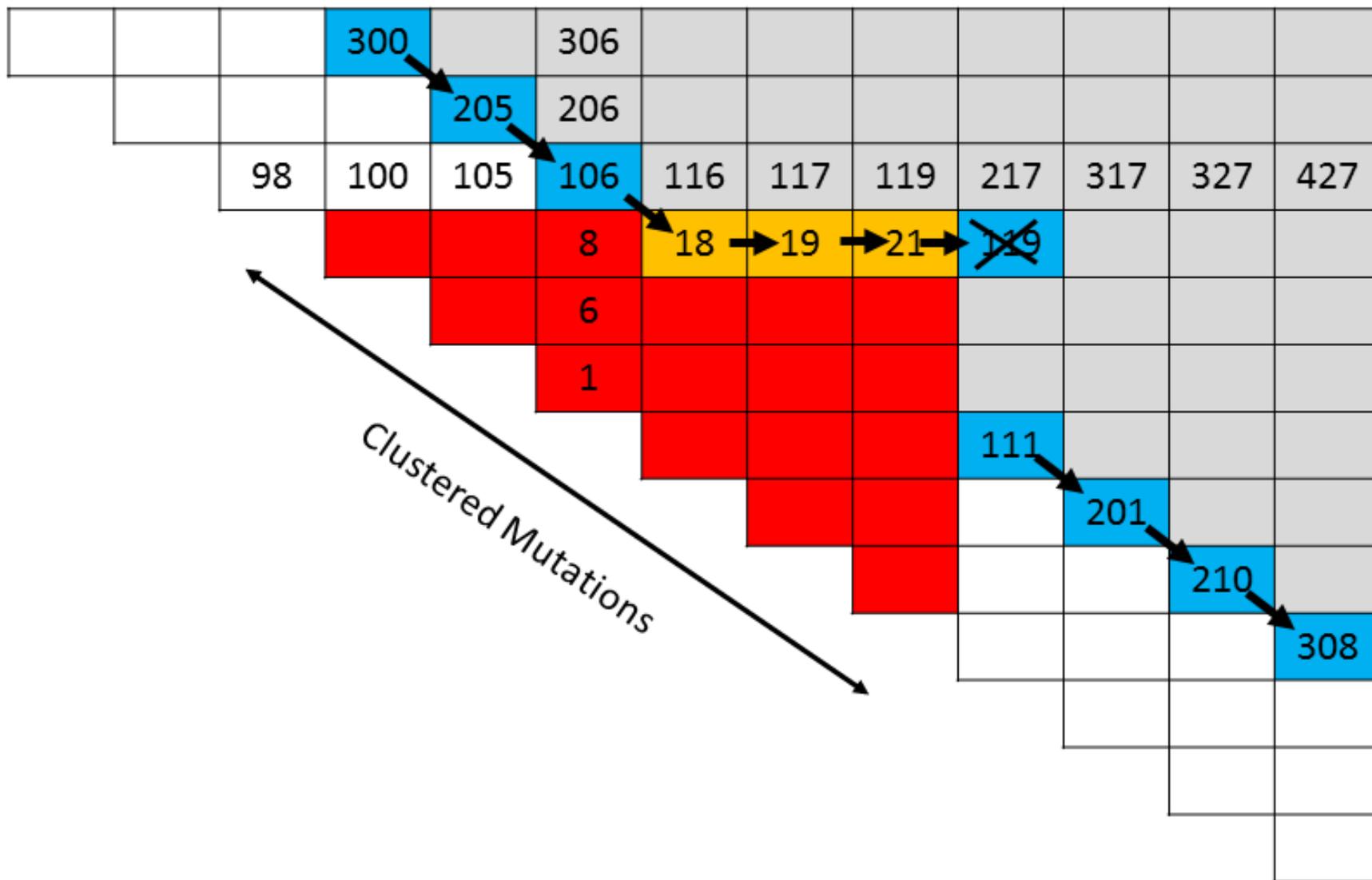


The position of somatic substitution mutations in the DNA of cancer cells.

1
101
201
299
301
306
307
317
318
320
418
518
528
628



Clustered Mutations

Symmetric dissimilarity mutation matrix. Each cell represents the distance in base pairs between the chromosomal position of somatic mutations. For this example, hyper-mutated region: min>=5, max=<100.