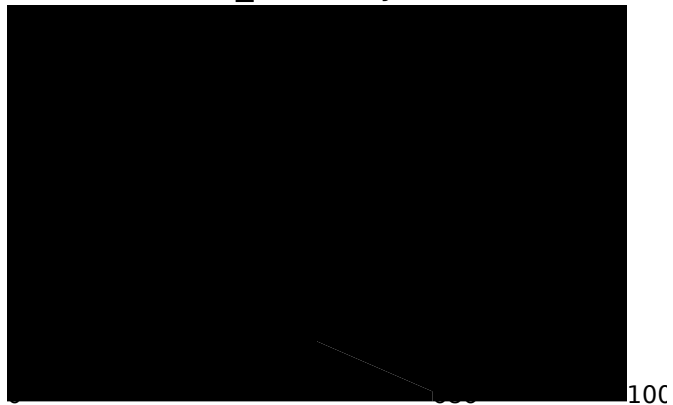


(A)

```
ts = sim_ancestry(3, ...)
```



Genome position

(B)

```
mts = sim_mutations(ts, ...)
```



Genome position