





Phylogenetic Network of Influenza A H1N1 pdm
By Network ver. 5.0.0.0

Sample Information

2009	2010	2011	2012	2013	2014	2015	2016	2017	Total
2033	736	550	323	478	260	164	464	57	5065

*All samples(FASTA format) were attained from IRD(<https://www.fludb.org/brc/home.spg?decorator=influenza>).

*Identical sequences were excluded using the option, “collapse identical sequences” from the database.

*Only the full-length of HA samples were included(length= 566).

*The samples attained were aligned by DNA Alignment version 1.3.3.2 (Fluxus Technology Ltd.)