be possible to reach all possible parameter vectors "eventually" by applying the "jump" function long enough.
- The jump function depends on which MCMC algorithm you select. I have used:
 - A naive "Metropolis" algorithm
 - The No-Uturn sampler that can be found in the adnuts package. We will discuss this sampler in Lecture G.



Example-I

- Compute posterior distributions for the parameters of a logistic regression.
 - Formulate the problem as maximum likelihood estimation.
 - Select prior distributions.
 - Choose a jump function.



Example-II

The likelihood formulation of the problem.

$$-\ell nL = n.\ell n\sigma + \frac{1}{2\sigma^2} \sum_{i} (logit(y_i) - logit \left[\left[1 + exp(-\ell n 19 \frac{x_i - x_{50}}{x_{95} - x_{50}}) \right]^{-1} \right]$$

- The prior: uniform on the logarithms of x_{50} , x_{95} and σ .
- The jump function: multivariate normal with mean zero and variance-covariance matrix from a TMB fit.
- Hint: You will need the mvtnorm library to run the mcmc function.
- The code is in LectF1.R.



Example-III (R Implementation)

```
f1 <- function(x)</pre>
                                                    Calculate the
A50 <- \exp(x[1])
                                                 likelihood function
A95 < -\exp(x[2])
Sigma <- exp(x[3])
pred <- MakePred(co$Length,A50,A95)</pre>
pred <- log(pred/(1-pred))</pre>
SS <- (pred-co$Prob2)^2
SS <- sum(SS)
LogLike <- length(pred)*log(Sigma) +</pre>
sum(SS)/(2.0*Sigma^2)
return(LogLike)
                                     MakePred<-function(Length,A50,A95)
                                     pred <- 1/(1+exp(-1*log(19)*(Length-
            Vector of
                                    ·A50)/(A95-A50)))
        predicted values
                                     return(pred)
```



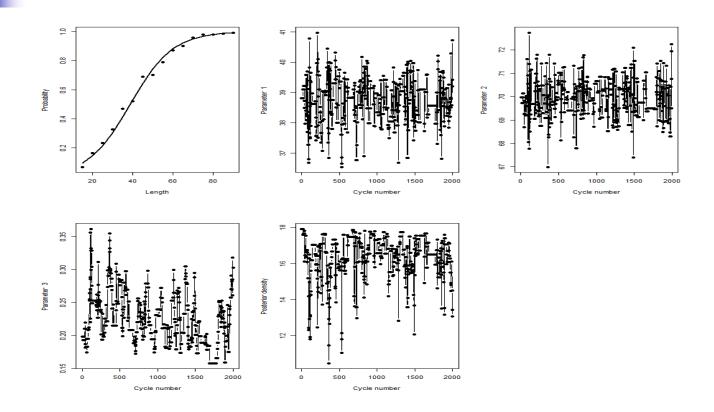
Example-IV (R Implementation)

```
DoMCMC<-
function(Xinit,Ndim,sd,cor,Nsim=1000,
Nburn=0,Nthin=1)
{
FileN <- "lect18.txt"
TheData<-scan(file=FileN,
what=list(Length=0,Prob=0))
co <- NULL
co$Length <- TheData$Length
co$Prob1 <- TheData$Prob
co$Prob2 <- log(TheData$Prob/(1-
TheData$Prob))
environment (f1) = environment()
```

```
Xcurr <- Xinit
Fcurr <- -1*f1(Xcurr)
Outs <- matrix(0,nrow=(Nsim-Nburn),ncol=(Ndim+1))
Ipnt <- 0; Icnt <- 0
for (Isim in 1:Nsim)
 Xnext <- rmvnorm(1, Xcurr, cov=cor, sd, rho, Ndim)</pre>
 Fnext <- -1*f1(Xnext)
 Rand1 <- log(runif(1,0,1))
 if (Fnext > Fcurr+Rand1)
 {Fcurr <- Fnext; Xcurr <- Xnext }
 if (Isim %% Nthin == 0)
  Ipnt <- Ipnt + 1
  if (Ipnt > Nburn)
   { Icnt <- Icnt + 1; Outs[Icnt,] <- c(Xcurr,Fcurr); }
```



