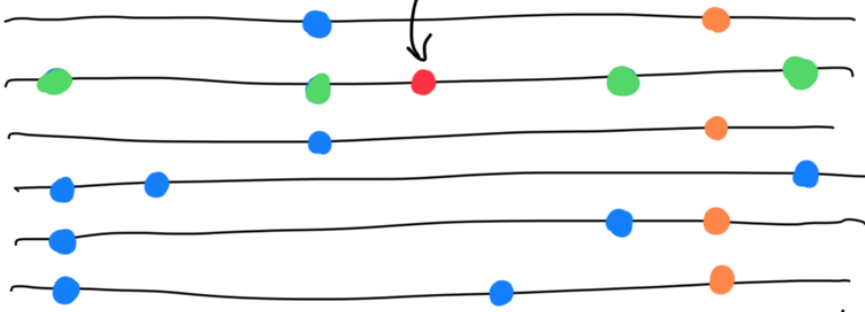


new beneficial mutation



not linked with
beneficial locus

