

Quantiles	log-Evidence	hyperparameter
10/80/10	-2.76×10^4	3.17
15/70/15	-2.54×10^4	2.33
20/60/20	-2.37×10^4	2.33
23/54/23	-2.36×10^4	2.33
25/50/25	-2.36×10^4	2.33
27/46/27	-2.40×10^4	1.84
30/40/30	-2.44×10^4	1.84

Table 1: Data discretisation for NASC gene clustering.