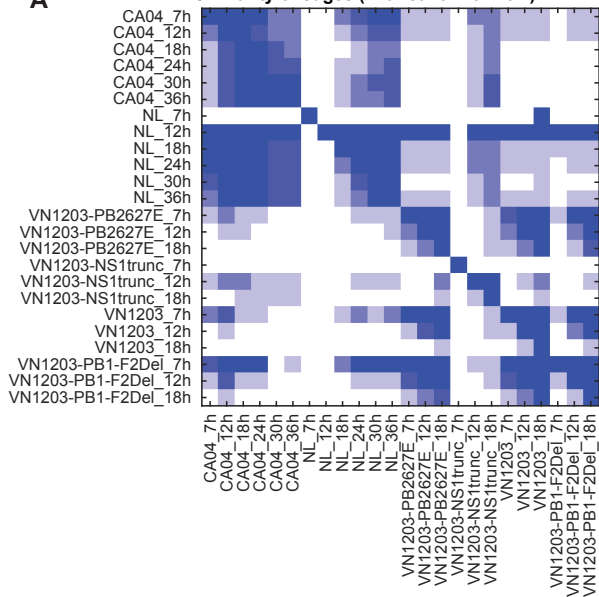


A**Similarity of edges (wrt network on row)**

Precision
|intersection|/|row network|

1 0.9 0.8 0.7 0.6 0.5 0.4 0.3 0.2 0.1 0

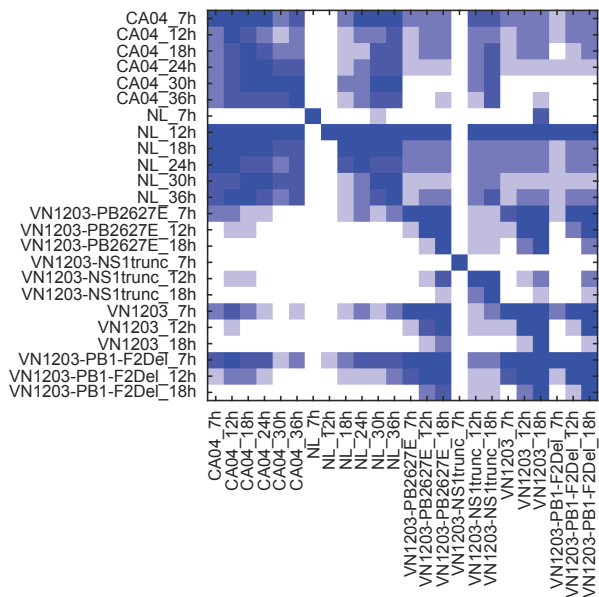


Low pathogenicity: CA04, NL (both H1N1)

H5N1 and mutants:

Medium pathogenicity: PB2-627E, NS1trunc

High pathogenicity: PB1-F2del, VN1203

B**Similarity of regulators (wrt network on row)****C****Similarity of targets (wrt network on row)**