## **Hierarchical Mixing Matrices**

The standard mixing matrix, defined in (Moss et al. 2019) as

"For each data set we constructed a family of mixing matrices M, under the assumption that some proportion of infections  $\delta_i^H$  arising from a resident of region  $\delta_i$  occur in the resident's home region i and that the remaining proportion  $\delta_i^* = 1 - \delta_i^H$  of infections occur outside of region i"

$$M = \begin{pmatrix} \delta^H & \delta^A & \cdots & \delta^A \\ \delta^A & \delta^H & \cdots & \delta^A \\ \vdots & \vdots & \ddots & \vdots \\ \delta^A & \delta^A & \cdots & \delta^H \end{pmatrix}$$

We simplify this by assuming that each region shares the proportion of infections that occur in the home region,  $\delta_H$ , and those that occur from outside the region,  $\delta^A = 1 - \delta^H$ , to produce the matrix to produce the single level mixing matrix

Similarly,  $\delta^A$ , the proportion of infections occurring outside an individuals home region can be divided into those infections which occur inside the home region at the next highest scale,  $\delta^{H_2}$ , and those that occur at outside this higher level home region  $\delta^{A_2}$ , to create the two level mixing matrix:

$$M = \begin{pmatrix} \delta^H & \delta^A \cdot L_{1,2}^2 & \cdots & \delta^A \cdot L_{1,j}^2 \\ \delta^A \cdot L_{2,1}^2 & \delta^H & \cdots & \delta^A \cdot L_{2,j}^2 \\ \vdots & \vdots & \ddots & \vdots \\ \delta^A \cdot L_{j,1}^2 & \delta^A \cdot L_{j,2}^2 & \cdots & \delta^H \end{pmatrix}$$

where

$$L_{i,j}^k = \begin{cases} \frac{\delta^H}{N_{j,k}} & \text{if } j \in R_i^k \\ \frac{\delta^A}{N_{j,k}} & \text{if } j \notin R_i^k \ \& \ k = n \\ \delta^A L_{i,j}^{k+1} & otherwise \end{cases}$$

Where  $R_i^k$ , is the set of first level regions in the same k level region as i, and  $N_{j,k}$  is the number of (first level) regions in the same level k region as j, and n is the total number of levels.

The sum of each row is given by

$$\sum_{i=1}^{N} M_{i,j} = \delta^H + \delta^A \delta^H + \dots + \delta^{H^{n-1}} \delta^A + \delta^{H^n}$$

Which factorises to

$$\sum_{j=1}^{N} M_{i,j} = \delta^H + \delta^A (\delta^H + \delta^A \delta^H + \dots + \delta^{A^{n-2}} \delta^H + \delta^{A^{n-1}})$$

iteratively until

$$\sum_{j=1}^{N} M_{i,j} = \delta^H + \delta^A (\delta^H + \delta^A (\delta^H + \delta^A (\cdots (\delta^H + \delta^A)))$$

since  $\delta^H + \delta^A = 1$ , then

$$\sum_{i=1}^{N} M_{i,j} = \delta^H + \delta^A = 1$$

As a real world example, we can look at a small group of suburbs in the greater melbourne region.

using DrWatson <a href="Qquickactivate">Qquickactivate</a> using Shapefile using Tables using Plots include(srcdir("Spa

Kensington, Racecourse, and West industrial are SA2 regions within the 'Melbourne City' SA3 region, Footscray and Maribyrnong are SA2 regions within the 'Maribyrnong' SA3 region, and Flemington and Ascot Vale are SA2 regions withing the 'Essendon' SA3 region. Furthermore, both 'Melbourne city' and Essendon are within the 'Inner Melbourne' SA4 region while Marybyrnong lies iside the 'West Melbourne' SA4 Region

 $V = \{Kensington, Racecourse, Industrial, Flemington, Ascot, Footscray, Maribyrnong\}$ 

$$M = \begin{pmatrix} \delta_H & \delta_A \cdot \frac{\delta_H}{2} & \delta_A \cdot \frac{\delta_H}{2} & \delta_A \cdot \frac{\delta_H}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_H}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_H}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} \\ \delta_A \cdot \frac{\delta_H}{2} & \delta_H & \delta_A \cdot \frac{\delta_H}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_H}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_H}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} \\ \delta_A \cdot \frac{\delta_H}{2} & \delta_A \cdot \frac{\delta_H}{2} & \delta_H & \delta_A \cdot \delta_A \cdot \frac{\delta_H}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_H}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} \\ \delta_A \cdot \delta_A \cdot \frac{\delta_H}{3} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} \\ \delta_A \cdot \delta_A \cdot \frac{\delta_H}{3} & \delta_A \cdot \delta_A \cdot \frac{\delta_H}{3} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{3} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} \\ \delta_A \cdot \delta_A \cdot \frac{\delta_A}{3} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} \\ \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} \\ \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} \\ \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} \\ \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} \\ \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} \\ \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2} & \delta_A \cdot \delta_A \cdot \frac{\delta_A}{2}$$

Moss, Robert, Elham Naghizade, Martin Tomko, and Nicholas Geard. 2019. "What Can Urban Mobility Data Reveal about the Spatial Distribution of Infection in a Single City?" *BMC Public Health* 19 (1): 656. https://doi.org/10.1186/s12889-019-6968-x.