List of substitution models and priors used for the cetaceans dataset.

Partition	Substitution model	Invariant proportion
Alb 12	$HKY + \Gamma + I$	0.32
Alb 3	$TN93 + \Gamma + I$	0.22
Bdnf 12	$TN93 + \Gamma + I$	0.96
Bdnf 3	$TN93 + \Gamma + I$	0.24
Cmos 12	$GTR + \Gamma + I$	0.31
Cmos 3	$HKY + \Gamma + I$	0.20
Co1 12	$GTR + \Gamma + I$	0.76
Co1 3	$TN93 + \Gamma$	
Cytb 12	$HKY + \Gamma + I$	0.54
Cytb 3	$GTR + \Gamma + I$	0.16
Irbp 12	$HKY + \Gamma + I$	0.53
Irbp 3	$GTR + \Gamma + I$	0.19
Nd3 12	$TN93 + \Gamma + I$	0.36
Nd3 3	TN93	
Nd4 12	$TN93 + \Gamma + I$	0.38
Nd4 3	$HKY + \Gamma$	
Nd4l 12	$TN93 + \Gamma + I$	0.31
Nd4l 3	K81	
Noncoding	$GTR + \Gamma$	
Prion 12	$GTR + \Gamma + I$	0.23
Prion 3	$GTR + \Gamma + I$	0.17
Rag2 12	$TN93 + \Gamma + I$	0.17
Rag2 3	$HKY + \Gamma + I$	0.18
Rrna	$GTR + \Gamma + I$	0.35
Sry 12	GTR	
Sry 3	GTR	
Tbx4 12	$TVM + \Gamma + I$	0.42
Tbx4 3	$K81 + \Gamma + I$	0.28

Parameter	Prior distribution
FBD diversification rate	Uniform $[0; \infty[$
FBD turnover	Uniform[0; 1[
FBD sampling proportion	Uniform[0; 1[
Origin of the FBD process	Uniform $[0; \infty[$
HKY κ parameter	LogNormal(1.0,1.25)
TN93 κ_1 , κ_2 parameter	LogNormal(1.0,1.25)
GTR AC, AT, CG, GT rates	Gamma($\alpha = 0.05, \beta = 10.0$)
GTR AG rate	Gamma($\alpha = 0.05, \beta = 20.0$)
Shape of the Γ distribution	Exponential (1.0)
Linked clock rate	Uniform $[0; \infty[$