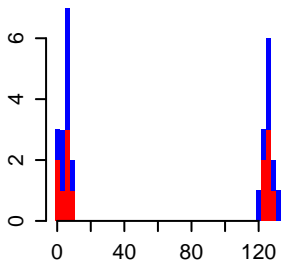
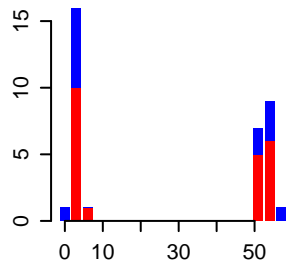


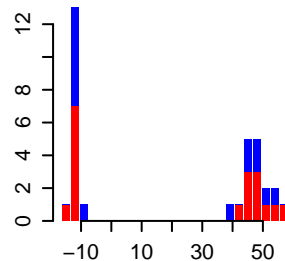
CTG: ATXN3:SCA3 coding  
chr14:92071010–92071040



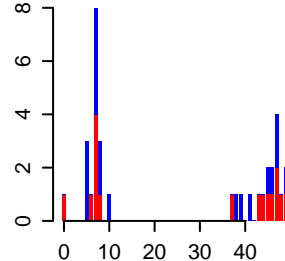
CAG: ATN1:DRPLA coding  
chr12:6936716–6936773



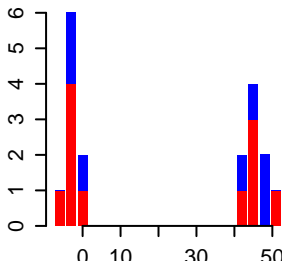
CAG: HTT:HD coding  
chr4:3074876–3074939



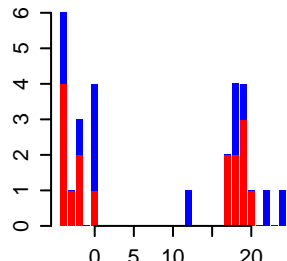
GCT: ATXN2:SCA2 coding  
chr12:111598950–111599019



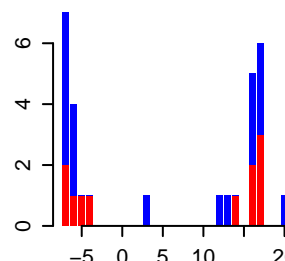
GCA: AR:SBMA coding  
chrX:67545317–67545386



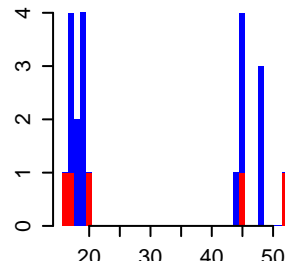
GCA: ATXN7:SCA7 coding  
chr3:63912685–63912715



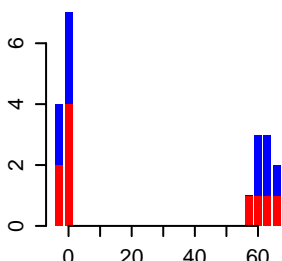
CAG: CACNA1A:SCA6 coding  
chr19:13207858–13207897



GCCCCG: C9orf72:FTDALS1 intron  
chr9:27573528–27573546



GCT: PPP2R2B:SCA12 intron  
chr5:146878728–146878758



CAGG: CNBP:DM2 intron  
chr3:129172576–129172656

