Fit of data for all possible interaction models on BRCA214 90000 -80000 Generalized Kullback–Leibler (GKL) 48 78 108 132 138 162 168 192 246 276 300 330 384 70000 -Signature parametrization Mono-nucleotide interaction 60000 -Di-nucleotide interaction Tri-nucleotide interaction 50000 -40000 -30000 12 Model