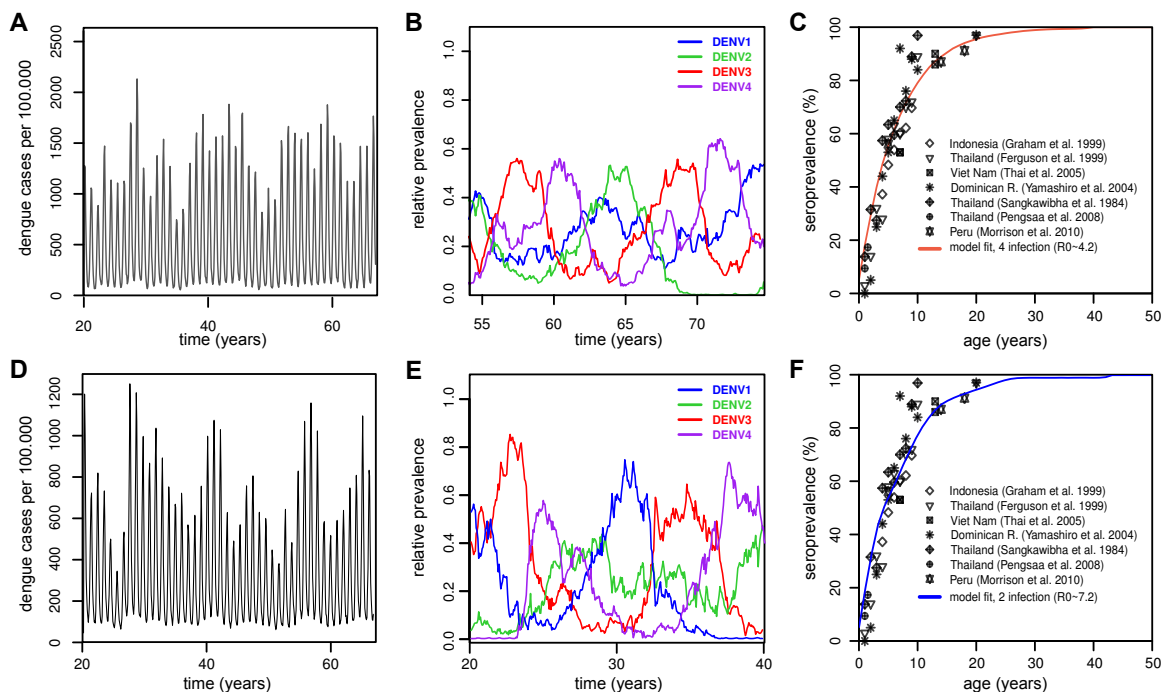


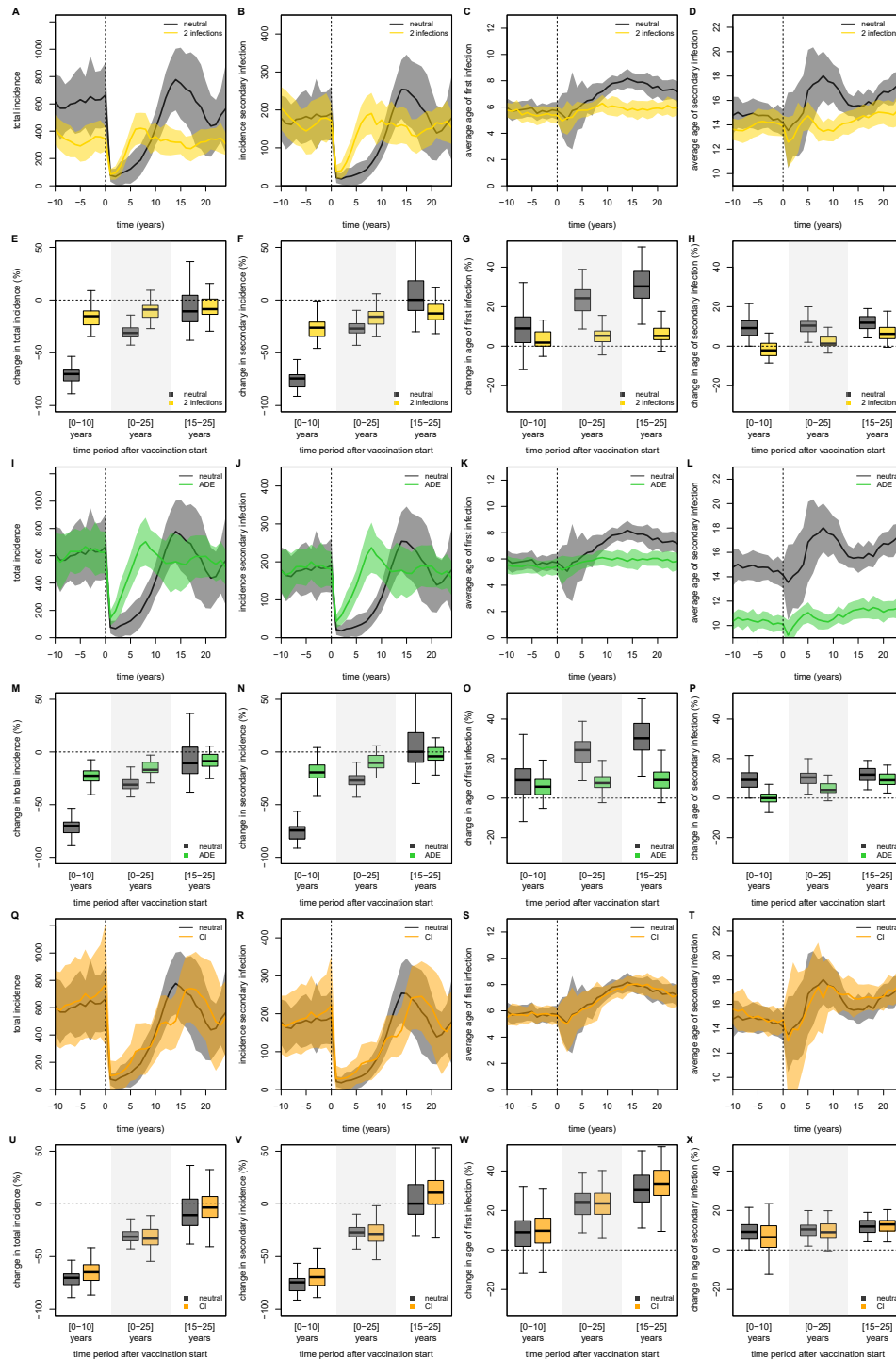
Dengue serotype immune-interactions and their consequences for vaccine impact predictions