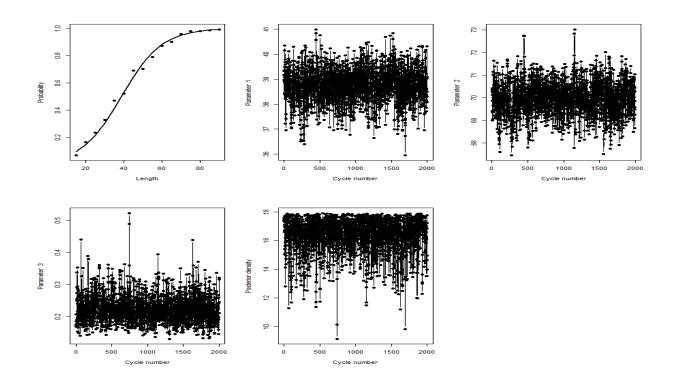
Example-V (Results-silly jump function)



sd <- c(0.1,0.1,0.1); cor <- matrix(c(1,0,0,0,1,0,0,0,1), ncol=3,nrow=3)

559

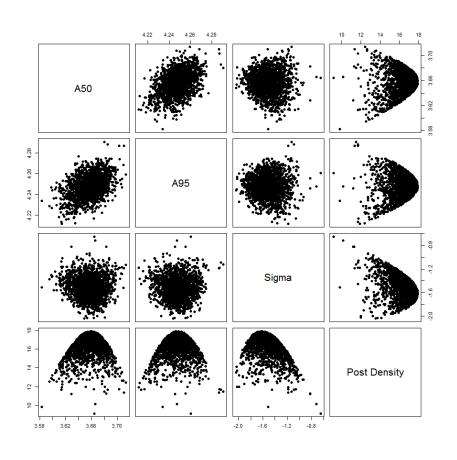
Example-VI (Results-Informative jump function)



sd <- c(0.015702,0.0093949,0.17678); cor <- matrix(c(1,0.3769,0,0.3769,1,0,0,0,1),ncol=3,nrow=3)



Example-VII (Results-Posterior correlations)





Example-IX

This problem can be done in one step because one can call TMB functions from R (see LectF2.R).

In general, the steps should be:

- Fit the model (maximum likelihood)
- Obtain the variance-covariance matrix (hessian=T)
- Call MCMC
- View MCMC diagnostics.



"Burn in" and "thinning"

- The impact of the initial parameter vector and the Markov nature of the algorithm are reduced by having a "burn in" period and by "thinning" the chain.
- To get x samples from the posterior:
 - Allow for a "burn-in" period (5-50% of the total chain length) to allow the algorithm to "set itself up". The results of this period are discarded.
 - "Thin" the chain by taking every n'th value in the chain.



Tricks of the (MCMC) trade - I

- The performance of MCMC deteriorates as the number of parameters increases.
 Therefore, integrate out any nuisance parameters analytically.
 - Examples (Walters and Ludwig, 1994).
 - The catchability coefficients;
 - The observation error standard deviations.

559

Tricks of the (MCMC) trade - II

- The performance of MCMC deteriorates if the parameters are highly correlated (often true for agestructured models).
- Use the .COR file to identify highly-correlated parameters ($\rho \ge 0.8$).
- Re-parameterize the model to reduce correlations:

Use

$$S_a = \left(1 + \exp\left[-\frac{(a - a_{50})}{\Delta}\right]\right)^{-1}$$

and not

$$S_a = \left(1 + \exp\left[-\frac{(a - a_{50})}{(a_{95} - a_{50})}\right]\right)^{-1}$$