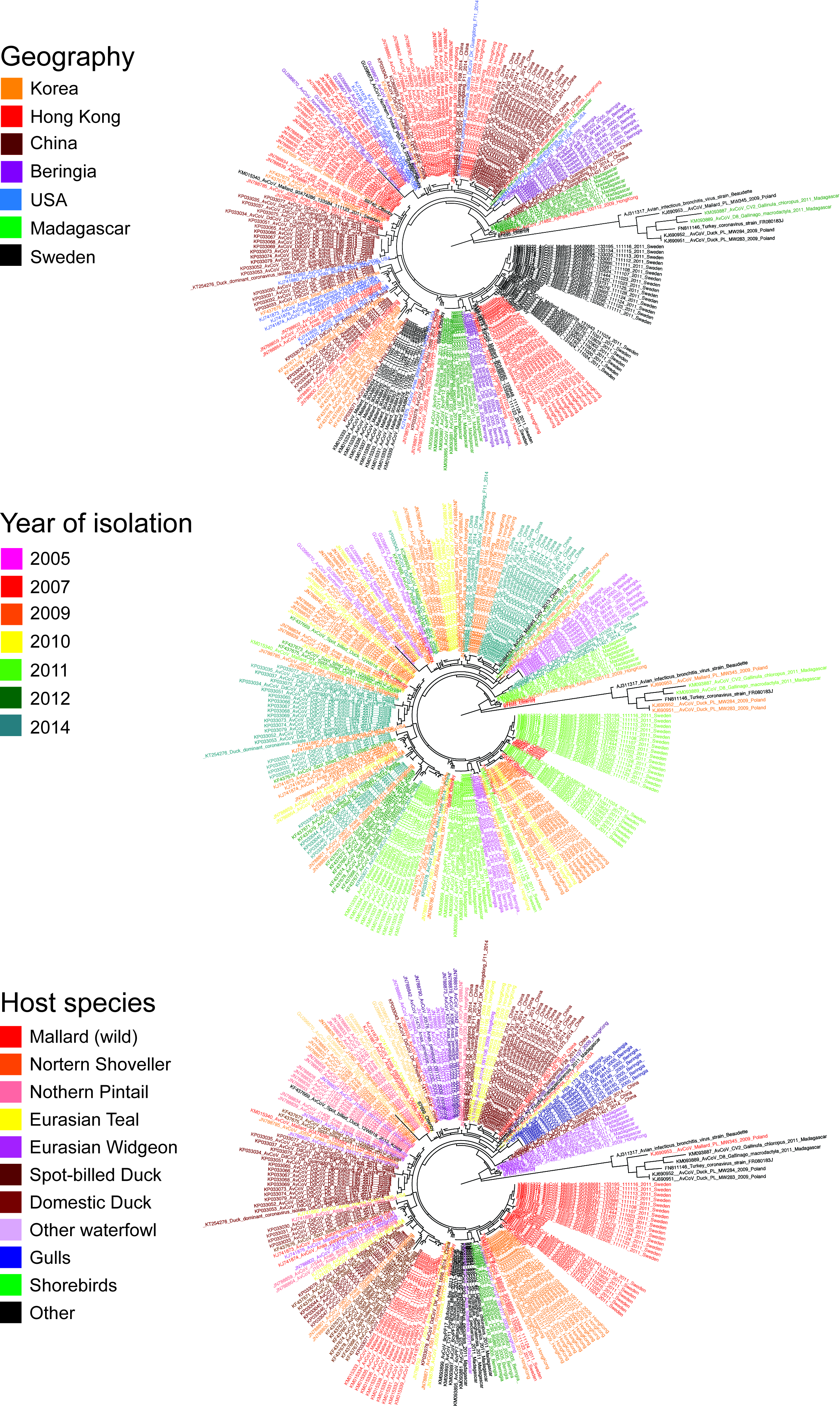
**High prevalence and putative lineage maintenance of avian coronaviruses in Scandinavian waterfowl**

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**S1 Fig**



S1 Fig: Avian coronavirus RNA-dependant RNA polymerase phylogeny shows no spatial, temporal or host species patterns. Sequence metadata are provided in Table S2