# Validation of a Static Model With a Dynamic Component by **Comparing With a Dynamic Model to Assess the Public Health Impact** of TAK-003 Vaccination: Case Studies in Thailand and Brazil

## INTRODUCTION

- Dynamic transmission models capture both direct and indirect effects of vaccination; however, they can be complex and computationally demanding
- Static models are easier to use, but do not capture indirect effects and may underestimate vaccine benefit
- A static model with a dynamic component (dyna-static model) was developed to approximate the dynamic model, and used to model the impact of TAK-003 on dengue epidemiology and the design of vaccination programs with TAK-003
- TAK-003 is a tetravalent dengue vaccine based on an attenuated dengue virus (DENV-2) backbone that is licensed for dengue prevention in the European Union, Great Britain, Brazil, Argentina, Thailand, and Indonesia, among other countries<sup>1,2</sup>

## **OBJECTIVES**

• To compare the results of the dyna-static model with the dynamic model and to validate the use of the dyna-static model to assess the impact of TAK-003 vaccination and inform vaccination program design, using 2 distinct dengue-endemic countries as case studies

## METHODS

## **DEVELOPMENT OF THE DYNA-STATIC MODEL**

- The dyna-static model is based on a static Markov model with 100 age cohorts (0-99 years) and 16 health states capturing up to 4 consecutive (not serotype specific) dengue infections (**Supplementary** Material) and 3 different levels of severity (asymptomatic, symptomatic nonhospitalized, and symptomatic hospitalized)
- The indirect effect was built into the structure by calculating the number of "infectious units," which combines the number of symptomatic and asymptomatic dengue infections and their relative contributions to transmission (Figure 1 and Supplementary Material)

## **FIGURE 1: DRIVERS OF THE NUMBER OF INFECTIOUS UNITS**



• At each cycle, the number of infectious units is calculated and compared with the reference number of infectious units (without vaccination); this ratio is then used to adjust the probability of infection in subsequent cycles

REFERENCES

1. Biswal S, et al. Lancet. 2020;395(10234):1423-1433. 2. Tricou V, et al. Lancet. 2020;395(10234):1434-1443.

the one-time vaccination of an additional age cohort in the year of vaccine introduction.

DISCLOSURES **EK:** External consultant for Putnam. **JS** and **RH:** Employees of Takeda Pharmaceuticals International AG and hold stock/stock options in Takeda Pharmaceuticals International AG. AT, JZ, and YGR: Employees of Putnam. SA: Employed at Putnam at time of study. Putnam received funding from Takeda Pharmaceuticals International AG for conducting relevant analysis for this study.





### **MODEL COMPARISON**

• The dyna-static model's performance is fine-tuned by varying cycle length and lag time (Figure 3)

• The cycle length is defined as the interval at which transitions between health states are evaluated and is considered semiflexible

• Lag time is defined as the number of cycles between a change in infectious units and a corresponding change in the probability of infection

### FIGURE 3: ELEVEN COMBINATIONS OF CYCLE AND LAG TIME WERE TESTED FOR EACH OF THE 12 SCENARIOS

Cycle	Length

o. of Cycles for Lag Time	1 wk	2 wk	1 mo	2 mo	3 mo
	1 wk	2 wk	1 mo	2 mo	3 mo
	2 wk	4 wk	2 mo	4 mo	6 mo
	3 wk	6 wk	3 mo	6 mo	9 mo
	4 wk	8 wk	4 mo	8 mo	12 mo
	5 wk	10 wk	5 mo	10 mo	15 mo
	6 wk	12 wk	6 mo	12 mo	18 mo
	7 wk	14 wk	7 mo	14 mo	21 mo
	8 wk	16 wk	8 mo	16 mo	24 mo
	9 wk	18 wk	9 mo	18 mo	27 mo
	10 wk	20 wk	10 mo	20 mo	30 mo
	11 wk	22 wk	11 mo	22 mo	33 mo
	12 wk	24 wk	12 mo	24 mo	36 mo

