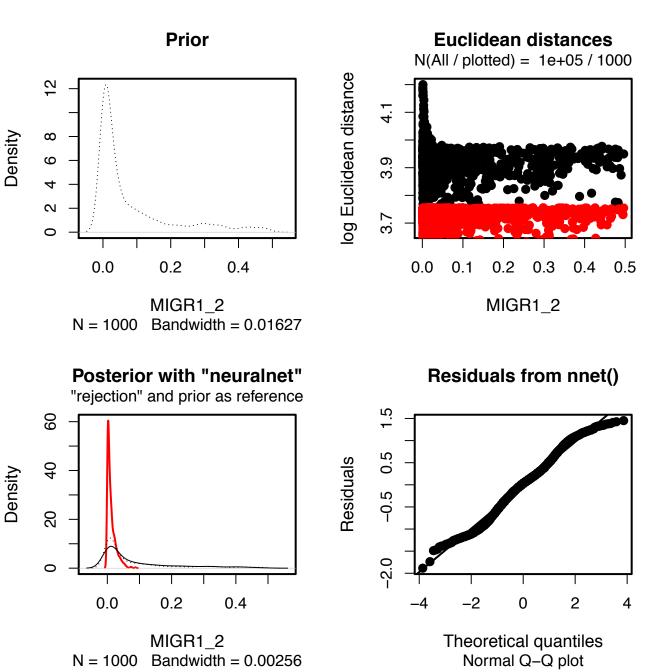
ABC model fit and posterior distributions

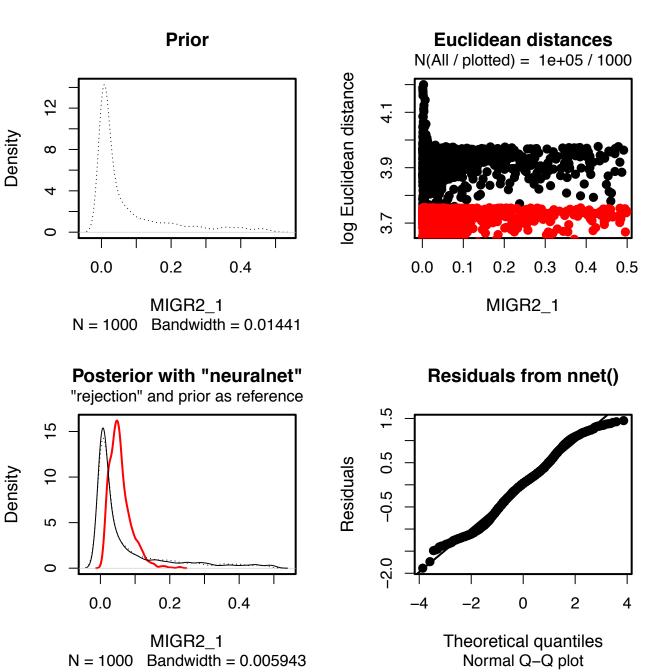
Supplementary File 3 for Carroll et al., Incorporating non-equilibrium situations into demographic history inferences of a migratory marine species

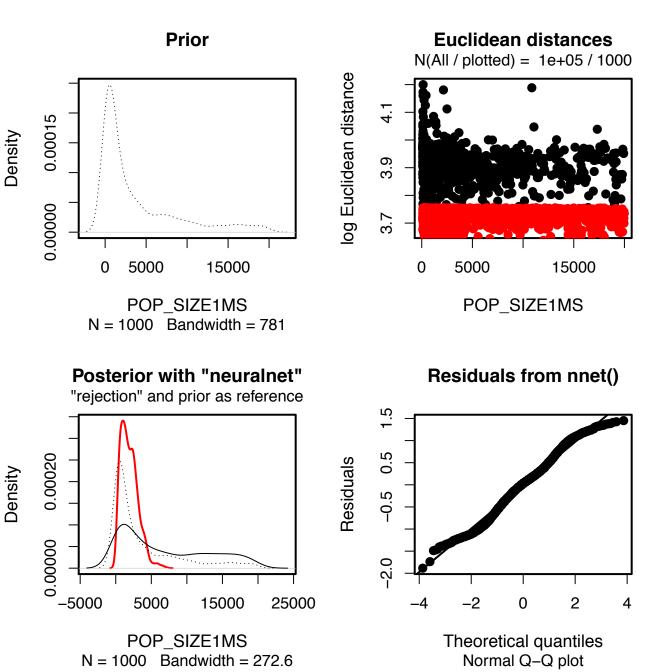
For each scenario, the following plots are displayed for each estimated parameter:

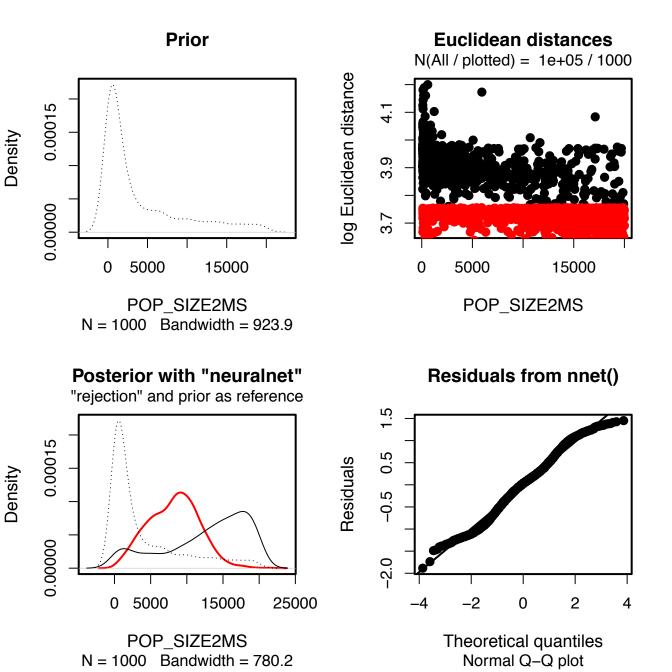
- 1. Top left: a density plot of the prior distribution
- 2. Bottom left: a density plot of the posterior distribution For the density plot of the posterior distribution, values estimated with the non-linear regression correction method or 'neural net' (red thick lines) and, for reference, using the simple rejection method (black fine lines) are displayed. The prior distribution (in the posterior distributions' range) is also displayed (dashed lines). For the scatter plot Euclidean
- 3. Top right: a scatter plot of the Euclidean distances as a function of the parameter values
- 4. Bottom right: Normal Q-Q plot of the residuals from the regression. Points corresponding to the accepted simulations are displayed in red.

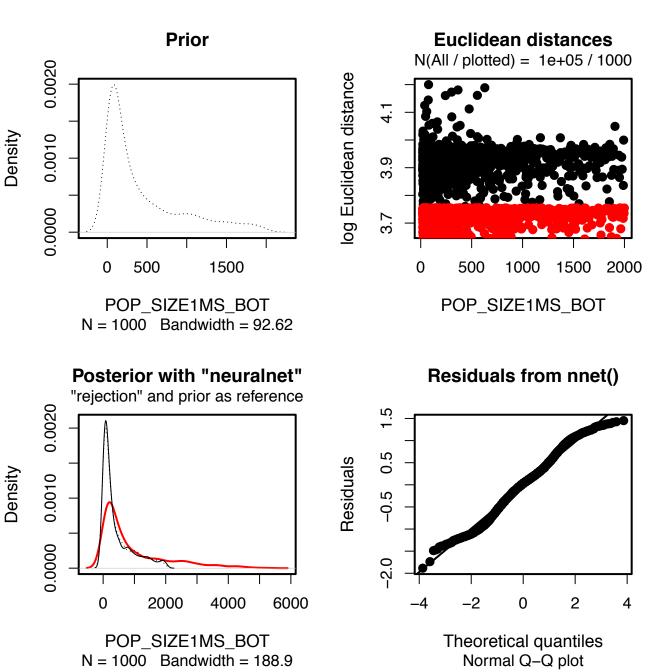
Scenario 1: random lineage sorting with continuous gene flow and a single migration rate M_H

