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Appendix

The structure of the model is represented by the following set of differential equations.

$$\begin{aligned}
 \frac{dX_{ki}^{00}}{dt} &= \Lambda_{ki}(t) - \sigma_{ki} X_{ki}^{01} - (\mu + \lambda_{1ki}(t) + \rho_{ki}(t)) X_{ki}^{00}(t) - C_{ki}^{01}(T2) + R_{ki}^{00}(t) \\
 \frac{dX_{ki}^{10}}{dt} &= \lambda_{1ki}(t) X_{ki}^{00}(t) - \sigma_{ki} X_{ki}^{11} - (\mu + \lambda_{2ki}(t) + \rho_{ki}(t)) X_{ki}^{10}(t) - C_{ki}^{11}(T2) + R_{ki}^{01}(t) \quad (0.2) \\
 \frac{dX_{ki}^{20}}{dt} &= \gamma_1 X_{ki}^{10}(t) - \sigma_{ki} X_{ki}^{21} - (\mu + \gamma_2 + \rho_{ki}(t)) X_{ki}^{20}(t) + C_{ki}^{21}(T2) \\
 \frac{dX_{ki}^{30}}{dt} &= \gamma_2 X_{ki}^{20}(t) - \sigma_{ki} X_{ki}^{31} - (\mu + \gamma_3 + \rho_{ki}(t)) X_{ki}^{30}(t) + C_{ki}^{31}(T2) \\
 \frac{dX_{ki}^{01}}{dt} &= \rho_{ki}(t) X_{ki}^{00}(t) - (\mu + \sigma_{ki} + \lambda_{2ki}(t)) X_{ki}^{01}(t) - C_{ki}^{01}(T1) \\
 \frac{dX_{ki}^{11}}{dt} &= \lambda_{2ki}(t) X_{ki}^{01}(t) - \rho_{ki}(t) X_{ki}^{10}(t) - (\mu + \sigma_{ki} + \gamma_2) X_{ki}^{11}(t) - C_{ki}^{11}(T2) \\
 \frac{dX_{ki}^{21}}{dt} &= \gamma_1 X_{ki}^{11}(t) - \rho_{ki}(t) X_{ki}^{20}(t) - (\mu + \sigma_{ki} + \gamma_2) X_{ki}^{21}(t) - C_{ki}^{21}(T2) \\
 \frac{dX_{ki}^{31}}{dt} &= \gamma_2 X_{ki}^{21}(t) - \rho_{ki}(t) X_{ki}^{30}(t) - (\mu + \sigma_{ki} + \gamma_3) X_{ki}^{31}(t) - C_{ki}^{31}(T2) \quad (0.8) \\
 \frac{dX_{ki}^{40}}{dt} &= \gamma_3 X_{ki}^{30}(t) - \sigma_{ki} X_{ki}^{41} - (\mu + \alpha) X_{ki}^{40}(t) - C_{ki}^{41}(T2) \quad (0.9) \\
 \frac{dX_{ki}^{41}}{dt} &= \gamma_3 X_{ki}^{31}(t) - (\mu + \sigma_{ki} + \alpha) X_{ki}^{41}(t) - C_{ki}^{41}(T2) \quad (0.10)
 \end{aligned}$$

Screening and treatment of Gc:

With: $C_{ki}^{h1}(T2) = C_{ki} \% \cdot F_{ki} X_{ki}^{h1}(t) \cdot \mathbb{1}_{\{t \geq \theta_1\}}, \forall h, k,$

$$\text{for } \theta: \begin{cases} t & \text{if } t \neq T2 \\ T2 & \end{cases}$$

Condom use per partnership:

$$\begin{aligned}
 \beta_{k,i,j}^{new,h}(t) &= (Cond_k(T2) \% \cdot \mathbb{1}_{\{t \geq \theta_2\}}) \cdot 1 - (Cond_{k',j,i}(T1) \% \cdot \mathbb{1}_{\{t \geq \theta_1\}}) \beta_{k,i,j}^h \\
 \xi_{new,k} &= Cond_{k',j,i}(T2) \% \cdot \mathbb{1}_{\{t \geq \theta_2\}} - (Cond_{k',j,i}(T1) \% \cdot \mathbb{1}_{\{t \geq \theta_1\}}) \xi_{k,i,j},
 \end{aligned}$$