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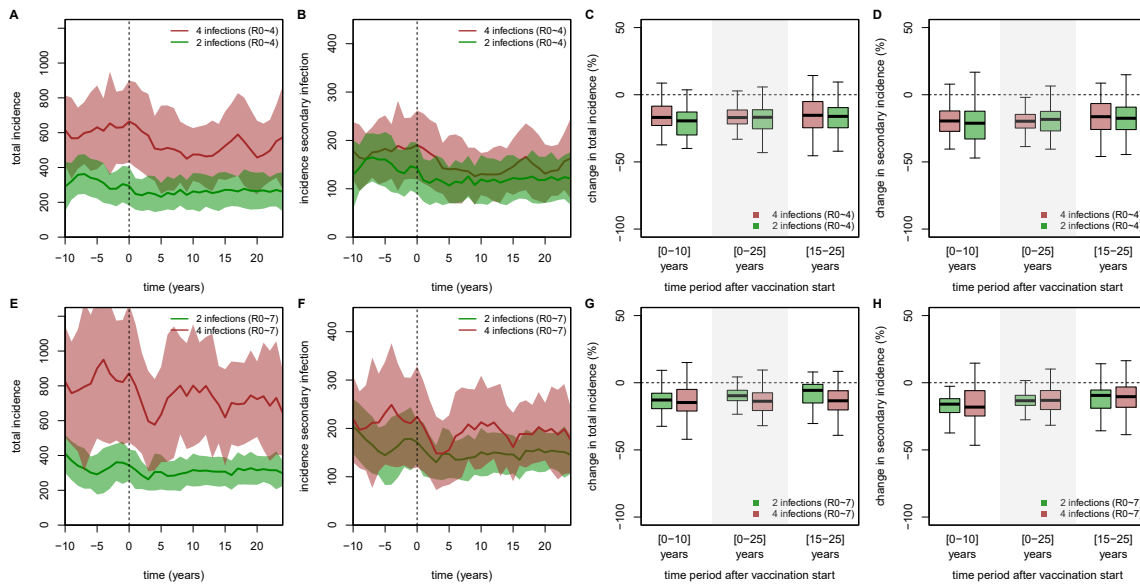
Supporting Information