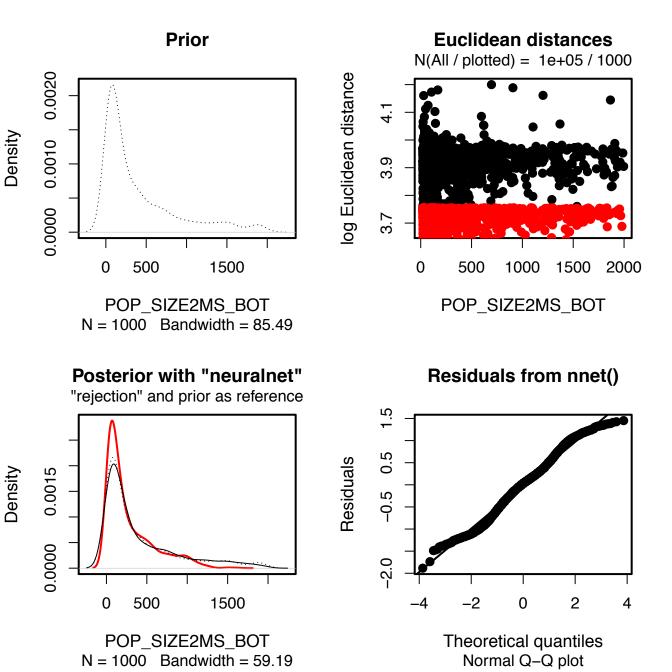


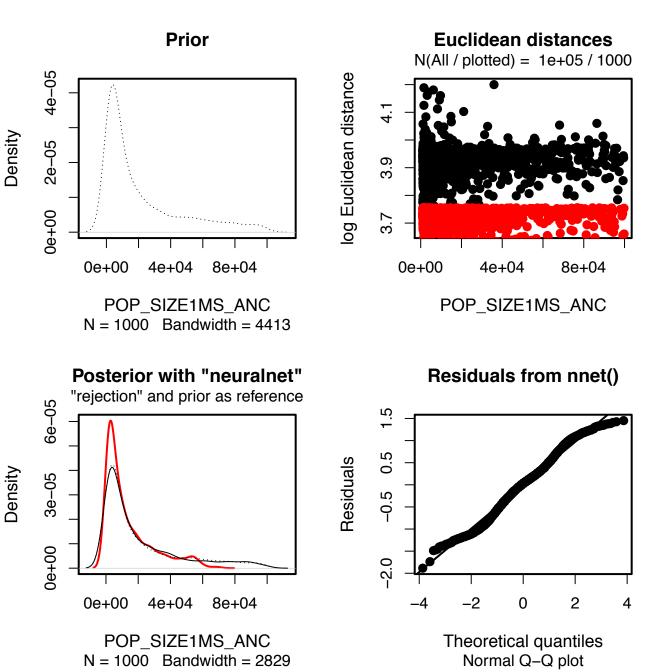


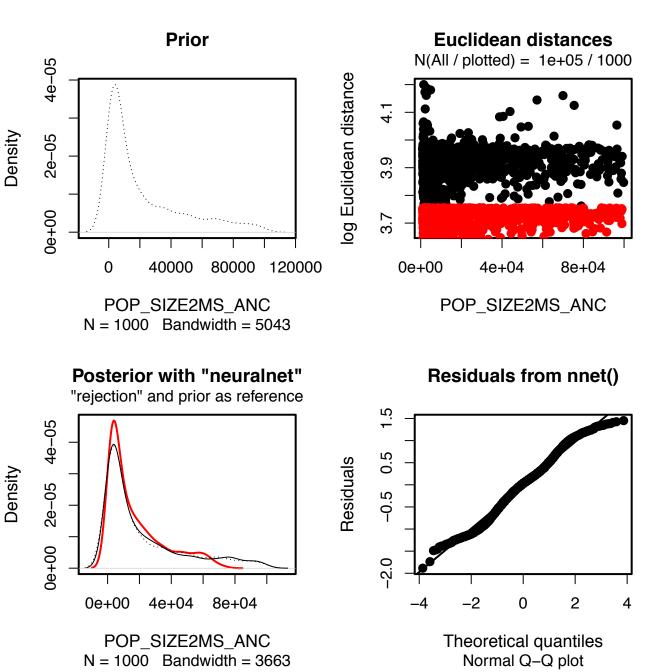


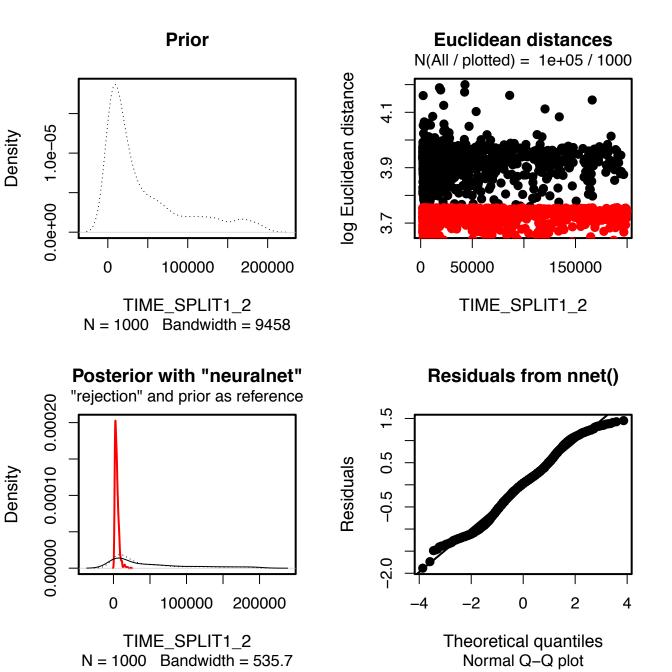




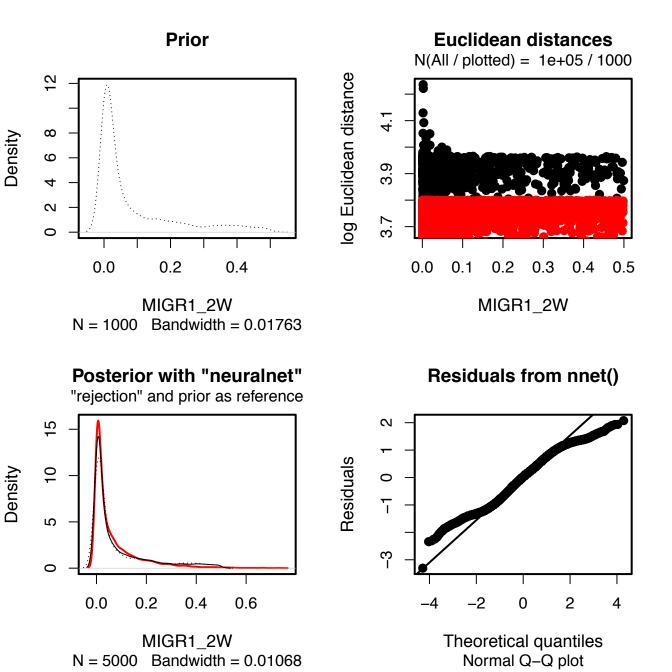


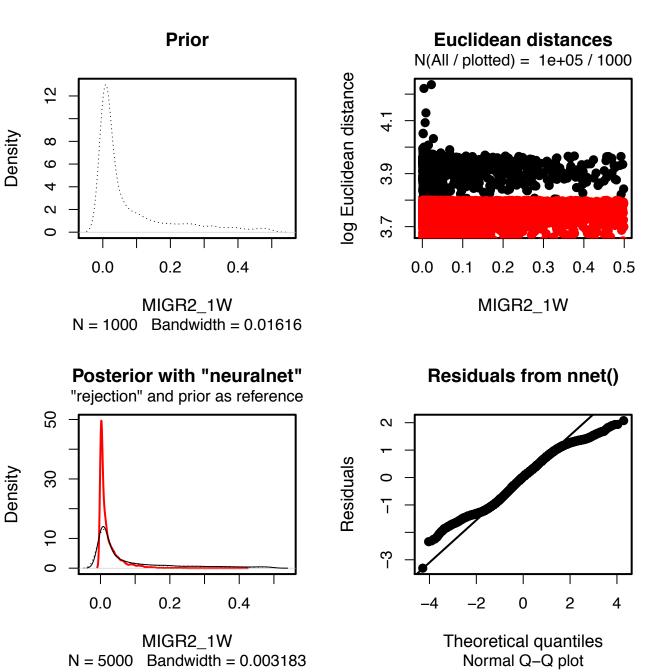


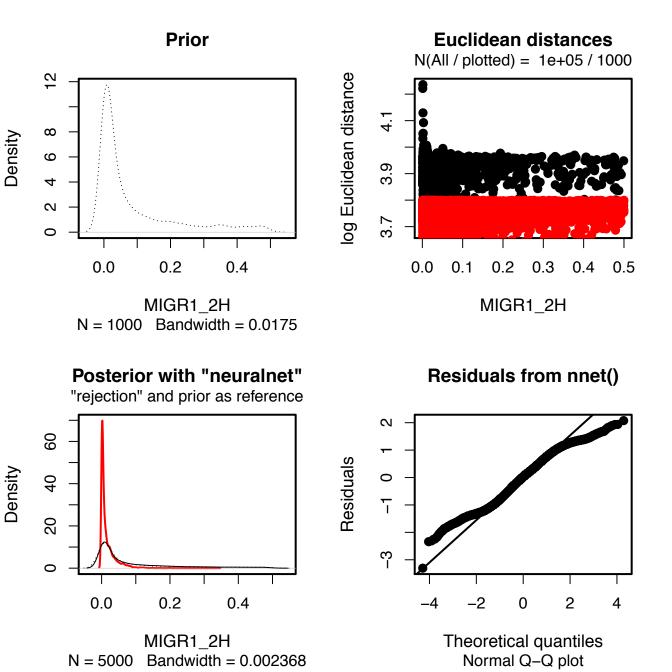


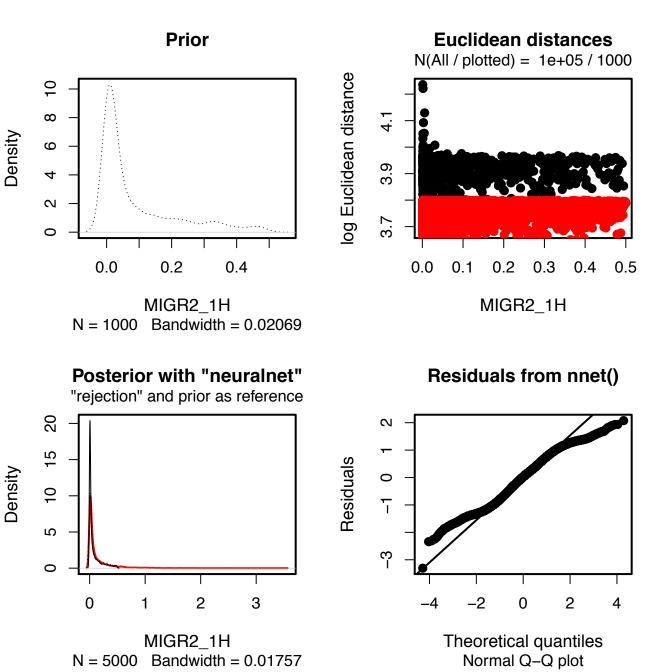


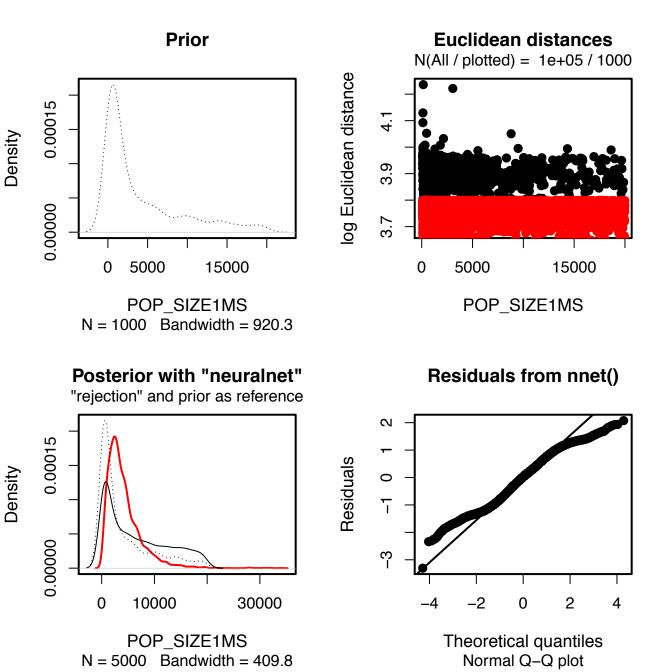
Scenario 2: random lineage sorting with continuous gene flow and two migration rates: one since divergence M_H and one since the whaling era M_W

