

Figure S1. Neighbour-joining tree for *Rana arvalis* MHC II exon 2 representing the 57 different alleles. Alleles present in the north are marked with an orange triangle and shared alleles between the northern and southern cluster are represented with a green rhombus. A natterjack toad sequence [Genbank HQ388291.1] from MHC II exon 2 was used as an outgroup. The scale bar represents substitution per site.

