**File S12 [A-G].** Prior (black) andposterior (blue) distribution for PopABC analysis of region two versus region four. Parameters investigated are mutation rate (mut rate [A]), migration into sampling regions two and four (labelled mig1 [B] and mig2 [C] respectively), effective population size of sampling regions two and four (labelled Ne1 [D] and Ne2 [E] respectively), effective population size of the ancestral population (labelled NeA [F]) and time since divergence of sampling regions two and four (labelled t1 [G]).

Macintosh HD:Users:stantondw:Documents:Okapi:Analysis:popABC:output:posterior_distributions_2v4.pdf

A

B

C

D

E

F

G

References

1. Lopes JS, Balding D, Beaumont M (2009) PopABC: a program to infer historical demographic parameters. Bioinformatics 25: 2747-2749.