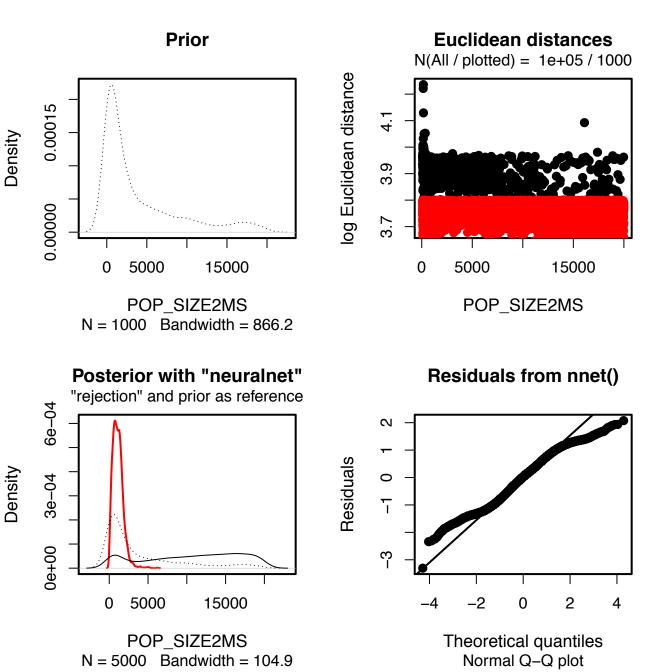


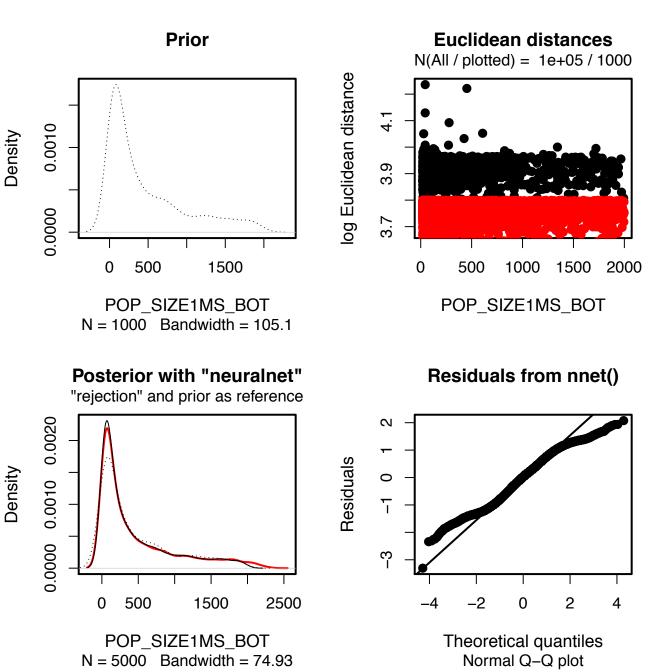


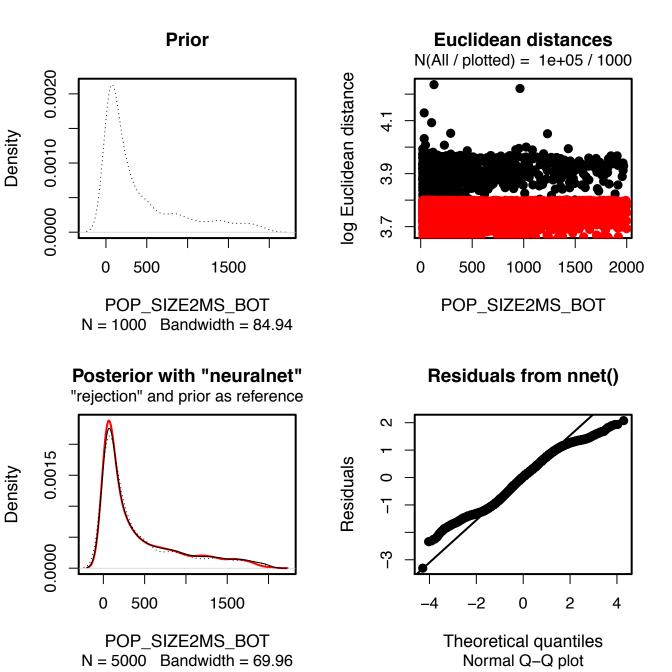


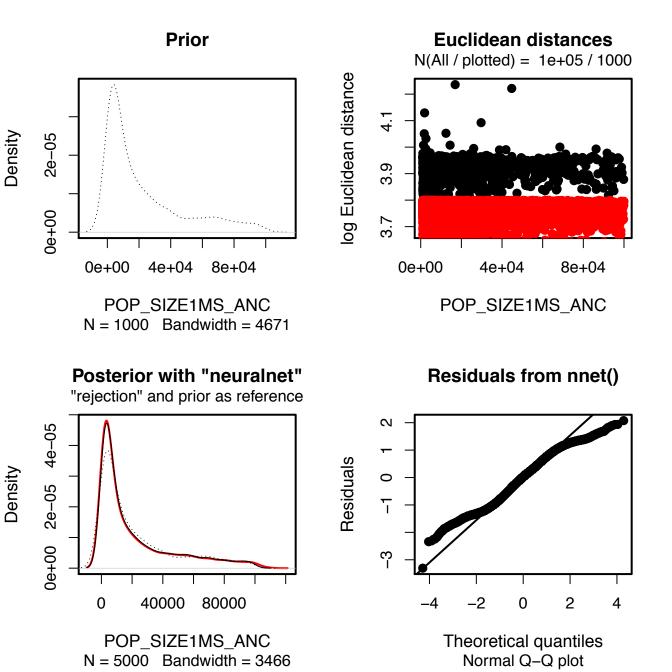


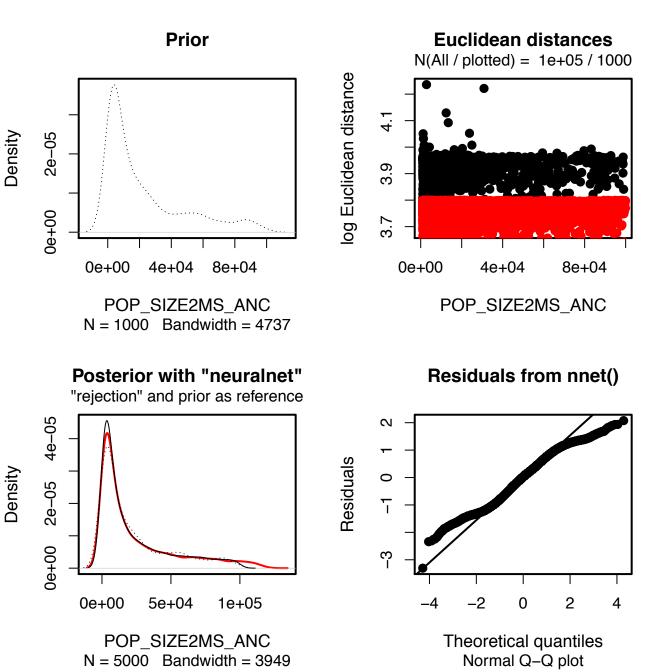


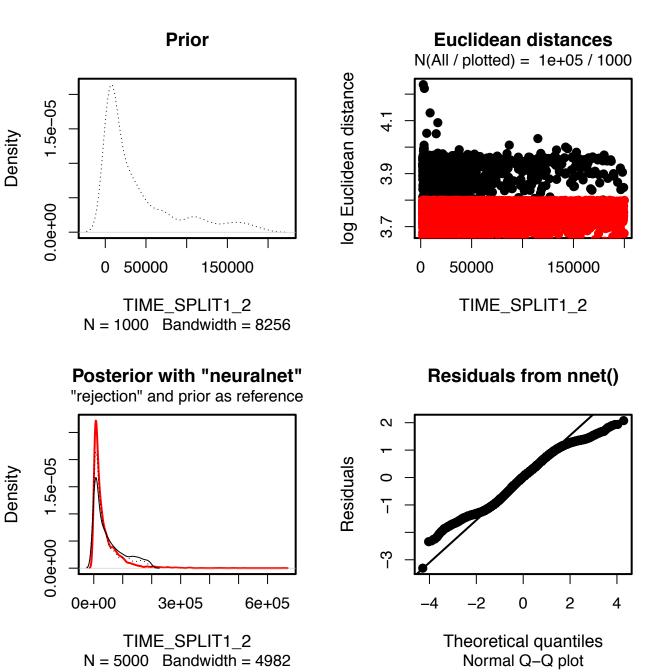




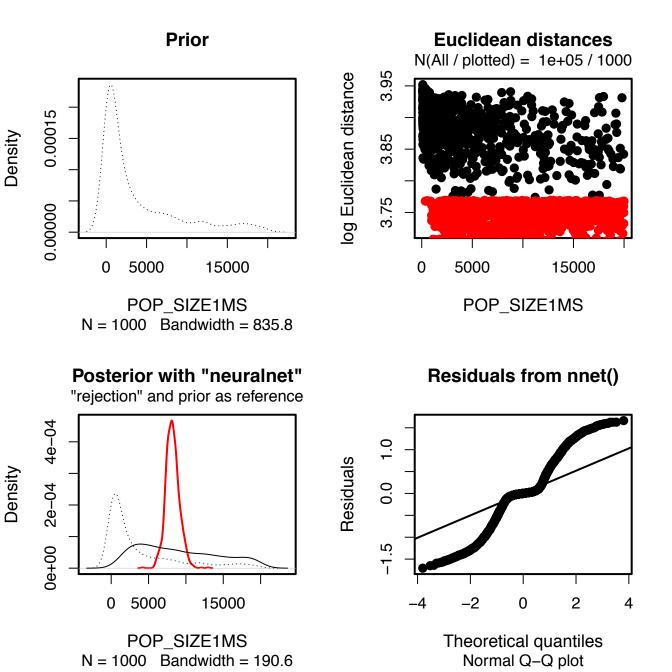


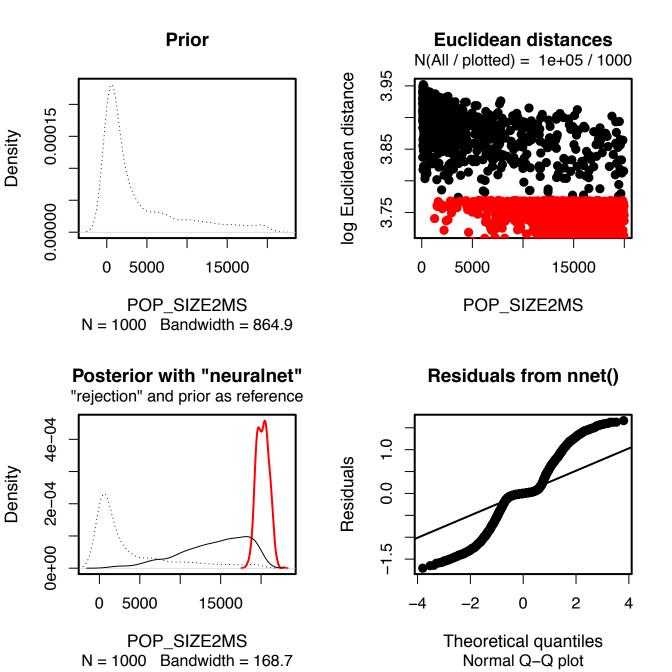


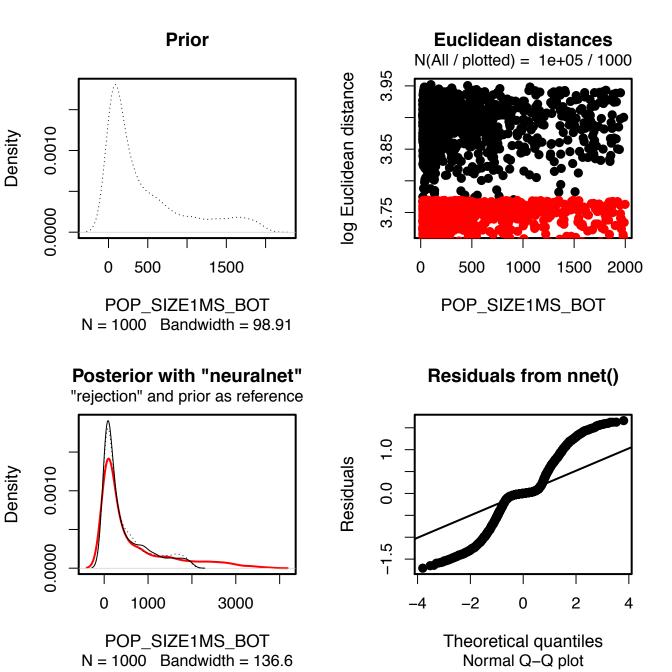


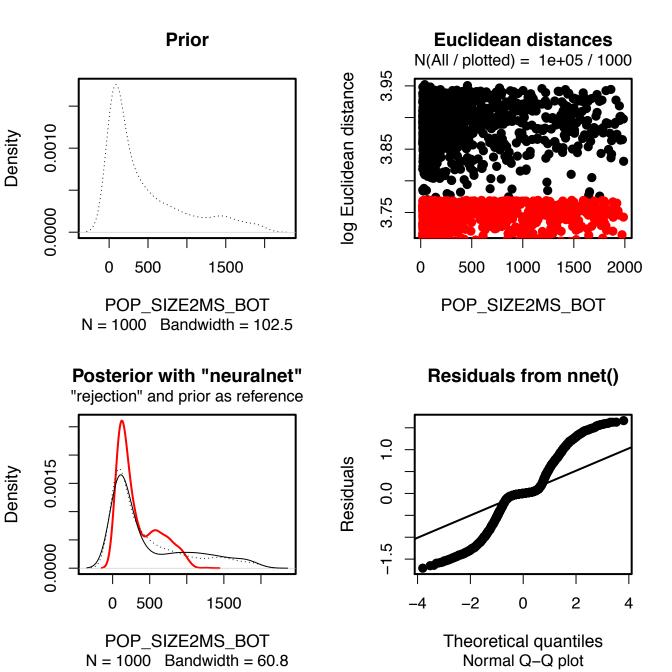


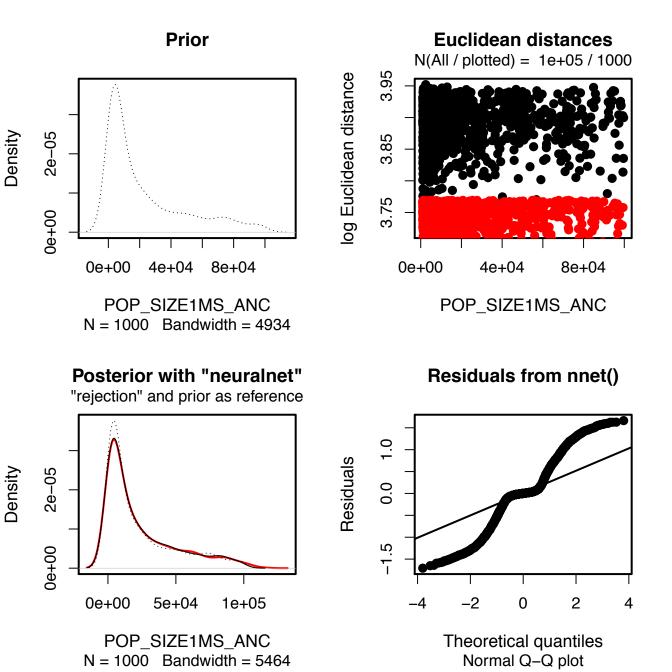