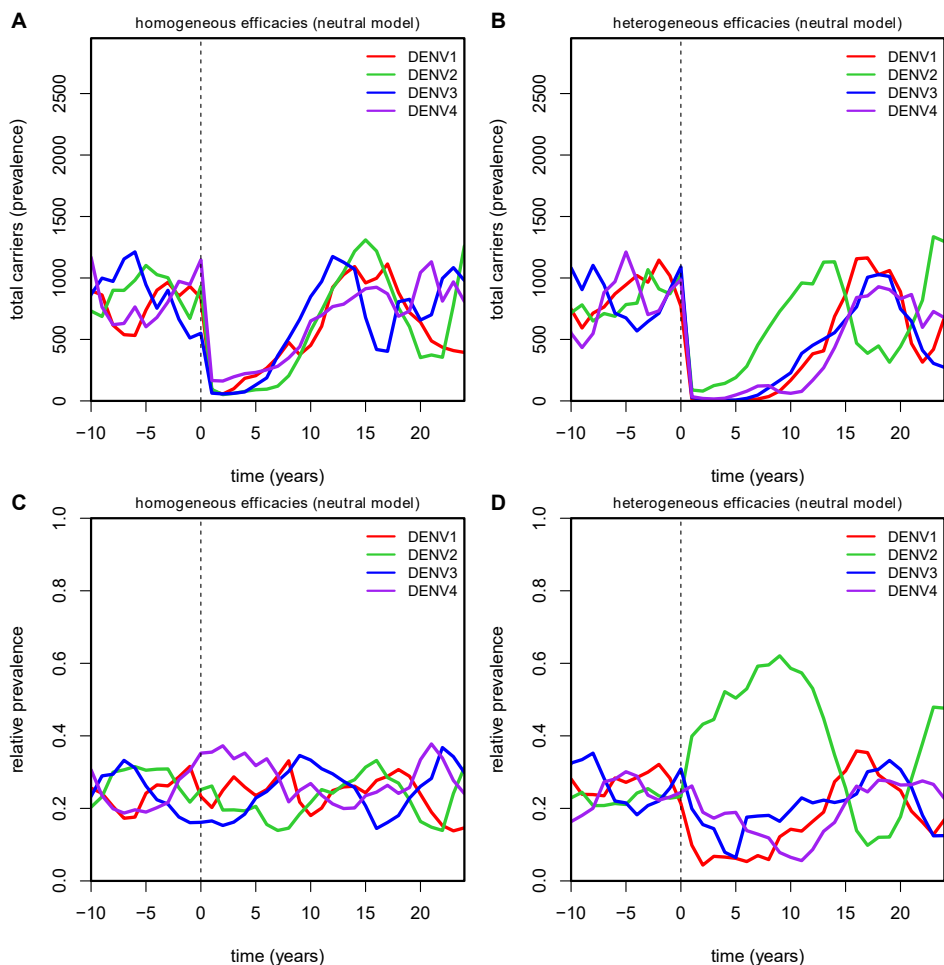
Supplementary Figure 1. Simulated epidemiologies and model fit. Examples of simulated timeseries of total dengue incidence (*A,D*) and relative serotype prevalence (*B,E*) for two models either allowing four consecutive infections (*top row*) or assuming full immunity after secondary infection (*bottom row*). In both cases the models were fit to seroprevalence data from highly endemic areas (*C,F*). The red and blue full lines in *C,F* are the average model behaviour for 200 simulations. Data taken from [1–7].



Supplementary Figure 2. Effect of immune-interactions on vaccination impact (full results). This figure completes the results presented on Figure 4 of the main text, by presenting the timeseries solution for secondary cases, as well as the ages of first and second infections. Results are based on 50 individual model runs. Please refer to the original figure legend for details.



Supplementary Figure 3. Effect of R_0 on the 2-infection and 4-infection models. (*A-D*). Results for the 4-infection (light red) and 2-infection (green) models when using low R_0 . (*E-H*) Results for the 4-infection (light red) and 2-infection (green) models when using high R_0 . The subplots present (*A,E*) 1-year average of total incidence; (*B,F*) 1-year average of secondary incidence; (*C,G*) changes in total incidence in the post-vaccination era; and (*D,H*) changes in secondary incidence in the post-vaccination era. Results are derived from 50 independent simulations. As the qualitative behaviour of both models is invariant to the level of R_0 , we note that the observations described in the main text between these two models are not simply due to their intrinsic differences in R_0 , as presented in the results section.



Supplementary Figure 4. Effect of heterogeneous vaccine efficacies on long-term serotype dynamics. (*A,C*) Serotype dynamics, presented as prevalence (*A*) and relative prevalence (*C*) for the neutral model with homogeneous efficacies as described in the main text. (*B,D*) Serotype dynamics, presented as prevalence (*B*) and relative prevalence (*D*) for the neutral model with heterogeneous efficacies (DENV1: 50%; DENV2: 35%; DENV3: 78%, DENV4: 75%). DENV2 is seen to escape earlier from the introduction of the vaccine at year 0 in the scenario with heterogeneous vaccine efficacies. Results are based on 25 simulations.

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