

**Fig S2: Bayesian Maximum clade credibility phylogeny of 400bp nucleoprotein gene from RABVs from Turkey and neighbouring countries in the Middle East.** The defined Middle East clades are labelled according to previous studies (Horton *et al* 2015), unlabelled viruses included to add context and include viruses from Europe and Africa. Posterior probabilities are indicated at each node. Turkey sequences are predominantly within clade C and are enlarged in the inset box.

