recombination events $x_i = x_1 + x_2$ x_3 $x_j = x_4 + x_5$ x_6 x_k tree-change events $x_u = x_1 + x_2 + x_3$

 \mathcal{G}_4

 \mathcal{G}_5

(B) Parameterized MSC model (
$$S$$
)
$$W_{12} = \begin{bmatrix} N_2 \\ N_1 \end{bmatrix}$$

(A) Proposed ARG

 \mathcal{G}_1

(C) Extract genealogies between each event type

(recombination events) $G_r = [\mathcal{G}_1, \mathcal{G}_2, \mathcal{G}_3, ...]$ $G_q = [\mathcal{G}_1, \mathcal{G}_3, \mathcal{G}_4, ...]$ (tree-change events)

 $G_t = [\mathcal{G}_1, \mathcal{G}_4, \mathcal{G}_7, ...]$ (topology-change events) (D) Extract waiting distances for each event type

$$X_r = [x_1, x_2, x_3, ...]$$
 (recombination events) $X_g = [x_i, x_3, x_j, ...]$ (tree-change events)

$$X_t = [x_u, x_v, x_z, ...]$$
 (topology-change events) (E) Calculate MS-SMC rate parameters (λ) for

each event type given S, G, and recomb rate

each event type given
$$\mathcal{S},\,\mathcal{G},\,$$
 and recomb rate $\Lambda_r=[_{\lambda r1},\lambda_{r2},\lambda_{r3},...]$ (recombination events)

 $\Lambda_{a} = [\lambda_{a1}, \lambda_{a3}, \lambda_{a4}, ...]$ (tree-change events)

$$\Lambda_g = [\lambda_{g1}, \lambda_{g3}, \lambda_{g4}, ...]$$
 (tree-change events

 $\Lambda_t = [\lambda_{t1}, \lambda_{t4}, \lambda_{t7}, ...]$ (topology-change events)

(F) Calculate likelihood of waiting distances $\mathcal{L}(X_r|\Lambda_r)$ (recombination events)

 $\mathcal{L}(X_q|\Lambda_q)$ (tree-change events)

 $\mathcal{L}(X_t|\Lambda_t)$ (topology-change events)

 $\mathcal{L}(X_q|\Lambda_q) * \mathcal{L}(X_t|\Lambda_t)$ (tree & topo-change events)