- w_{AA} = the probability that a drive homozygous embryo survives to adult-hood
- w_{Aa} = the probability that a drive heterozygous embryo survives to adulthood
- w_{aa} = the probability that a wild-type embryo survives to adulthood

•
$$\overline{w(t)} = u^2(t)w_{AA} + 2u(t)(1 - u(t))w_{Aa} + (1 - u(t))^2w_{aa}$$

•
$$E[u(t+1)] = \frac{u^2(t)w_{AA} + u(t)(1-u(t))w_{Aa}}{\overline{w(t)}}$$

•
$$E[u_{AA}(t+1)] = \frac{w_{AA}}{w(t)}u^2(t)$$

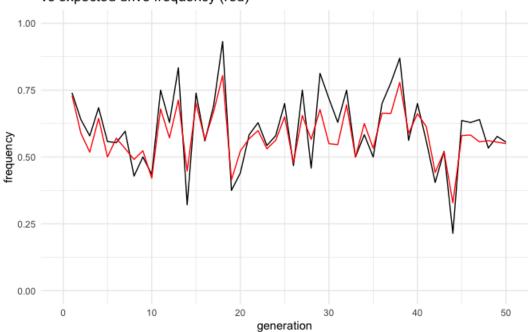
•
$$E[u_{Aa}(t+1)] = \frac{w_{Aa}}{w(t)} 2u(t)(1-u(t))$$

•
$$E[u_{aa}(t+1)] = \frac{w_{aa}}{w(t)}(1-u(t))^2$$

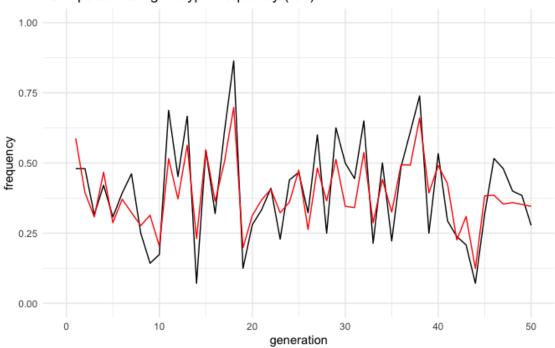
 In SLiM, the drive frequency before viability selection is applied is the drive frequency during an early() event, after all embryos are created but before fitness() is applied.

x < -a/2:

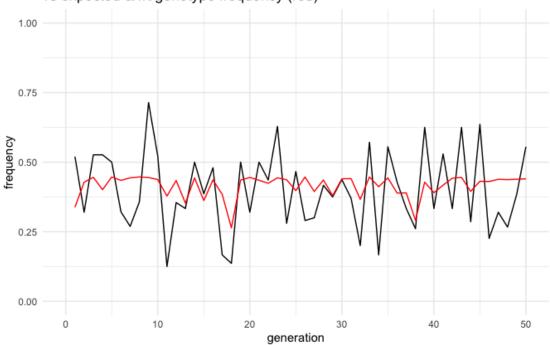
a = 0.0688 & x = 0.47 observed drive frequency (black) vs expected drive frequency (red)



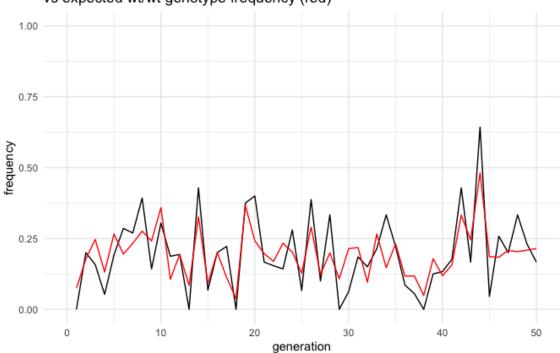
a = 0.0688 & x = 0.47 observed d/d genotype frequency (black) vs expected d/d genotype frequency (red)



a = 0.0688 & x = 0.47 observed d/wt genotype frequency (black) vs expected d/wt genotype frequency (red)

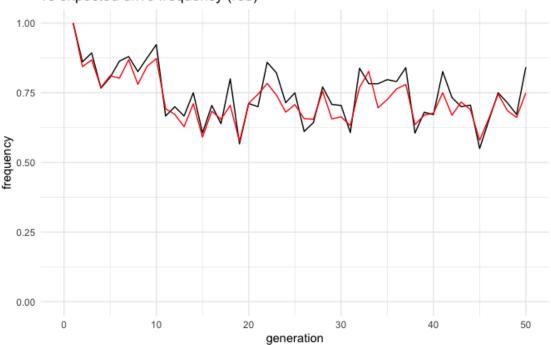


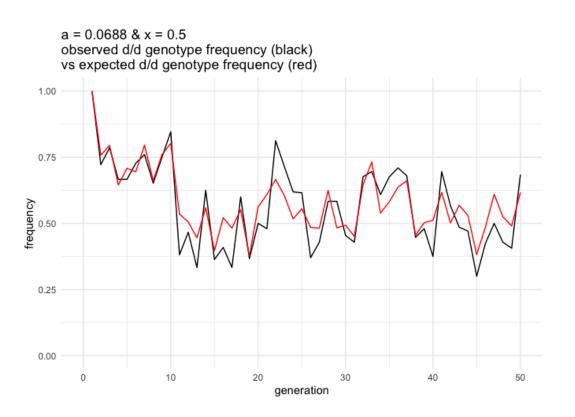
a = 0.0688 & x = 0.47 observed wt/wt genotype frequency (black) vs expected wt/wt genotype frequency (red)



