

Reassortment of American and Eurasian genes in an influenza A virus isolated from a Great Black-backed Gull (*Larus marinus*), a species demonstrated to move between these regions

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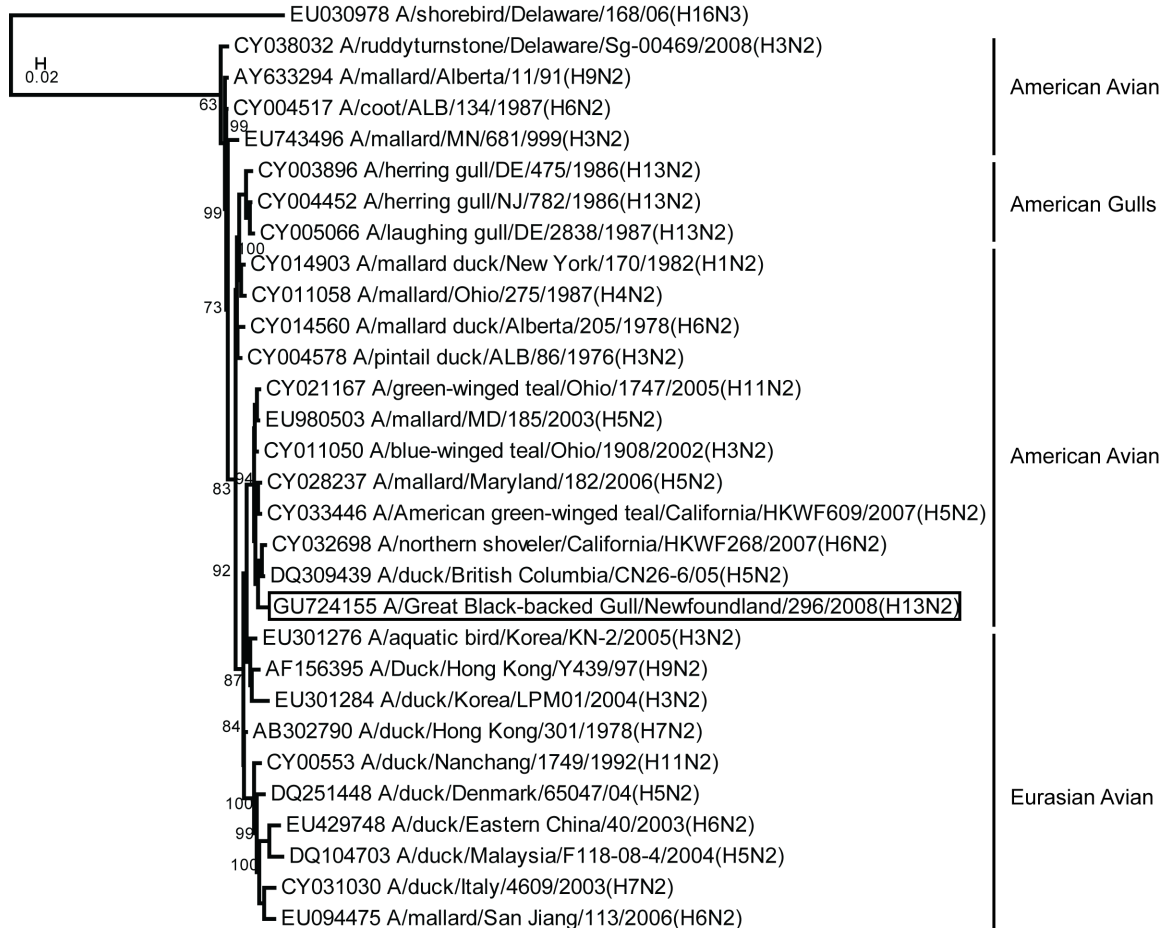
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Note: Nucleotide sequence data reported are available in the GenBank database under the accession numbers: GU724150 – GU72415



Online Resource 3: Phylogenetic analysis of the N2 neuraminidase gene of A/Great Black-backed Gull/Newfoundland/296/2008(H13N2). The Bayesian inference tree is rooted with A/influenza/shorebird/Delaware/168/06(H16N3). A/Great Black-backed Gull/Newfoundland/296/2008(H13N2) is indicated by a box. Bayesian posterior probabilities are indicated as percentages. The scale bar indicates the number of nucleotide substitutions per site. Sequences from the American avian, American gull and Eurasian avian clades are indicated.