Scenario 3: isolation following divergence, with no subsequent gene flow

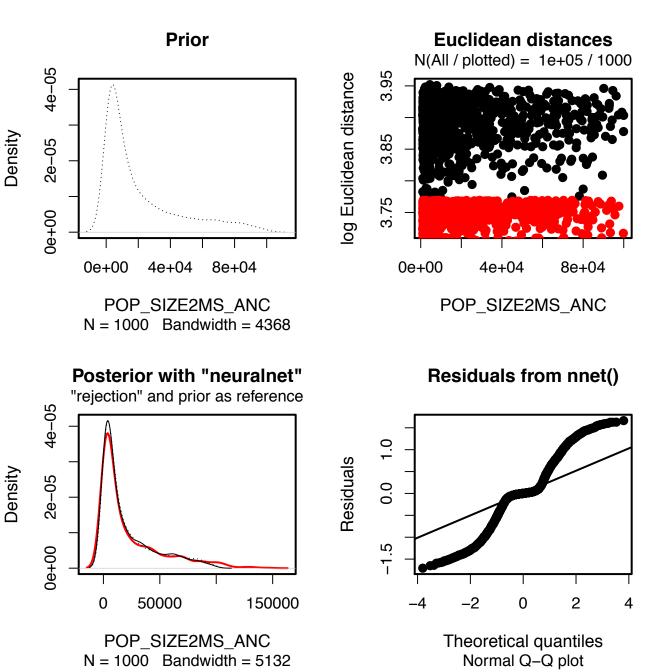


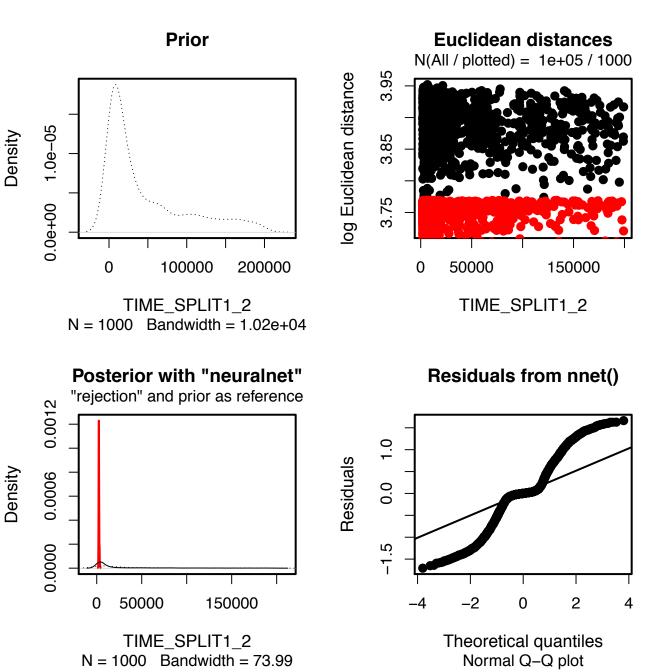




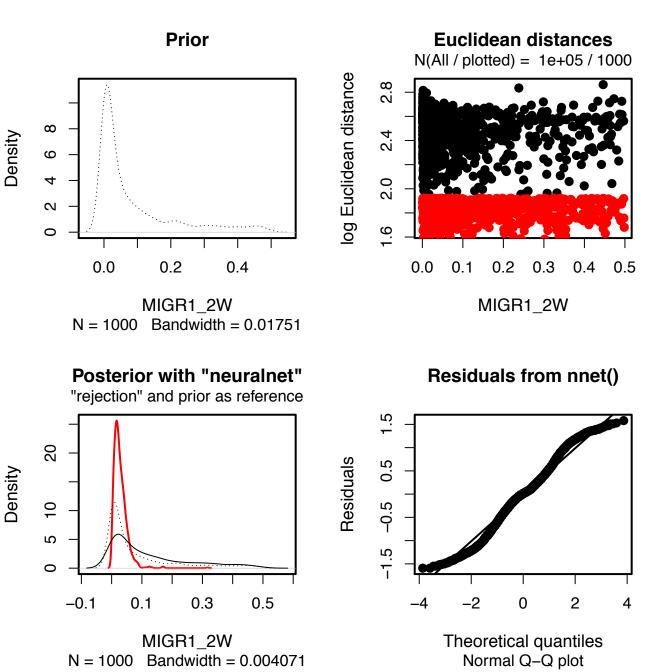


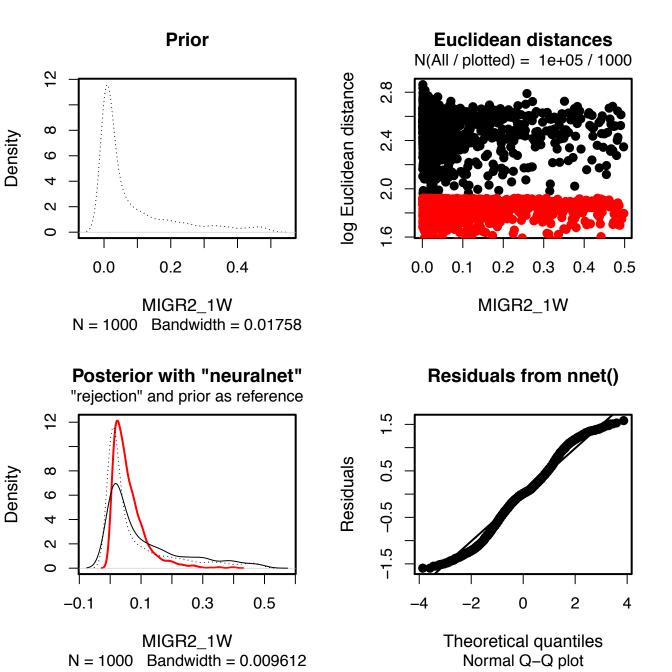


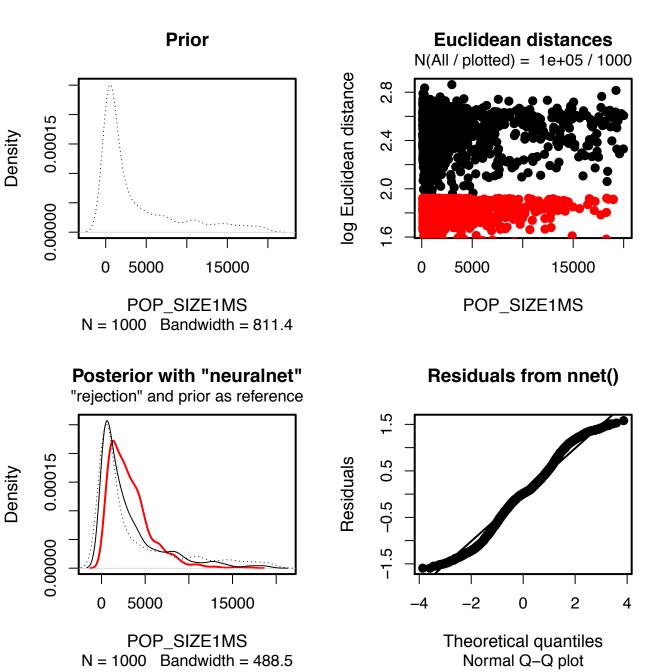


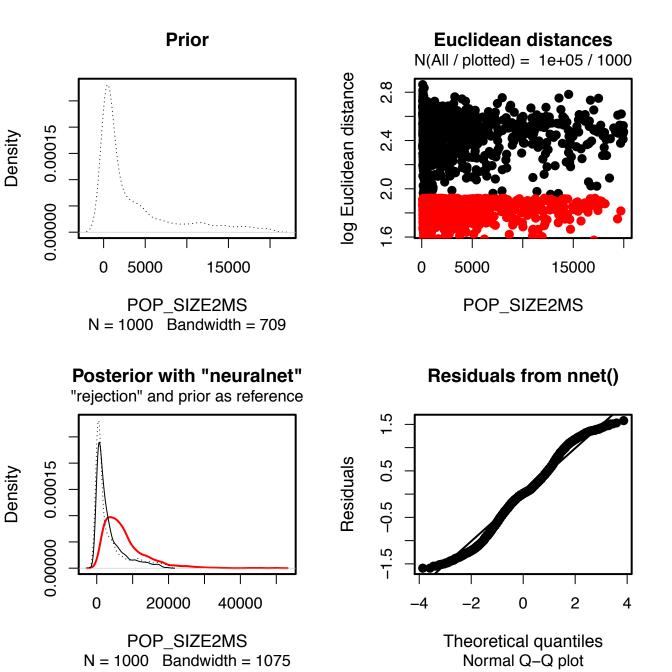


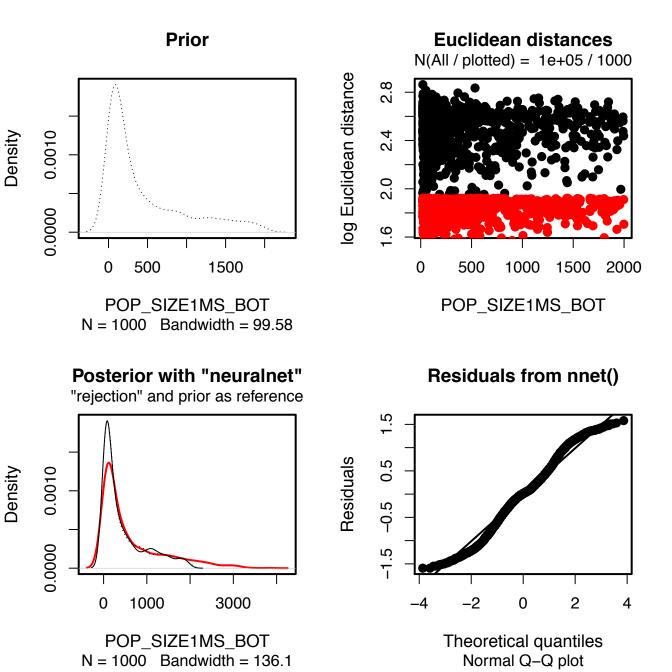
Scenario 4: isolation following divergence, with gene flow with one migration rate since the whaling era M_W

