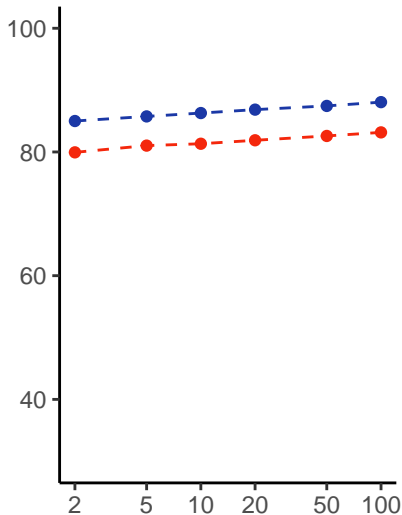


%Variance explained



phenotype

—●— abundance/folding

—●— binding

validation

-- test_fitness

%Retained double AA mutants