Approximately 1 mo Approximately 2 mo Approximately 3 mo Lag time:

• Eleven combinations of cycle length and lag time were tested across 12 scenarios (Figure 2), plus 2 no-vaccination scenarios, to find the optimal combination that minimizes deviations between the dyna-static and dynamic models across all assessed outcomes (symptomatic cases, hospitalized cases, and deaths) and for all time frames between 1 and 30 years

• The optimal combination of cycle length and lag time was determined using a set process, as described in the **Supplementary Material** 

ACKNOWLEDGMENTS

## RESULTS

• A cycle length of 2 months and a lag time of 2 months (1 cycle) are considered the optimal combination. Model comparison results are produced using this combination in the dyna-static model

• For both outcomes of symptomatic (Figures 4 and 6) and hospitalized (Figures 5 and 7) infections, the dyna-static model produces similar results to the dynamic model and in both types of dengue epidemiology

• The magnitude of the dyna-static model's deviations is generally associated with the expected level of indirect effects • The dyna-static model produces similar results to the dynamic model in both types of dengue epidemiology for time frames longer than ~5-8 years (Figures 4, 5, 6, and 7)

#### FIGURE 4: DYNA-STATIC MODEL VERSUS DYNAMIC MODEL WITH VACCINATION FIGURE 5: DYNA-STATIC MODEL VERSUS DYNAMIC MODEL WITH VACCINATION TOTAL SYMPTOMATIC INFECTIONS: THAILAND TOTAL HOSPITALIZED INFECTIONS: THAILAND

## FIGURE 6: DYNA-STATIC MODEL VERSUS DYNAMIC MODEL WITH VACCINATION TOTAL SYMPTOMATIC INFECTIONS: BRAZIL

### % Deviation From the Dynamic Model in Number of Cases Averted **Compared With No Vaccination**



asympt, asymptomatic; eff, efficacy; RX, routine vaccination at X years old (X=10 for Brazil); RX+5CU, routine vaccination with 5 years of catch-up; RX+10CU, routine vaccination with 10 years of catch-up; sympt, symptomatic.

## LIMITATIONS

• These findings are specific to Thai- and Brazilian-like epidemiological settings and may not be transferable to other types of dengue epidemiology

RX+10CU, routine vaccination with 10 years of catch-up; sympt, symptomatic.

- The optimal combination of cycle length and lag time was determined by testing all time horizons simultaneously (from 1 to 30 years); the conclusions may change if a local adaptation prioritized a specific time frame (short, medium, or long)
- The dyna-static model performs best when focused on longer timescales and is not well suited for shorter-term predictions (<5 years)

## CONCLUSIONS

- dynamics and vaccination impact

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asympt, asymptomatic; eff, efficacy; RX, routine vaccination at X years old (X=8 for Thailand); RX+5CU, routine vaccination with 5 years of catch-up; RX+10CU, routine vaccination with 10 years of catch-up; sympt, symptomatic

### FIGURE 7: DYNA-STATIC MODEL VERSUS DYNAMIC MODEL WITH VACCINATION TOTAL HOSPITALIZED INFECTIONS: BRAZIL

### % Deviation From the Dynamic Model in Number of Cases Averted **Compared With No Vaccination**



sympt, asymptomatic; eff, efficacy; RX, routine vaccination at X years old (X=10 for Brazil); RX+5CU, routine vaccination with 5 years catch up; RX+10CU, routine vaccination with 10 years catch up; sympt, symptomatic

• We developed a new static model with a dynamic component (dyna-static model) to approximate dynamic dengue transmission to assess the impact of dengue vaccination and inform vaccination program design

• The dyna-static model's validity was tested in 2 distinct dengue-endemic countries across scenarios with different levels of indirect effects; in both case studies, the dyna-static model provided good approximations to the dynamic model over longer-term time horizons (approximately 10 years and longer); long-term validity is key when considering infectious disease

 The dyna-static model is a valuable tool for evaluation of dengue vaccines in settings with limited resources and technical capabilities



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