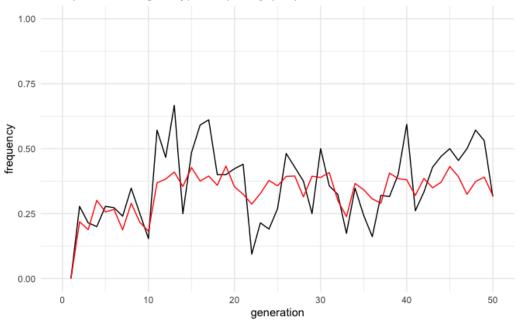
$-a/2 \le x \le a/2$:

a = 0.0688 & x = 0.5 observed drive frequency (black) vs expected drive frequency (red)

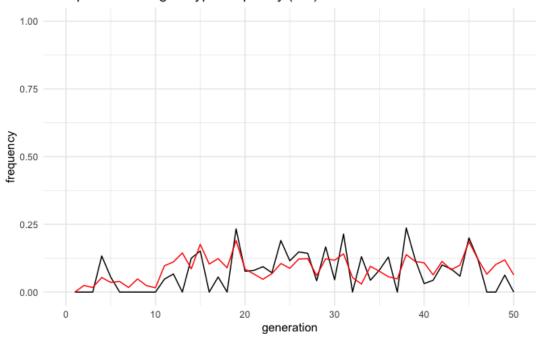




a = 0.0688 & x = 0.5 observed d/wt genotype frequency (black) vs expected d/wt genotype frequency (red)

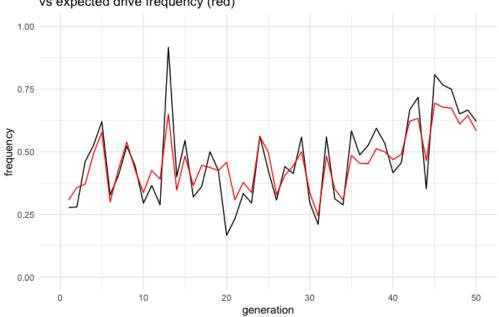


a = 0.0688 & x = 0.5 observed wt/wt genotype frequency (black) vs expected wt/wt genotype frequency (red)

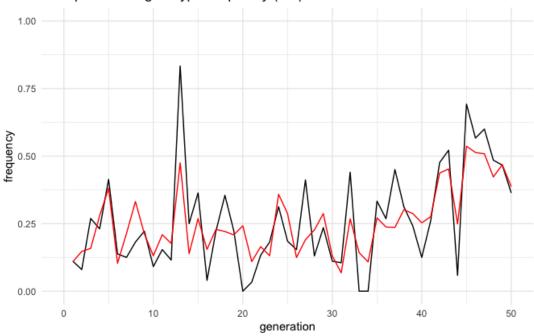


x > a/2:

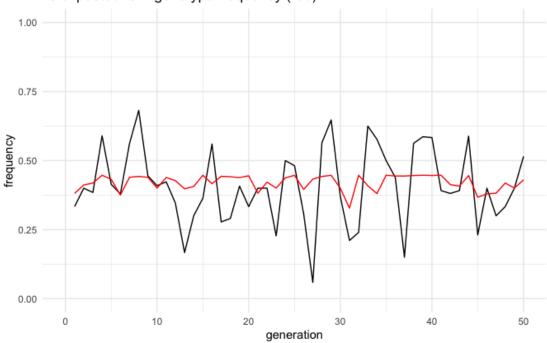
a = 0.0688 & x = 0.54 observed drive frequency (black) vs expected drive frequency (red)



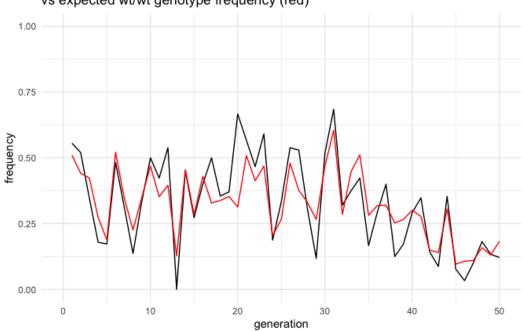
a = 0.0688 & x = 0.54 observed d/d genotype frequency (black) vs expected d/d genotype frequency (red)



a = 0.0688 & x = 0.54 observed d/wt genotype frequency (black) vs expected d/wt genotype frequency (red)

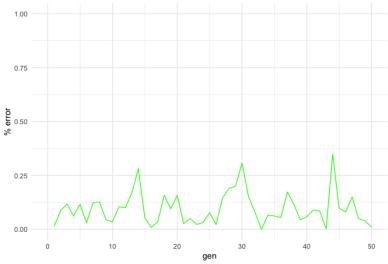


a = 0.0688 & x = 0.54 observed wt/wt genotype frequency (black) vs expected wt/wt genotype frequency (red)

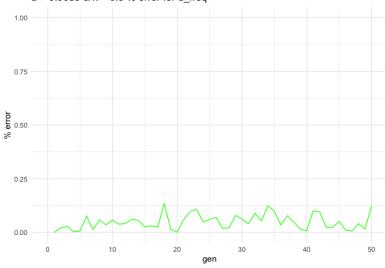


Percent errors in drive frequency predictions:





a = 0.0688 & x = 0.5 % error for d_freq



a = 0.0688 & x = 0.54 % error for d_freq

