1   **Supplementary Information S1**   BaseSpace Correlation Engine. This Microsoft Excel file

2   contains five spreadsheets; Combined, Canonical Pathways, Gene Ontology, Protein Families

3   and Regulatory Motifs. Within each of the Canonical Pathways, Gene Ontology, Protein

4   Families and Regulatory Motifs spreadsheets are the data generated by the BaseSpace

5   Correlation Engine analyses of differentially expressed genes at 36hpi with pOka, gB[Y881F],

6   gB[Y920F] and gB[Y881/920F] or the genes associated with the wild type-like or hyperfusogen

7   phenotypes. The combined spreadsheet compiles all of the analyses into a single table.