

PEMODELAN RAMALAN TREND PENYAKIT INFLUENZA H1N1 MENGUNAKAN MODEL BERSEPADU SEIQRE

(Modelling Predicting of Novel Influenza H1N1 Trend Disease Using Integrated
SEIQRE Model)

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ABSTRACT

The H1N1 influenza outbreak, which initially infected over 15,000 individuals in Malaysia, served as a stark warning to healthcare professionals and all stakeholders in Malaysia and global health sectors. This study proposes SEIQRE model to manage and prevent the spread of influenza through quarantine strategies and natural recovery mechanisms. By employing the Runge-Kutta method, the study provides deeper insights into the dynamics of H1N1 influenza and develops effective strategies to mitigate transmission and reduce risk in high-prevalence areas. Analysis of key epidemiological parameters reveals that reinfection plays a significant role in triggering a backward bifurcation, challenging the conventional reliance on a basic reproduction number less than one. A modified model excluding reinfection confirms a forward bifurcation. These findings underscore the importance of incorporating quarantine, bifurcation effects, and scaling factors into effective infection control strategies. This research contributes to global efforts to combat the H1N1 pandemic and safeguard public health in Malaysia.

Keywords: novel influenza H1N1; disease dynamics; preventive measures; transmission risk; epidemiological parameters; basic reproduction number

ABSTRAK

Wabak influenza H1N1, yang pada mulanya menjangkiti lebih daripada 15,000 individu di Malaysia, bertindak sebagai amaran tegas kepada para profesional kesihatan dan semua pemegang kepentingan dalam sektor kesihatan Malaysia dan global. Kajian ini mencadangkan model SEIQRE untuk mengurus dan mencegah penularan influenza melalui strategi kuarantin dan mekanisme pemulihan semula jadi. Dengan menggunakan kaedah *Runge-Kutta*, kajian ini mampu memperoleh pemahaman yang lebih mendalam tentang dinamik influenza H1N1 dan membangunkan strategi yang berkesan untuk mengurangkan penularan dan mengurangkan risiko di kawasan dengan prevalen tinggi. Analisis parameter epidemiologi utama menunjukkan bahawa jangkitan berulang memainkan peranan penting dalam mencetuskan bifurkasi terbalik, mencabar ketergantungan konvensional kepada nombor reproduksi asas yang kurang daripada satu. Model terubah suai yang tidak termasuk jangkitan berulang mengesahkan bifurkasi ke hadapan. Penemuan-penemuan ini menggariskan kepentingan memasukkan kuarantin, kesan bifurkasi, dan faktor skala ke dalam strategi kawalan jangkitan yang berkesan. Penyelidikan ini menyumbang kepada usaha global untuk memerangi pandemik H1N1 dan menjaga kesihatan awam di Malaysia.

Kata kunci: influenza H1N1 novel; dinamik penyakit; langkah pencegahan; risiko penularan; parameter epidemiologi; nombor reproduksi asas

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