

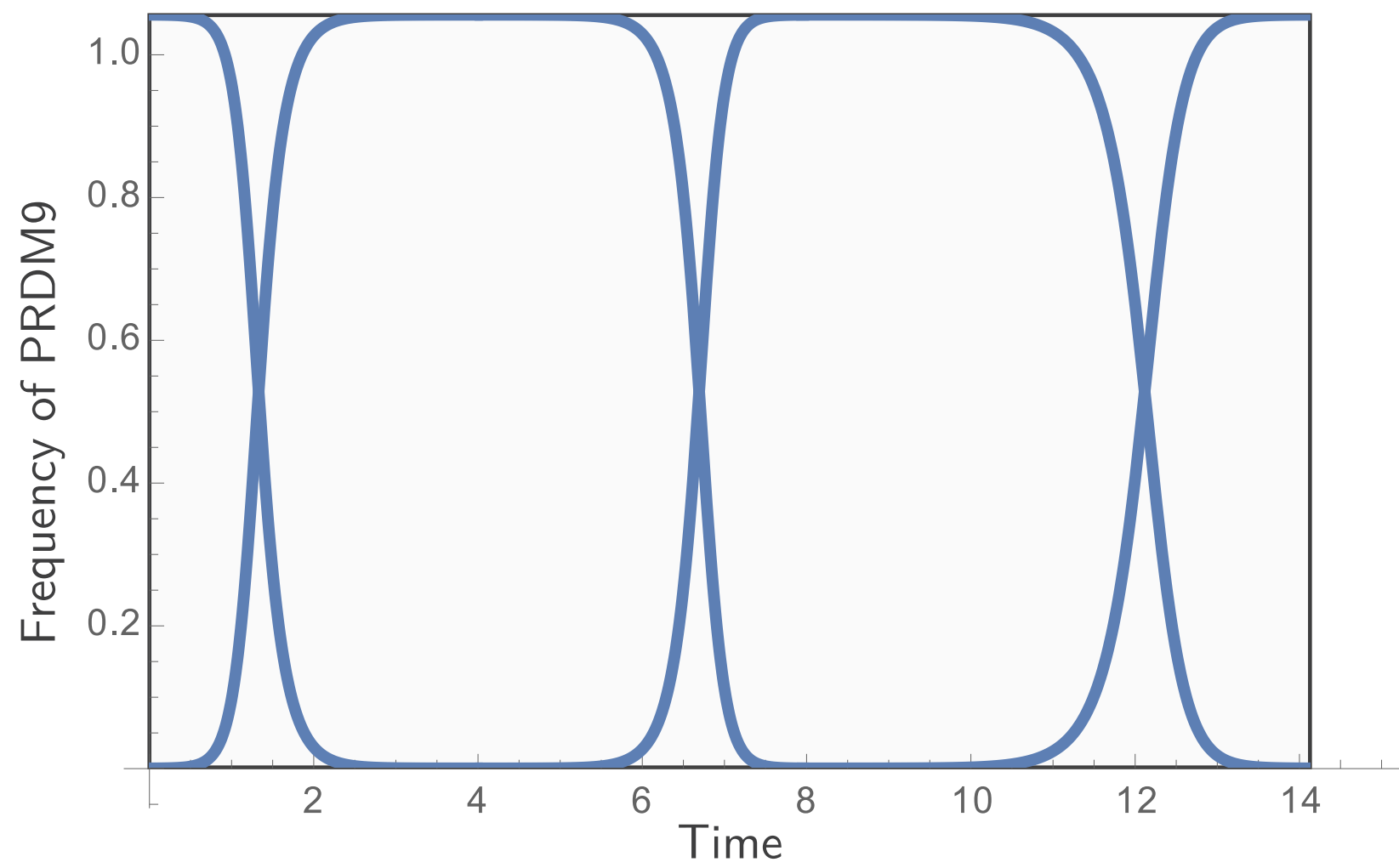
$$N_e, u, v, r_0, f$$

$$N_e u$$

The expected number of new PRDM9 variants in the population at each generation.

$$N_e u \ll 1$$

Succession of  
PRDM9 variants



$$N_e u \gg 1$$

Polymorphism of  
PRDM9 variants