$\forall h, k$

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with $Cond_{k',j,i}(t) = Cond_{k,i,j}(t) \forall k, i, j$

$$\text{for } \theta_1: \begin{cases} t & \text{if } t \neq T1 \\ T1 - 1 & \text{if } t \geq T1 \end{cases}$$

$$\text{for } \theta_2: \begin{cases} t & \text{if } t \neq T2 \\ T2 - 1 & \text{if } t \geq T2 \end{cases}$$

Here $X_{ki}^{hs}(t)$ represents the number of individuals of sex k ($k = 1$: female; $k = 2$: male) and activity class i ($i = 1$: low rate of partner change; $i = 2$: higher rates of partner change) in disease states h (HIV status) and s (STD status) at time t . An individual is in state $h = 0$ if susceptible to HIV, in state $h = 1$ if in phase 1 of HIV infection, in state $h = 2$, or $h = 3$ if in phase 2 and 3 of HIV infection respectively. An individual in state $h = 4$ is diagnosed with full blown AIDS. An individual is susceptible to the STD if $s = 0$ and currently infected with the STD if $s = 1$. C_{ki} represents the fraction of individuals in risk group k, i screened/diagnosed and treated at time $T2$ (coverage). F_{ki} is the frequency of screening in activity class i of sex k per year. $Cond_{k,i,j}(T1)$ and $Cond_{k,i,j}(T2)$ are the fraction of individuals k, i who start using condoms in partnership with someone of opposite sex k' and activity class j at time $T1$ and $T2$.

An individual susceptible to both infections ($X_{ki}^{00}(t)$) can get infected either by the STD or HIV at a rate $\rho_{ki}(t)$ (eqn 0.7) and $\lambda_{1ki}(t)$ (eqn 0.8) respectively. It is assumed that HIV positives and negatives are equally likely to acquire an STD infection. However, HIV negative individuals ($X_{ki}^{01}(t)$ or $X_{ki}^{00}(t)$) are more likely to get HIV from HIV-STD $X_{ki}^{11}(t)$, $X_{ki}^{21}(t)$, $X_{ki}^{31}(t)$ partners. STD positive individuals ($X_{ki}^{01}(t)$) are more susceptible to infection with HIV ($\lambda_{2ki}(t)$) (eqn 0.9). At time $t = 0$ (i.e. 1980), $\sum_{i=1}^2 X_{1i}^{00}(0) = 122605$

and $\sum_{i=1}^2 X_{2i}^{00}(0) = 115558$ and the HIV epidemic is seeded with $X_{11}^{10}(0) = 1, X_{12}^{10}(0) = 2, X_{21}^{10}(0) = 1, X_{22}^{10}(0) = 1$ while the STD is already at its initial equilibrium prevalence before the introduction of HIV. The other state variables are equal to 0. Once infected, an STD positive of sex k and activity class i remains infected and infectious for a period of time of $1/\sigma_{ki}$ respectively after which period, the individual joins the susceptible class again. Whether STD positive or negative, HIV seroconverters remain symptomless for an average period of time $(1/\gamma_1 + 1/\gamma_2 + 1/\gamma_3)$ before progressing to the AIDS stage. Individuals in the population die at a natural rate μ ($\mu = \text{average sexual life expectancy}$). The AIDS related mortality, α is equal to $1/\text{life expectancy of patient with full blown AIDS}$.

The per capita rate of STD infection is defined as follows:

$$\rho_{ki}(t) = m_{ki}(t) \sum \xi_{new_{k'ji}} \varphi_{kij}(t) \frac{\sum_{h=0}^3 X_{k'j}^{h1}(t)}{\sum_{hs} X_{k'j}^{hs}(t) + \sum_s X_{k'j}^{4s}(t)} \quad (0.11)$$

The per capita rates of HIV infection for those without (STD) and with (STD) current infection with the cofactor STD are defined as follows:

$$\lambda_{1ki}(t) = m_{ki}(t) \sum \varphi_{kij}(t) \frac{\sum_{h=1}^3 \beta_{new_{k'ji}^h} (X_{k'j}(t)^{h0} + RR_{HIV/STD} \cdot X_{k'j}(t)^{h1}(t))}{\sum_{hs} X_{k'j}^{hs}(t) + \sum_s X_{k'j}^{4s}(t)} \quad (0.12)$$

$$\lambda_{2ki}(t) = RR_{HIV/STD} \cdot \lambda_{1ki}(t) \quad (0.13)$$

As seen in equations (0.7-0.8), the rates of infection depend on the annual rate of partner acquisition ($m_{ki}(t)$) at time t of an individual of sex k and sexual activity level i ; the probability of choosing an infected partner, which is a function of the mixing pattern $\varphi_{kij}(t)$

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(the probability that a member of class i and sex k has selected a partner of sex k' and class j at time t); the probability that this partner is infected (prevalence of infection in risk group k,i); and the per partner transmission probability of the STD ($\xi_{new_{k'ji}}$) or HIV ($\beta_{new_{ji}}$) from a partner of sex k' , activity class j to a susceptible of sex k and activity class

AIDS patients are assumed not to contribute to transmission because of the severity of their illness. The term $RR_{HIV/STD}$ in (0.13) and in the right hand side of (0.12) represents the relative increase in HIV susceptibility of HIV⁻ STD⁻ and in infectivity of STD⁺/HIV⁺ respectively.

