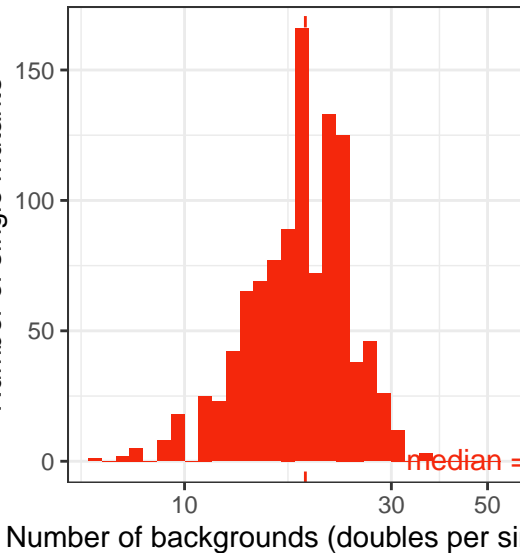


Number of single mutants



phenotype



binding

median =