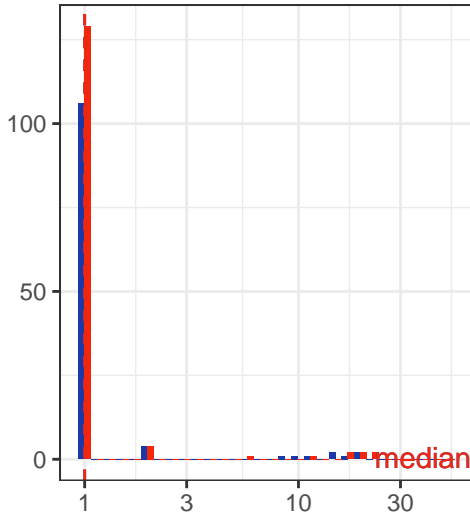


Number of single mutants



phenotype



abundance



binding

median

Number of backgrounds (doubles per single)