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09.10.2016 Data from Confirmed Cases of Influenza A (H1N1) in Australia, 2009-2011 Contains information about the type, number and age distribution of cases, date of onset, and date of death. Where the age of the person at onset of symptoms is known, the median age of symptom onset is provided as standard unit age (i.e. from 0 to 44 years). Other data includes: the type of specimen tested (i.e. throat swab, nasopharyngeal swab, or urine), number of days from onset of illness to specimen collection, and death date. Systematic Study of the Proteins of Influenza Virus - Influenza A (H3N2) The Influenza A Virus is a member of the Orthomyxoviridae family. The Influenza A virus has a segmented single strand negative sense RNA genome. There are 3 types of Influenza A viruses: influenza A(H1N1), influenza A(H1N1pdm09), and Influenza A(H3N2). The influenza A(H1N1) virus is the most common cause of human influenza and can affect all ages. The influenza A(H3N2) virus has been shown to cause significant morbidity and mortality and is currently predominating in human influenza viruses. The Influenza A(H1N1)pdm09 virus is a reassortant virus with genes from both Influenza A(H1N1) and Influenza A(H3N2). Viral Protein Analysis of Influenza A(H3N2) Virus We performed a systematic analysis of the whole proteome of Influenza A(H3N2) virus using 2D gel electrophoresis. The viral proteins were isolated and separated by two-dimensional electrophoresis. The electrophoretic patterns were recorded with silver staining, analyzed, and compared to the whole proteome of the virus. Ten viral protein spots were differentially expressed in the virus. The protein spots could not be identified by mass spectrometry but there were a few proteins that were identified by mass spectrometry or that had a high probability score. Systematic Study of the Proteins of Influenza A(H3N2) Virus The Influenza A virus is a member of the Orthomyxoviridae family. The Influenza A virus has a segmented

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