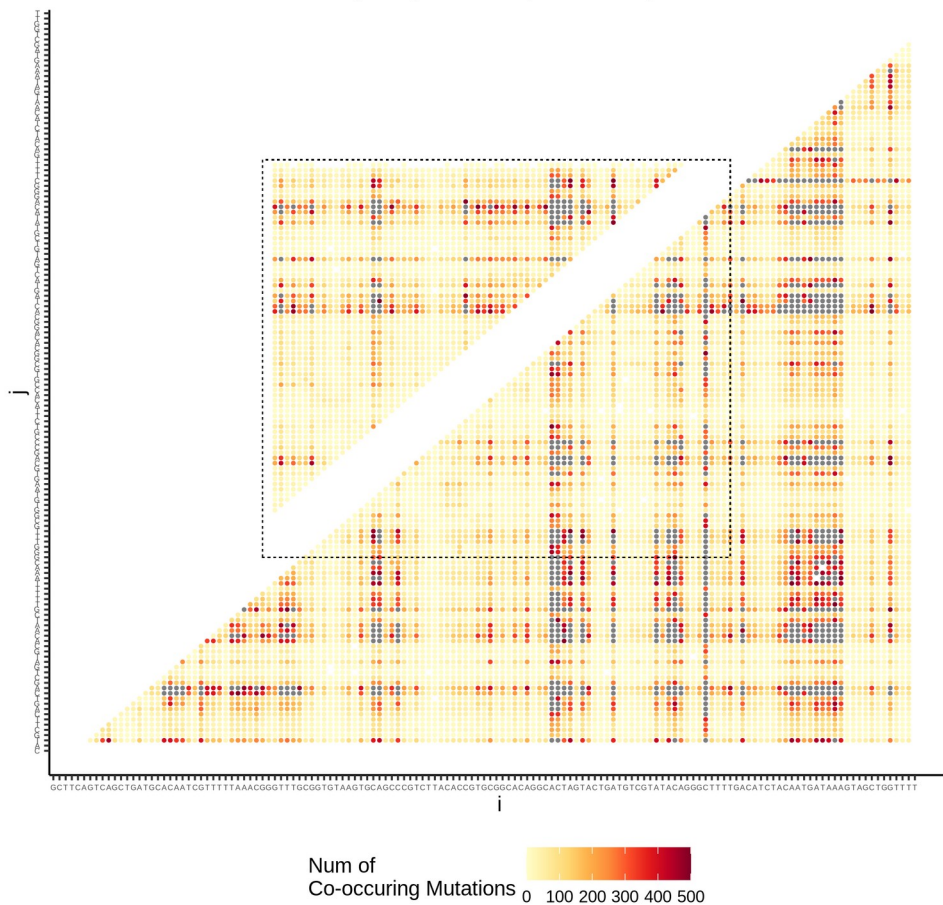


Raw mutation count. Tri-Sup: Experiment1, Tri-Inf: Experiment2



R.diff. Tri-Sup: Experiment1, Tri-Inf: Experiment2