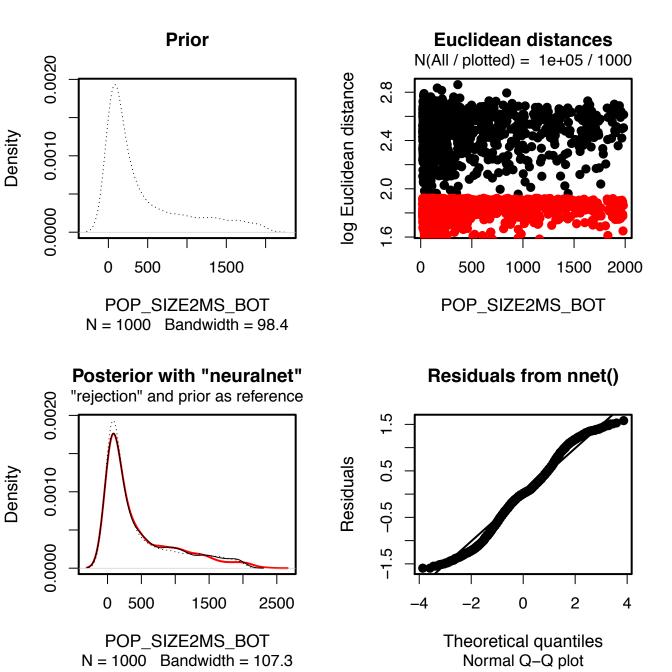


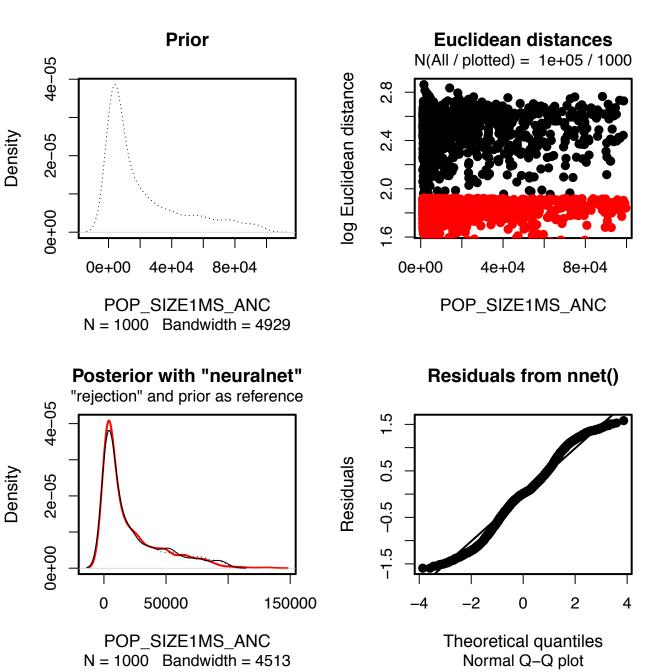


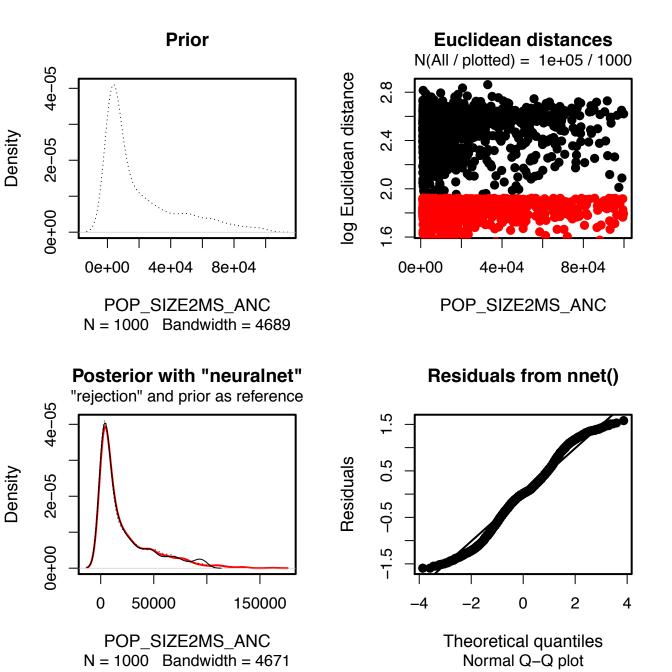


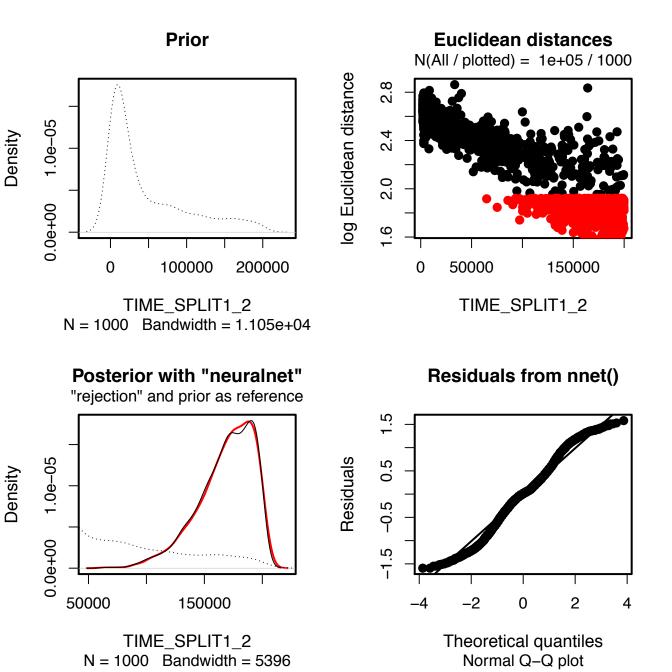




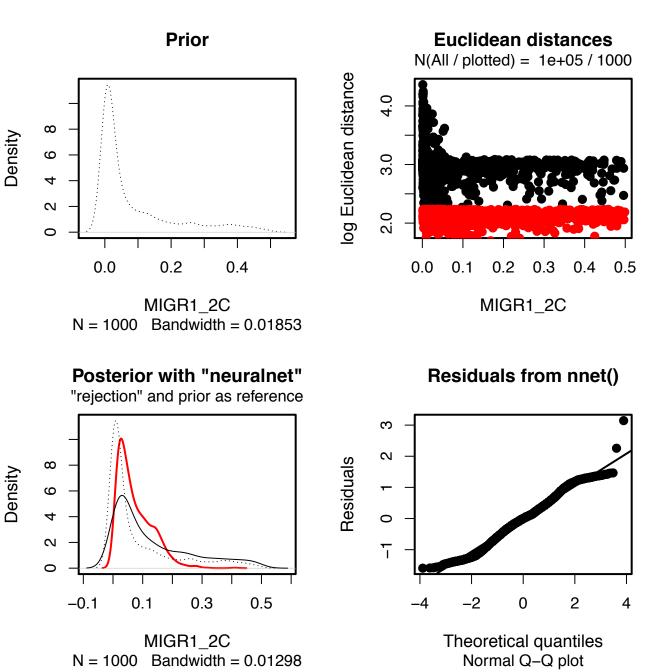


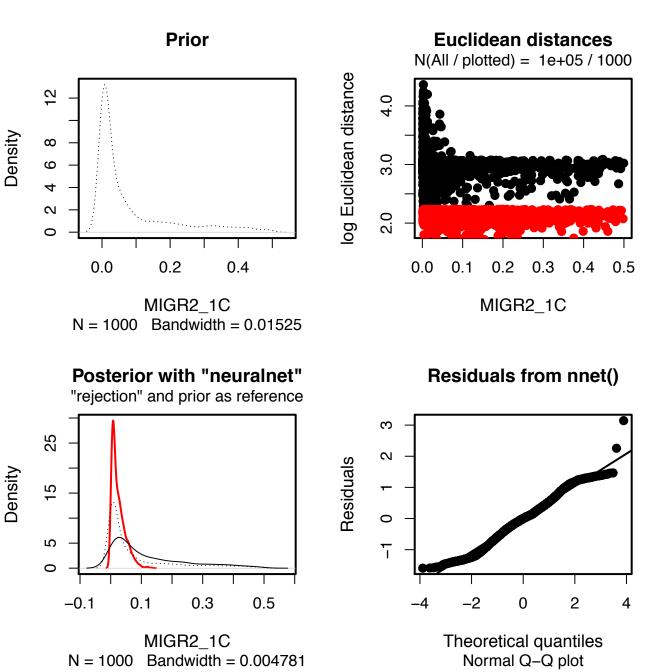


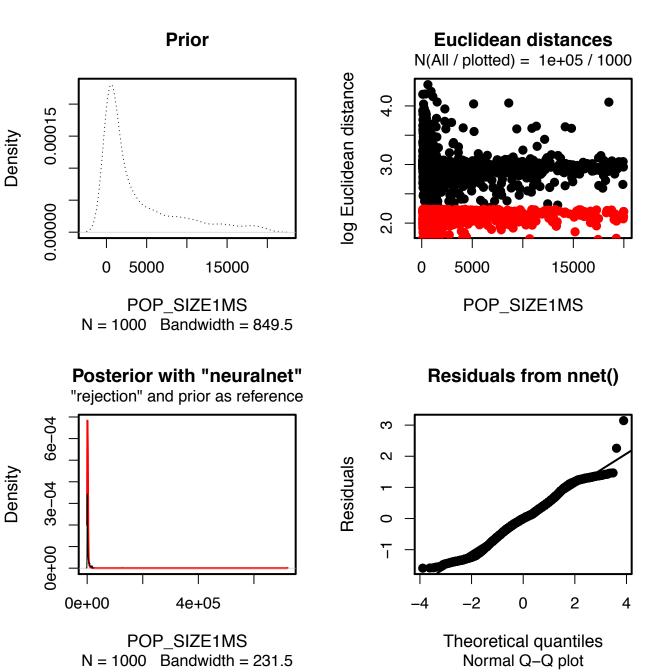


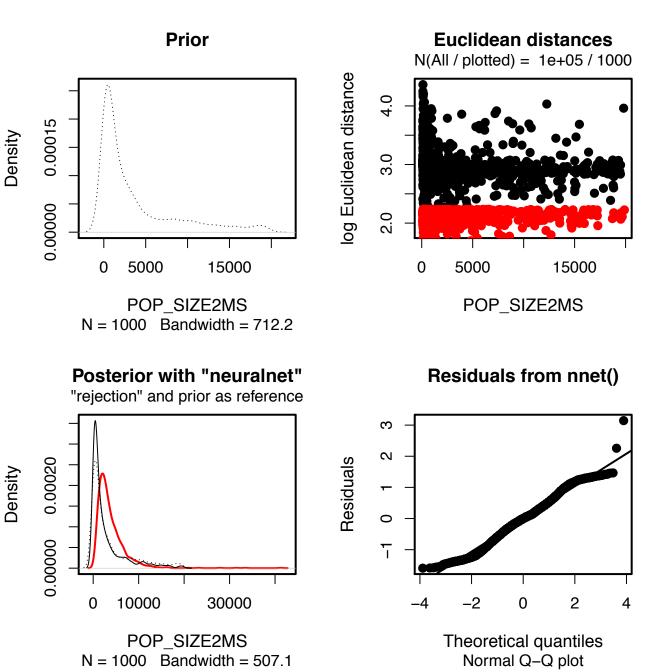


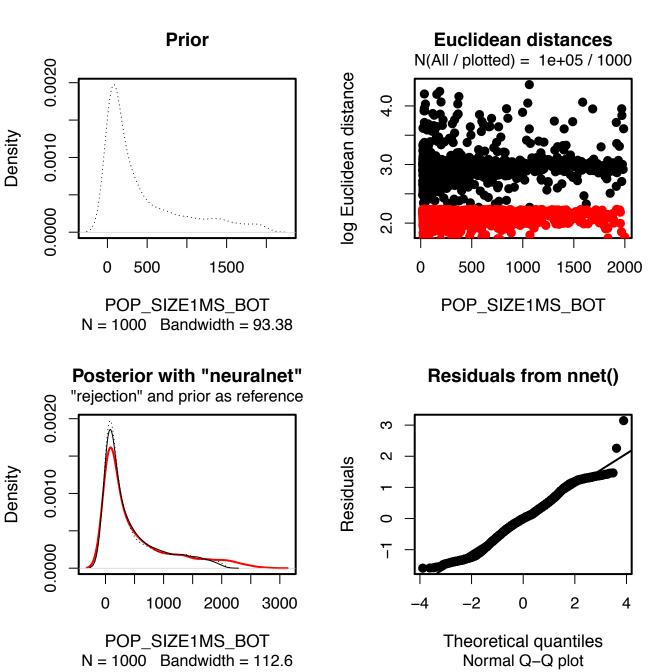
Scenario 5: isolation following divergence, with one migration rate since secondary contact, M_C

