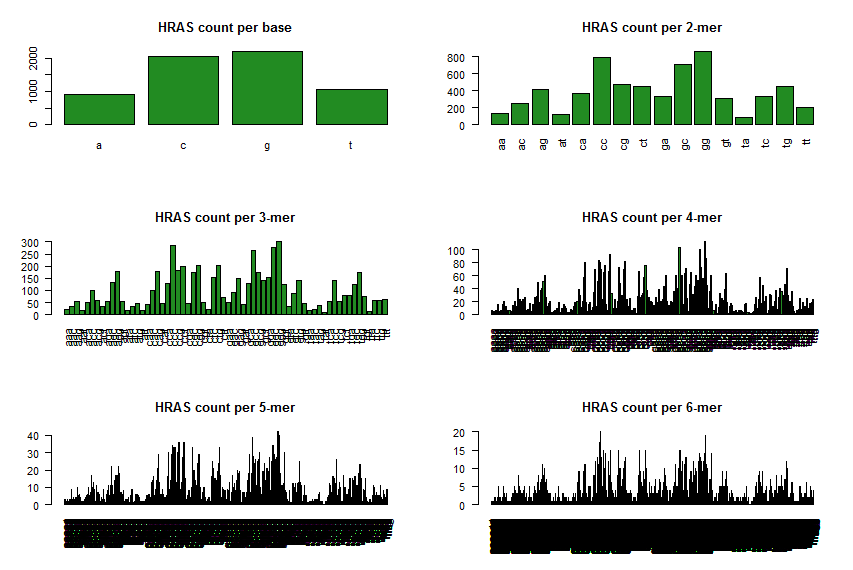
Digesting the 6 mer approach with HRAS, KRAS and TP53 genomic sequence

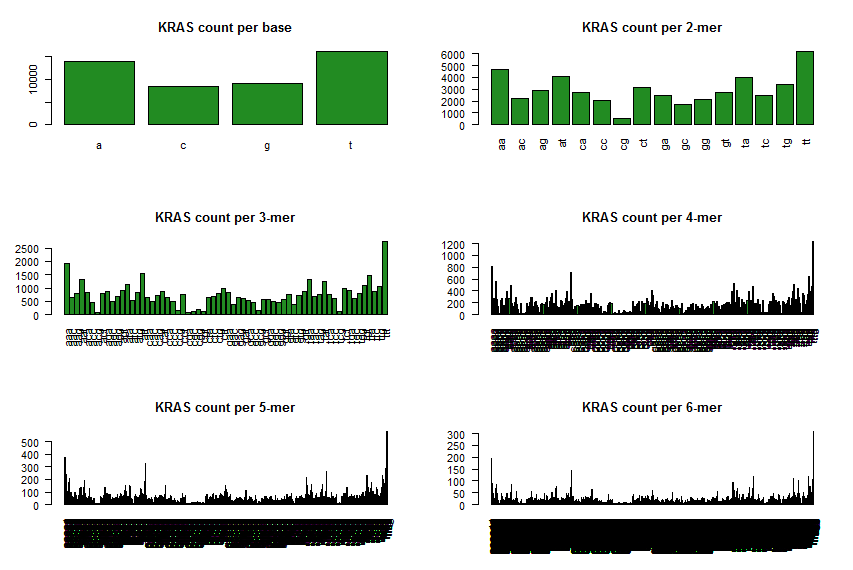
If I take a gene HRAS and plot the n mers frequency what do we see?

1. To do this I downloaded the seqinr package
2. Run the code in the readme

1MER TO 6 MER PLOTS X GENE

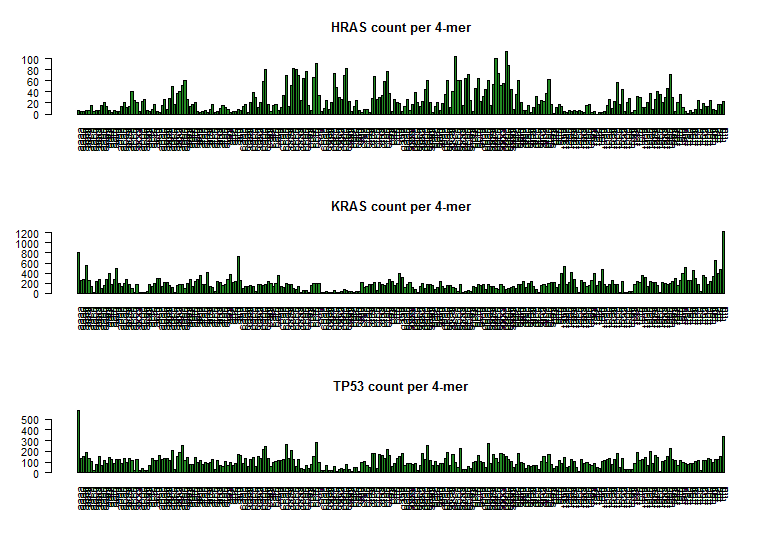
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4 MER PLOTS X GENE



6 MER PLOTS X GENE

