Table S7: Posterior probability of four putative scenarios, and type I and type II errors for scenario 1 in DIYABC[1] analysis of *Dipterocarpus turbinatus*

Scenario	1	2	3	4
Posterior probabilities	0.5269	0.14	0.1748	0.1583
Confidence intervals	0.4974-0.5564	0.0801-0.1998	0.1455-0.2041	0.1262-0.1905
Type I error	0.397			
Mean type II error	0.123 (0.059-0.159)			

[1] Cornuet J.M., Pudlo P., Veyssier J., Dehne-Garcia A., Gautier M., Leblois R., Marin J-M., Estoup A. (2014) DIYABC v.2.0: a software to make approximate Bayesian computation inferences about population history using single nucleotide polymorphism, DNA sequence and microsatellite data. *Bioinformatics* 30:1187-1189. doi: 10.1093/bioinformatics/btt763