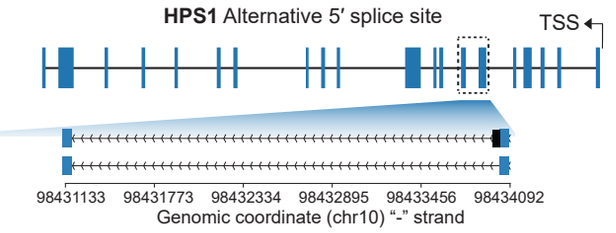
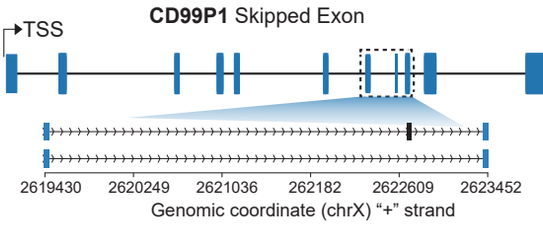
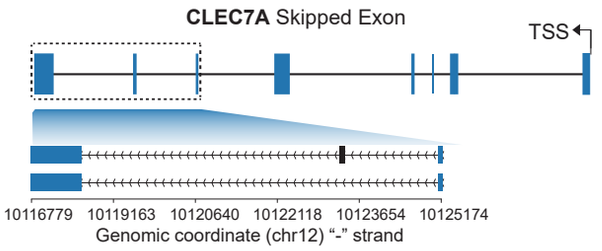
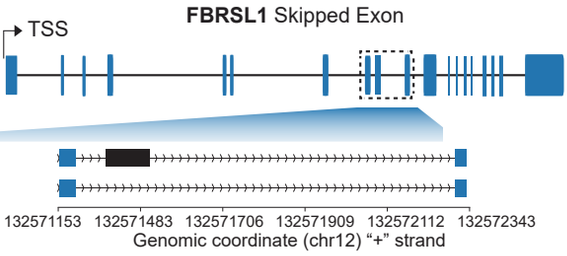
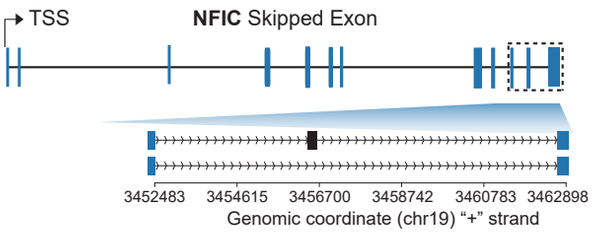
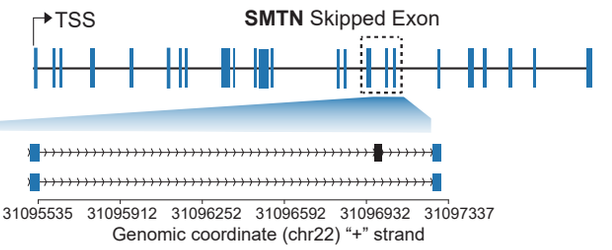

HPS1 Alternative 5′ splice site
TSS
98431133 98431773 98432334 98432895 98433456 98434092
Genomic coordinate (chr10) "-" strand

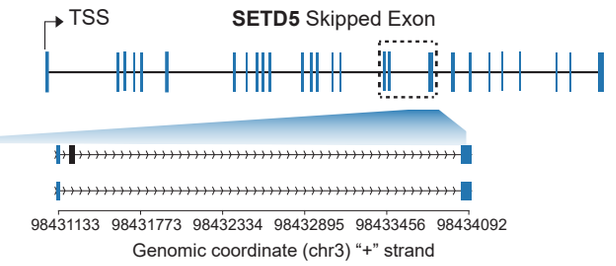

TSS
CD99P1 Skipped Exon
2619430 2620249 2621036 262182 2622609 2623452
Genomic coordinate (chrX) "+" strand

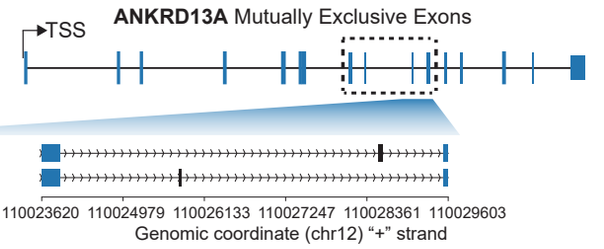

CLEC7A Skipped Exon
TSS
10116779 10119163 10120640 10122118 10123654 10125174
Genomic coordinate (chr12) "-" strand

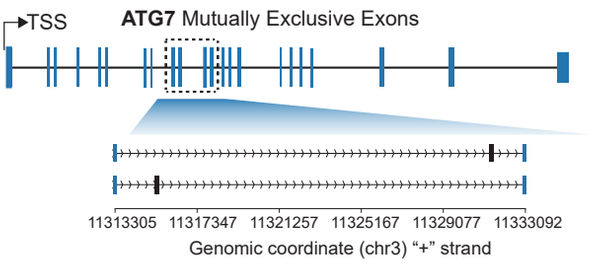

TSS
FBRSL1 Skipped Exon
132571153 132571483 132571706 132571909 132572112 132572343
Genomic coordinate (chr12) "+" strand