New recruits join the susceptible sexually active population at a rate $\Lambda_{ki}(t)$. The expression for the new recruits to the sexually active population is:

$$\Lambda_{ki}(t) = Q_{ki} R P f \left[X_{1i}^{01}(t - \tau) + X_{1i}(t - \tau) (1 - \gamma) \left(X_{ki}^{10}(t) + X_{ki}^{11}(t) + X_{1i}^{20}(t - \tau) + X_{1i}^{21}(t - \tau) \right) \right] \quad (0.14)$$

Here Q_{ki} is the initial distribution in sexual activity for each sex in the absence of AIDS induced mortality. R denotes the sex ratio (assumed to be 1:1 at birth which gives $R = 0.5$). f denotes the per capita birth rate of sexually active women ($f=0.3$). P is the proportion of uninfected infants who survive to join the sexually active age classes [49] ($P = \exp^{-b\tau}$ where τ is the age at sexual maturity (15 years) and b is the death rate over the interval $[0,\tau]$). The net rate of prostitute renewal is given by

$$R_{ki}^{00} = \gamma \cdot X_{ki}^{30}(t)/2, \text{ for } k = 1, i = 2 \quad (0.15)$$

$$R_{ki}^{00} = 0, \text{ otherwise} \quad (0.16)$$

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$$R_{ki}^{01} = \gamma_3 X_{ki}^{31}(t)/2 \quad \text{or } k = i \quad (0.17)$$

$$R_k^0 = 0, \text{ otherwise} \quad (0.18)$$

We define the elements of the mixing matrix as [49]

$$\varphi_{kij} = \frac{W_{kij} (N_{k'j}^+(t) - N_{k'j}^-(t)) m_{k'j}(0)}{\sum W_k (N_{k'j}^+(t) - N_{k'j}^-(t)) m_{k'j}(0)} \quad (0.19)$$

The W_k define a set of weights which represent the preference of someone of sex k in activity class i for someone of the opposite sex in activity class

The elements of the mixing matrix $\varphi_{kij}(t)$ must satisfy the following constraint [49]

$$\sum_i \varphi_{kij}(t) = 1 \quad (0.20)$$

$$\sum_j \varphi_{kij}(t) = 1$$

$$N_{ki}^-(t) - N_{ki}^+(t) = \sum_j \varphi_{kij}(t) (N_{k'j}^+(t) - N_{k'j}^-(t)) m_{k'j}(0) \quad (0.22)$$

As described in ref [49] we adopt a procedure which the mean rate of partner change in the lowest female activity group remains unchanged (for all t) to act as a reference point to define a rate of change as mortality influences population size and structure (condition (0.22) to hold) and k at the cross product the preference matrix (hence the mixing matrix) should be equal for males and females (details in ref [49])

For Cotonou we used the weights: $W_{111}(1980) = 3.335$, $W_{112}(1980) = 3.335$, $W_{121}(1980) = 3.335$, $W_{122}(1980) = 3.335$, $W_{211}(1980) = 17$, $W_{212}(1980) = 17$, $W_{221}(1980) = 6.5$, $W_{222}(1980) = 6.5$. The corresponding elements of the mixing matrix are given in Table 1. We also use the following

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recovery rate of Gc: $\sigma_{11} = 5.5, \sigma_{12} = 8.8, \sigma_{21} = 5.5, \sigma_{22} = 8.8$ For the high prevalence scenario we used the weights: $W_{111}(1980) = 2, W_{112}(1980) = 1, W_{211}(1980) = 2, W_{212}(1980) = 1$ and $W_{221}(1980) = 1, W_{222}(1980) = 1$. To obtain the same Gc equilibrium prevalence in 1980 than for Cotonou, the following Gc recovery rates were used: $\sigma_{11} = 6.5, \sigma_{12} = 8.4, \sigma_{21} = 6.5, \sigma_{22} = 8.4$.

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