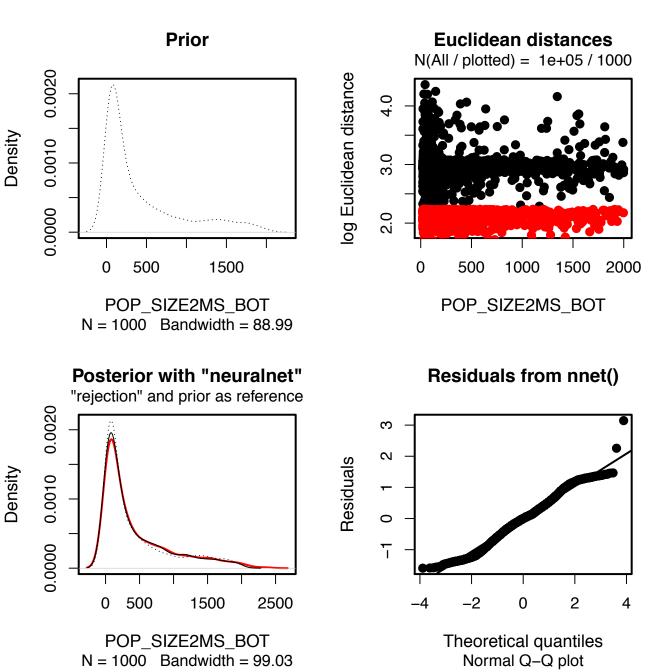


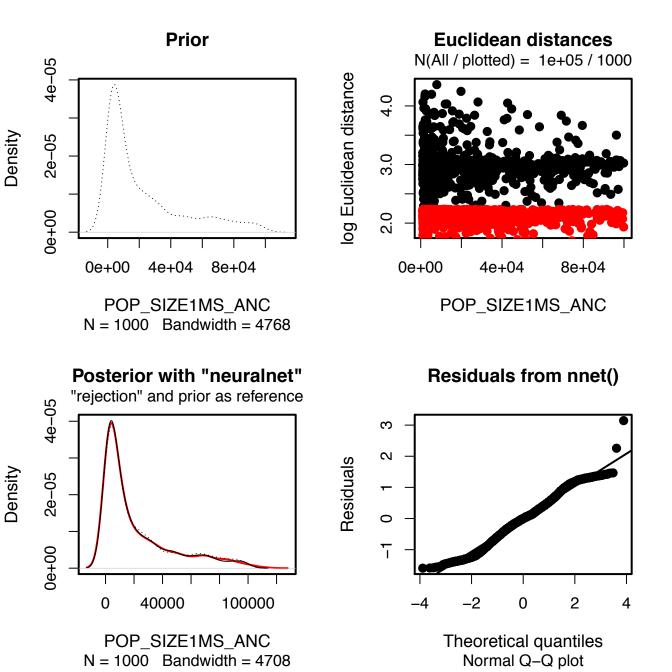


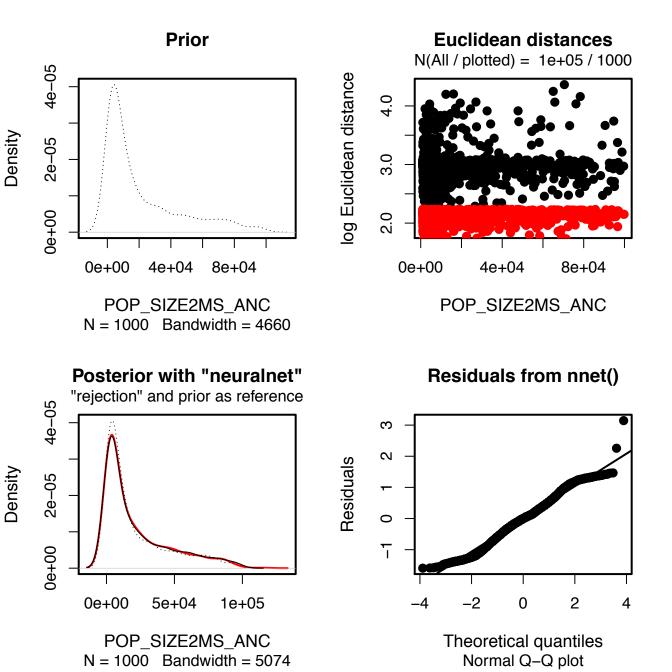


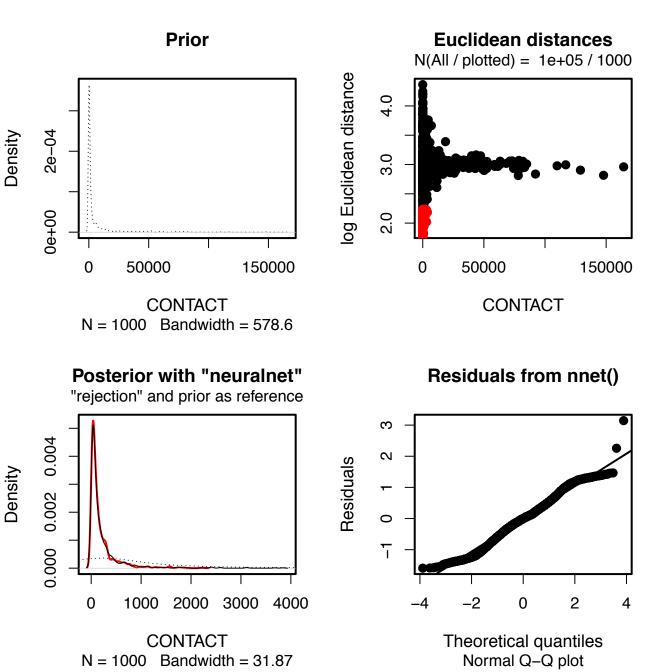


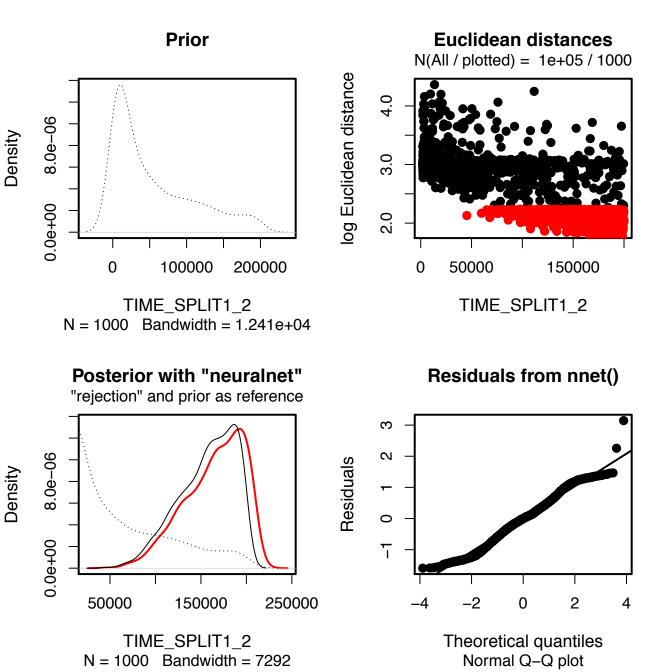




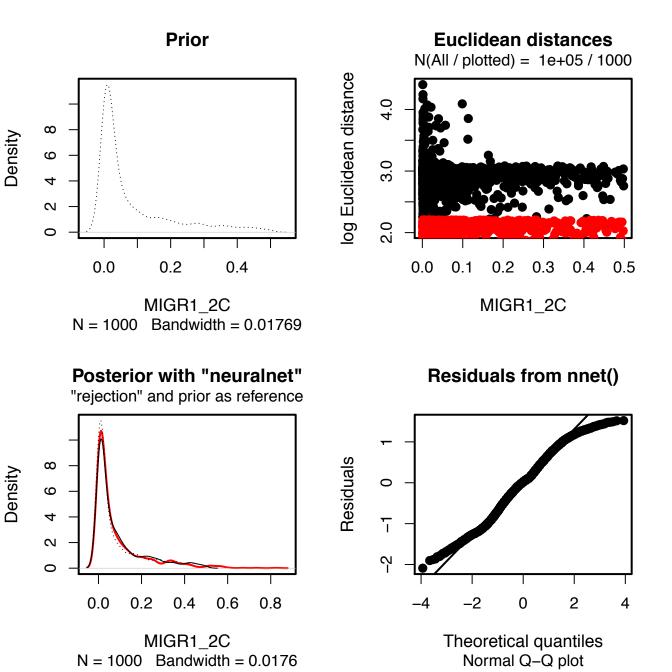


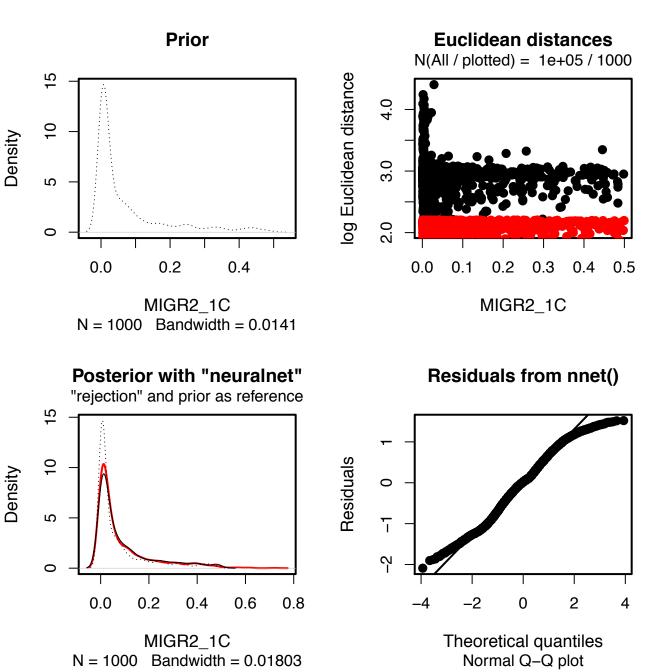


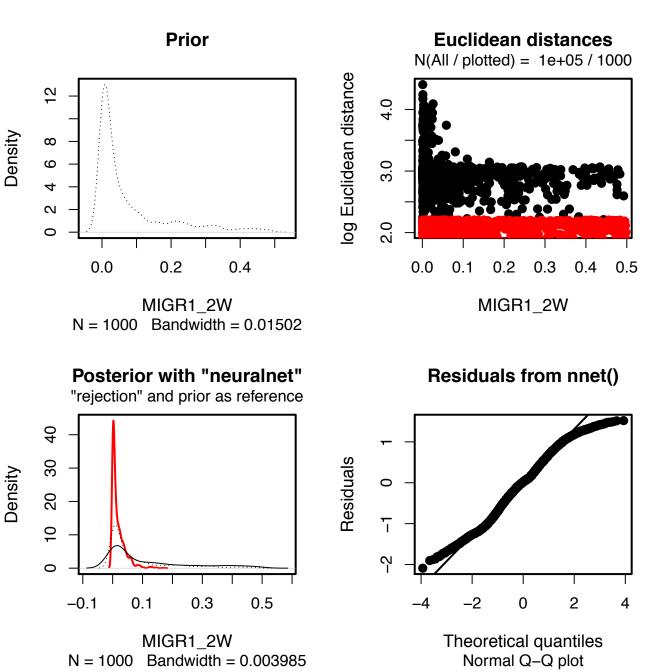


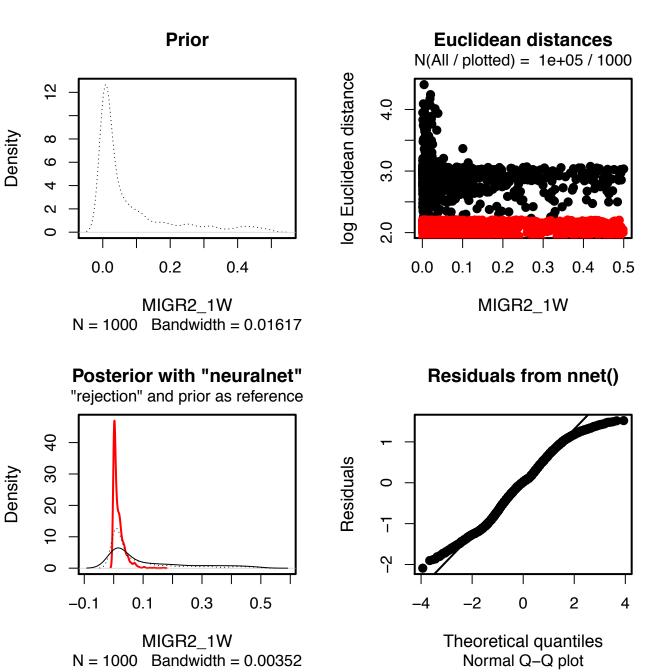


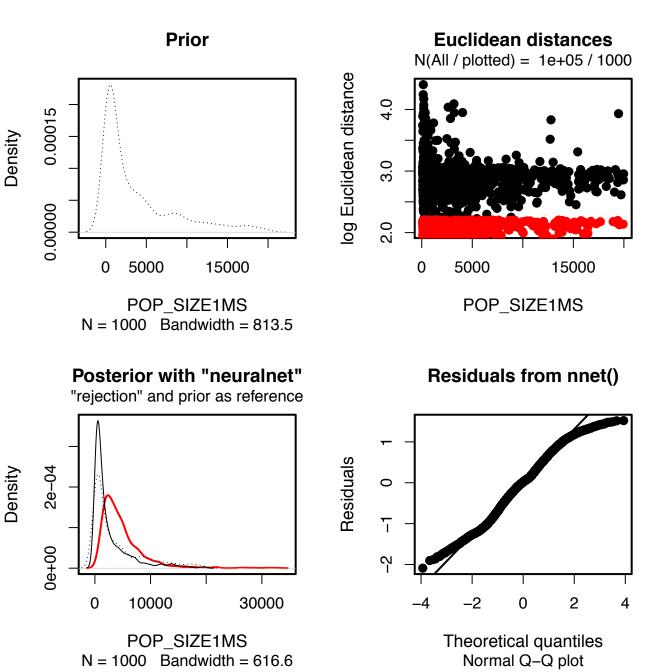
Scenario 6: isolation following divergence, with two migration rates: one since secondary contact, M_C , and one since the whaling era, M_W