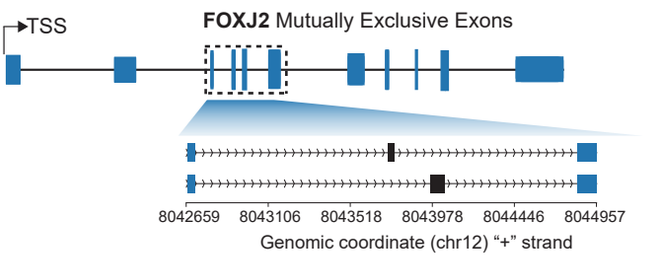

TSS
NFIC Skipped Exon
3452483 3454615 3456700 3458742 3460783 3462898
Genomic coordinate (chr19) "+" strand

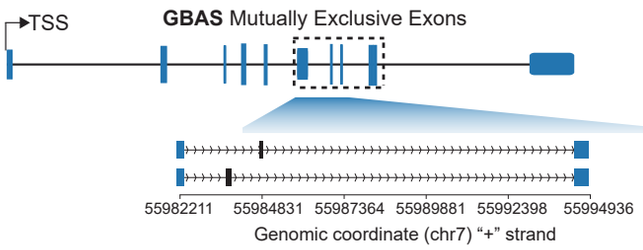

TSS
SMTN Skipped Exon
31095535 31095912 31096252 31096592 31096932 31097337
Genomic coordinate (chr22) "+" strand

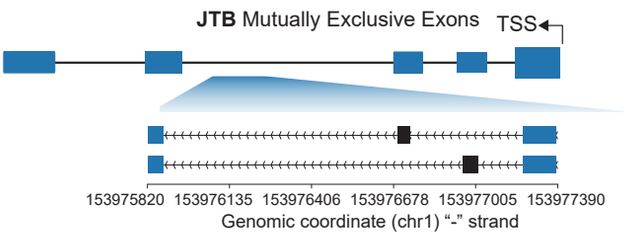

TSS
SETD5 Skipped Exon
98431133 98431773 98432334 98432895 98433456 98434092
Genomic coordinate (chr3) "+" strand

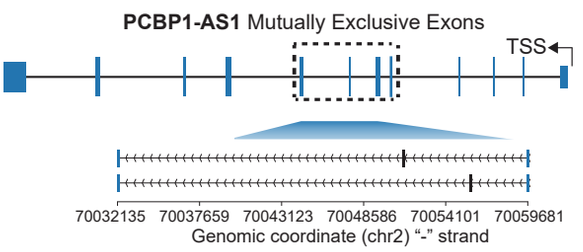

TSS
ANKRD13A Mutually Exclusive Exons
110023620 110024979 110026133 110027247 110028361 110029603
Genomic coordinate (chr12) "+" strand

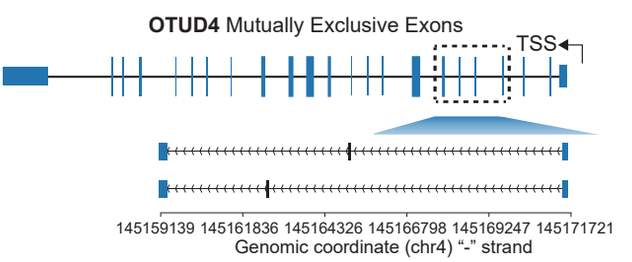

TSS
ATG7 Mutually Exclusive Exons
11313305 11317347 11321257 11325167 11329077 11333092
Genomic coordinate (chr3) "+" strand

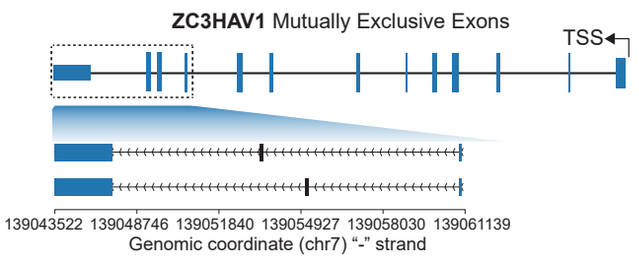

TSS
FOXJ2 Mutually Exclusive Exons
8042659 8043106 8043518 8043978 8044446 8044957
Genomic coordinate (chr12) "+" strand


TSS
GBAS Mutually Exclusive Exons
55982211 55984831 55987364 55989881 55992398 55994936
Genomic coordinate (chr7) "+" strand


JTB Mutually Exclusive Exons
TSS
153975820 153976135 153976406 153976678 153977005 153977390
Genomic coordinate (chr1) "-" strand


PCBP1-AS1 Mutually Exclusive Exons
TSS
70032135 70037659 70043123 70048586 70054101 70059681
Genomic coordinate (chr2) "-" strand


OTUD4 Mutually Exclusive Exons
TSS
145159139 145161836 145164326 145166798 145169247 145171721
Genomic coordinate (chr4) "-" strand


ZC3HAV1 Mutually Exclusive Exons
TSS
139043522 139048746 139051840 139054927 139058030 139061139
Genomic coordinate (chr7) "-" strand