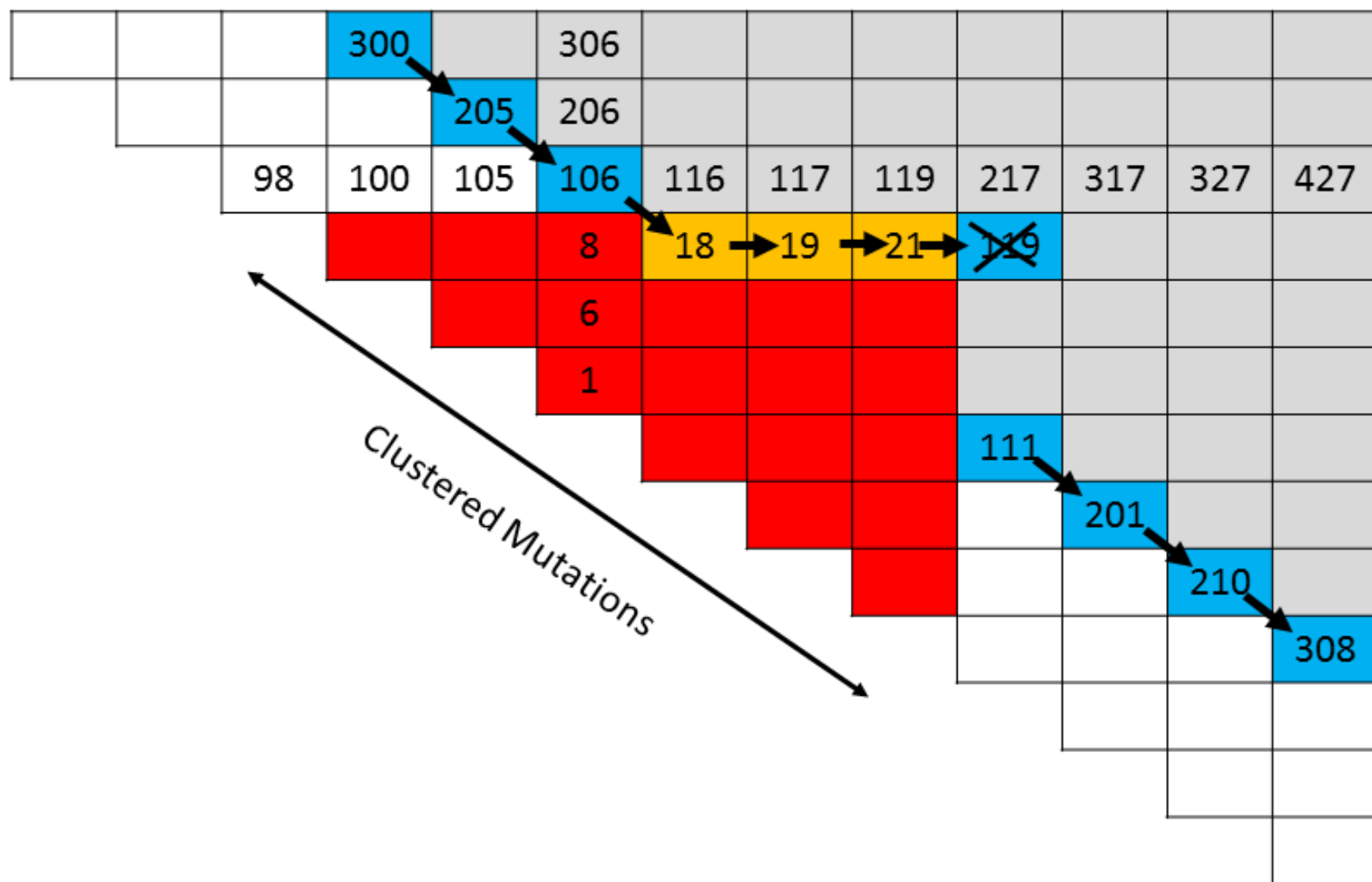


The position of somatic substitution mutations in the DNA of cancer cells.

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



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.