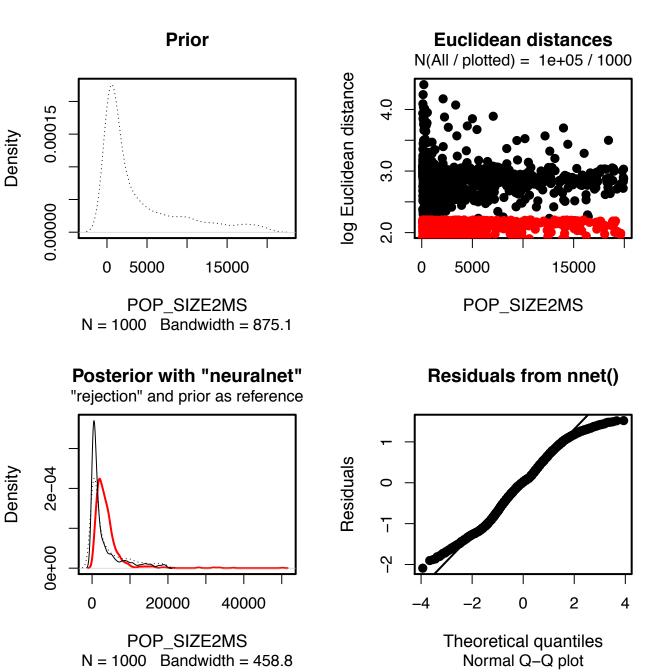


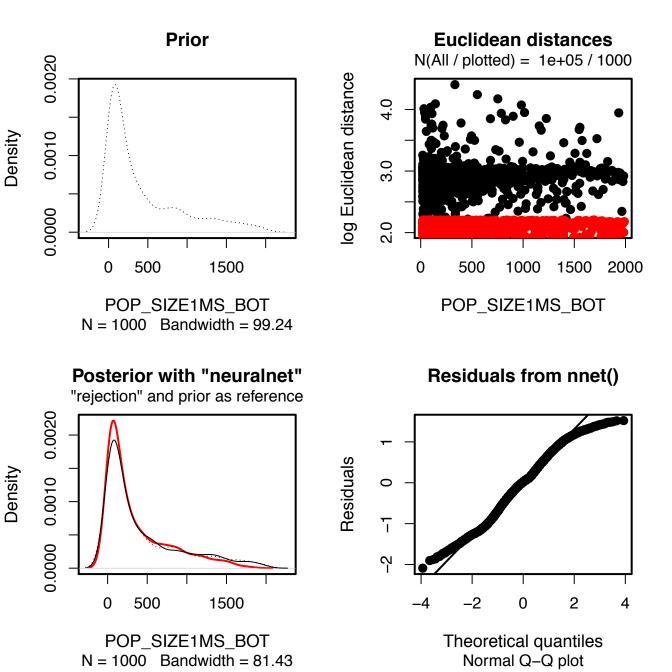


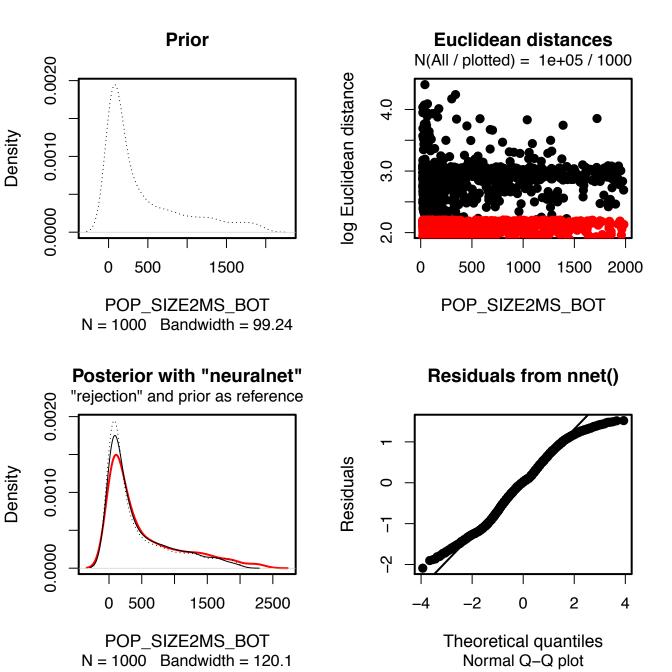


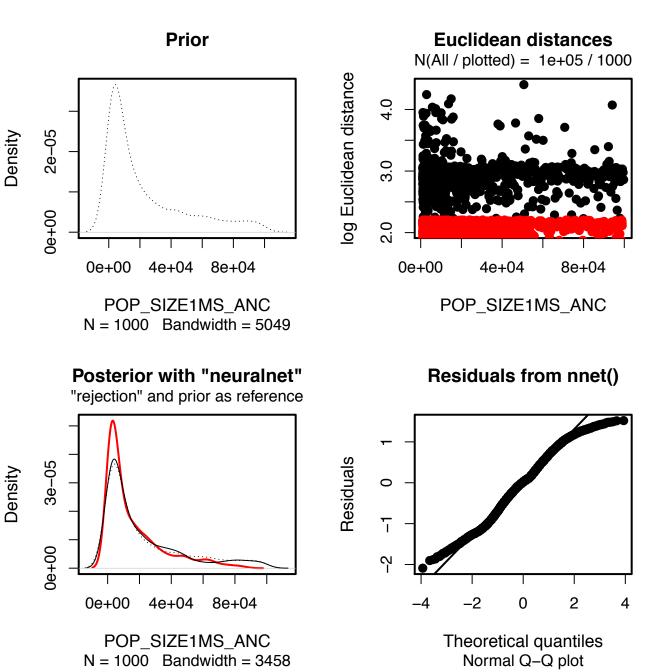


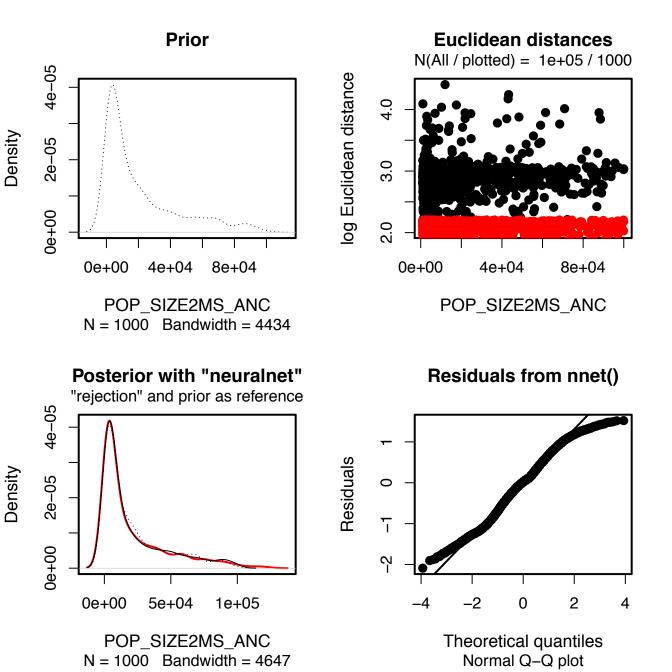


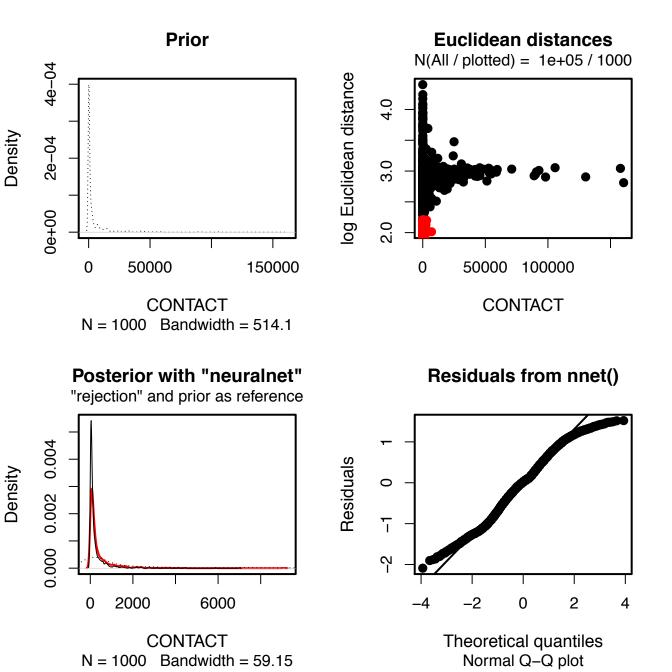


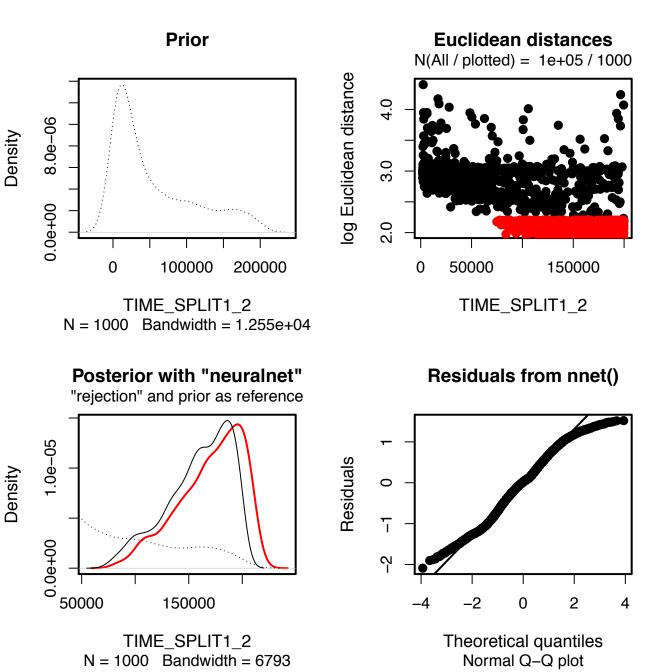






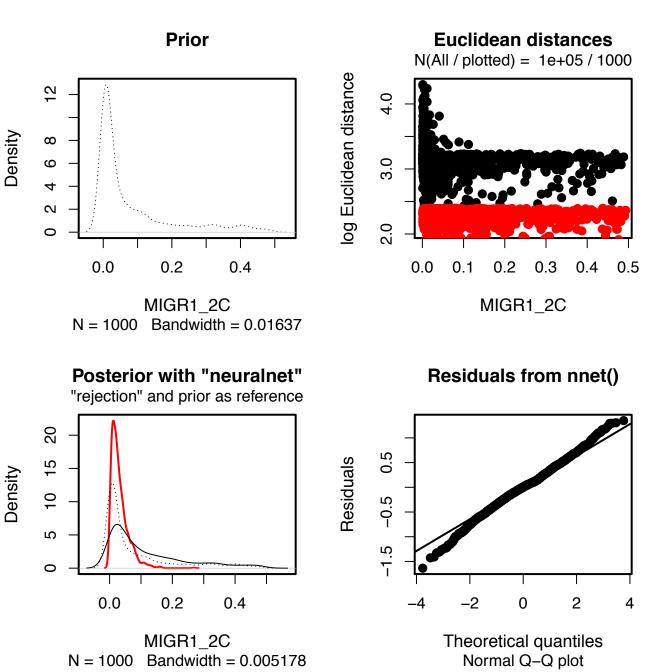


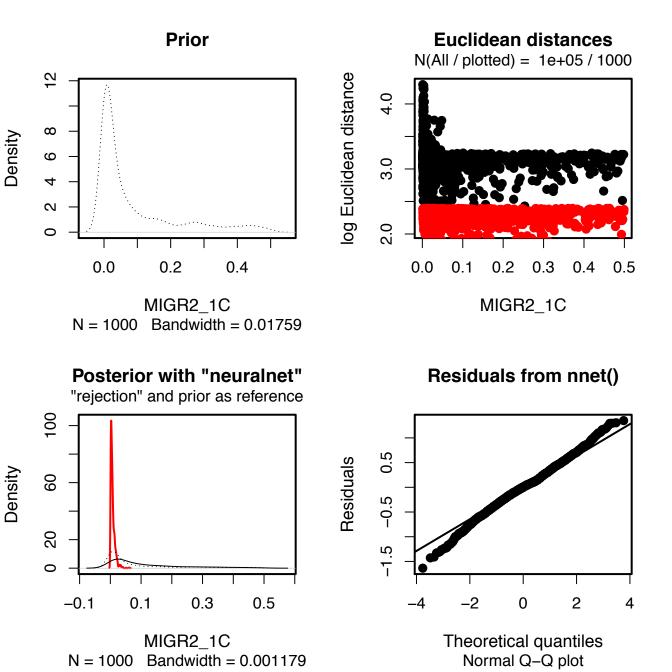


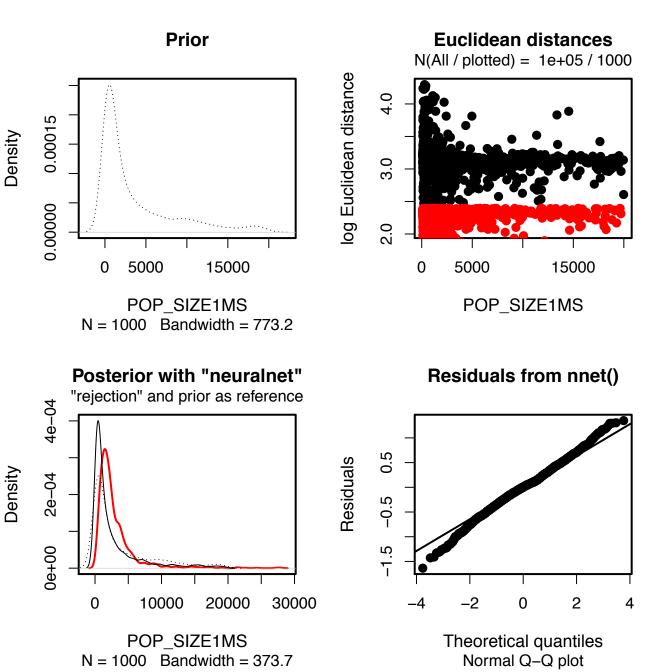


