## Modeling Insights into *Haemophilus influenzae* Type b Disease, Transmission, and Vaccine Programs

## **Technical Appendix 1**

## **Model Structure**

The following set of partial differential equations defines the rates at which the simulated population moves between model states:

$$\frac{\partial NS}{\partial t} + \frac{\partial NS}{\partial a} = \mu(t, a)X(t) + \omega_L LS(t, a) + \omega_{BPIG}I(t, a) - V(t) + \lambda(t, a) + \gamma(t, a)\varepsilon(a) + \delta_{BPIG}(t, a) - V(t) + \lambda(t, a) + \gamma(t, a)\varepsilon(a) + \delta_{BPIG}(t, a) - V(t) + \lambda(t, a) + \gamma(t, a)\varepsilon(a) + \delta_{BPIG}(t, a) - V(t) + \lambda(t, a) + \gamma(t, a)\varepsilon(a) + \delta_{BPIG}(t, a) - V(t) + \lambda(t, a) + \gamma(t, a)\varepsilon(a) + \delta_{BPIG}(t, a) - V(t) + \lambda(t, a) +$$

$$\frac{\partial I}{\partial t} + \frac{\partial I}{\partial a} = \delta_{BPIG}(t, a)NS(t, a) - \omega_{BPIG}I(t, a)$$

$$\frac{\partial NC}{\partial t} + \frac{\partial NC}{\partial a} = \lambda(t, a)NS(t, a) - \sqrt{(t) + \rho_C + \sigma(a) + \gamma(t, a)\varepsilon(a)} \frac{NC}{NC}(t, a)$$

$$\frac{\partial LS}{\partial t} + \frac{\partial LS}{\partial a} = \omega_H HS(t, a) - \Psi(t) + \omega_L + \lambda(t, a)(1 - \alpha_L) + \gamma(t, a)\varepsilon(a) \underline{L}S(t, a)$$

$$\frac{\partial LC}{\partial t} + \frac{\partial LC}{\partial a} = \lambda(t, a)(1 - \alpha_L)LS(t, a) - \left[ \mathbf{f}(t) + \rho_C + \sigma(a)(1 - \beta_L) + \gamma(t, a)\varepsilon(a) \right] \underline{L}C(t, a)$$

$$\frac{\partial HS}{\partial t} + \frac{\partial HS}{\partial a} = \rho_C VC(t,a) + LC(t,a) + HC(t,a) + \rho_D D(t,a) + \gamma(t,a)\varepsilon(a) VS(t,a) + LS(t,a) - V(t) + \omega_H(a) + \lambda(t,a)(1-\alpha_H) - \frac{1}{2}HS(t,a)$$

$$\frac{\partial HC}{\partial t} + \frac{\partial HC}{\partial a} = \lambda(t, a)(1 - \alpha_H)HS(t, a) + \gamma(t, a)\varepsilon(a)[NC(t, a) + LC(t, a)] - (t) + \rho_C + \sigma(a)(1 - \beta_H)HC(t, a)$$

$$\frac{\partial D}{\partial t} + \frac{\partial D}{\partial a} = \sigma(a) VC(t, a) + (1 - \beta_L)LC(t, a) + (1 - \beta_H)HC(t, a) - V(t) + \rho_D D(t, a)$$

## In which:

- NS, NC, LS, LC, HS, HC, D, and I are population states, where N=No antibody, L = Low antibody, H = High antibody, S = Susceptible, C = Colonized, D = Diseased, and I = Immune; X(t) is the total population.
- $\mu(t,a)$  and  $\nu(t)$  are time-dependent birth and death rates, respectively. Birth rate also depends on age as individuals are only born into the age=0 group
- $\omega_L$  is the rate at which low antibody wanes to no antibody and  $\omega_H(a)$  is the age-dependent rate at which high antibody wanes to low antibody
- $\lambda(t,a)$  is the time- and age-dependent force of infection
- $\gamma(t,a)$  is the time- and age-dependent rate of vaccination, and  $\epsilon(a)$  is the age-dependent vaccine take rate
- $\sigma(a)$  is the age-dependent rate of invasive disease among colonized persons
- $\alpha_L$  and  $\alpha_H$  are the efficacy of low and high antibody at preventing colonization
- $\beta_L$  and  $\beta_H$  are the efficacy of low and high antibody at preventing invasive disease
- $\rho_C$  and  $\rho_D$  are the rates of recovery from colonization and invasive disease, respectively
- $\delta_{BPIG}(t,a)$  is the time- and age-dependent rate of BPIG use (for Alaska Native populations only), and  $\omega_{BPIG}$  is the rate of waning of BPIG protection.