Supplementary Figure 2

Bayesian 50% majority-rule consensus phylogenetic tree illustrating the relationships among MHC DQB/DRB from New Zealand sea lion, and other pinnipeds: Phocarctos hookeri, red; Zalophus wollebaeki, blue; Zalophus californianus, grey; Halichoerus grypus, green; Mirounga leonina and Mirounga angustirostris, black; Arctocephalus gazella, purple) . The tree is unrooted, but for illustrative purposes is rooted by midpoint. Details of the sequences used and their accession numbers are provided in the Supplementary Table 5. Bayesian posterior probabilities are shown for each node.