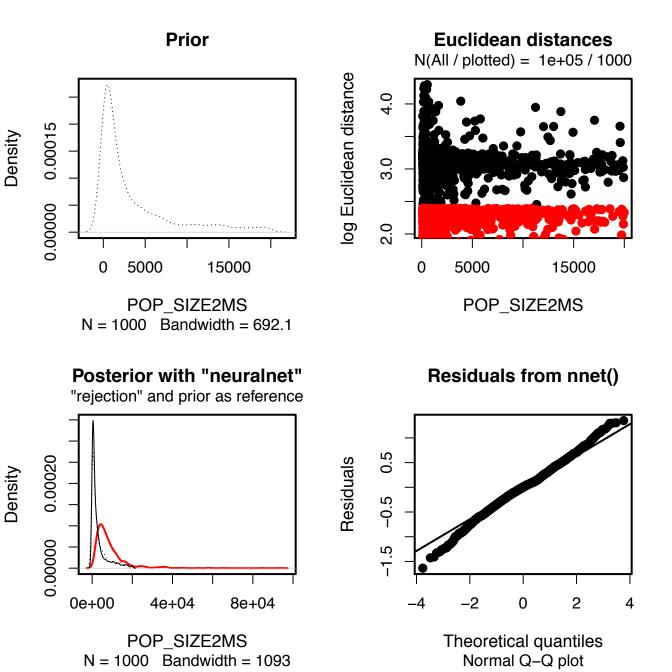
Scenario 5NB: isolation following divergence, with one migration rate since secondary contact, M_C

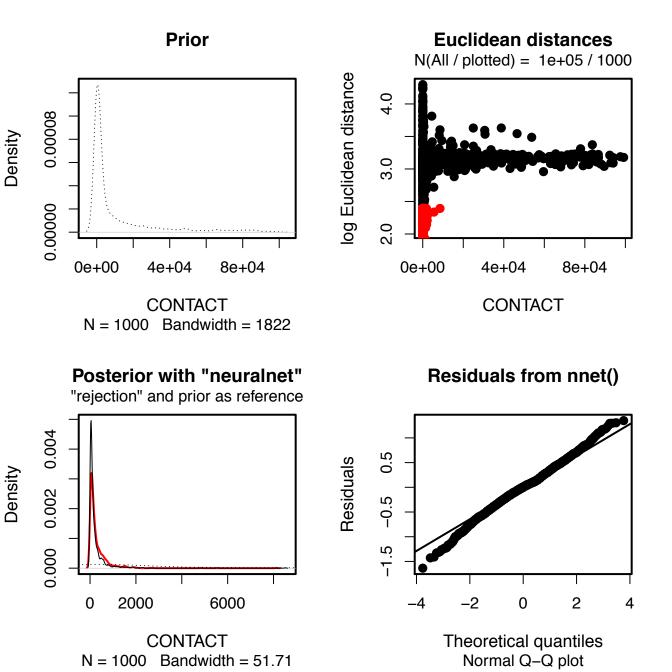
No whaling bottleneck

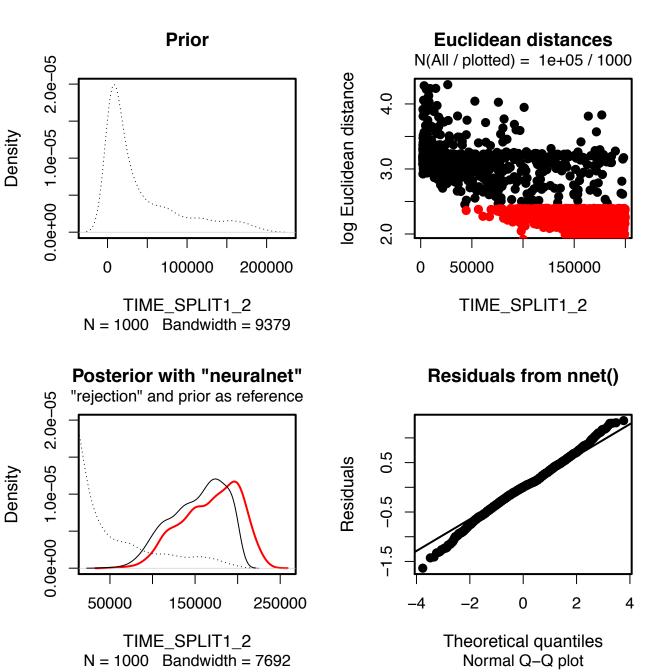












Scenario 6NB: isolation following divergence, with two migration rates: one since secondary contact, M_C , and one since the whaling era, M_W

No whaling bottleneck

