## Scripts for Bayesian Phylogenetic Analysis Workshop

File conversion: <a href="https://www.sing-group.org/ALTER/">https://www.sing-group.org/ALTER/</a>

## **Example of Bayes Block: Burning phase+ no of chains**

Log start filename=bayes.log;				
Lset nst=6 rates=invgamma;				
<pre>prset statefreqpr=dirichlet(1,1,1,1);</pre>				
mcmc ngen=100000000 printfreq=100 samplefreq=100 burninfrac=0.25 nruns=2 nchains=4				
Log stop;				
end;				
<b>Example of Bayes Block: For ITS MSA</b>				
begin mrbayes;				
Log start filename=bayes.log;				
Lset nst=6 rates=propinv;				
Prset statefreqpr=dirichlet(1,1,1,1);				
mcmc ngen=10000000 printfreq=1000 samplefreq=1000 burninfrac=0.25 nruns=2 nchains=4 savebrlens=yes stoprule=yes stopval=0.01;				
Log stop;				
end;				

## **Analysis with multiple data partitions**

begin mrbayes;
Log start filename=bayes.log;

begin mrbaves:

```
charset locus2 = 496-943;
charset locus3 = 944-1902;
partition alldata = 3: locus1, locus2, locus3;
set partition = alldata;
Lset applyto= (1) nst=6 rates=propinv;
Lset applyto= (2) nst=2 rates=gamma;
Lset applyto= (3) nst=6 rates=gamma;

prset applyto= (all) statefreqpr=dirichlet(1,1,1,1);
unlink statefreq=(all) revmat=(all) shape=(all) pinvar=(all);

mcmc ngen= 100000000 relburnin=yes burninfrac=0.25 printfreq=1000 temp=0.15 samplefreq=1000 nchains=4 savebrlens=yes stoprule=yes stopval=0.01;
Log stop;
end;
```

## **MrModeltest2**

```
ITS
BEGIN MRBAYES;
       Lset nst=6 rates=propinv;
       Prset statefreqpr=dirichlet(1,1,1,1);
END;
Note: Substitution model: General Time Reversible (GTR)
BTUB
BEGIN MRBAYES;
       Lset nst=2 rates=gamma;
       Prset statefreqpr=dirichlet(1,1,1,1);
END;
Note: K80
EF
BEGIN MRBAYES;
       Lset nst=6 rates=gamma;
       Prset statefreqpr=dirichlet(1,1,1,1);
END;
Note: Substitution model: General Time Reversible (GTR)
COMBINE
BEGIN MRBAYES;
       Lset nst=6 rates=invgamma;
       Prset statefreqpr=fixed(equal);
END;
```