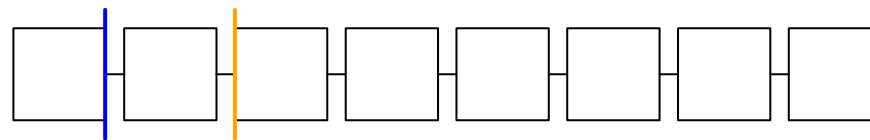


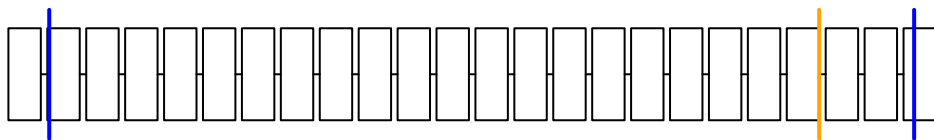
MGA--NUTM1

Spanning.pairs=146
Spanning.unique.reads=37
Predicted.effect=in-frame

GeneA.transcript=ENST00000219905
GeneB.transcript=ENST00000537011



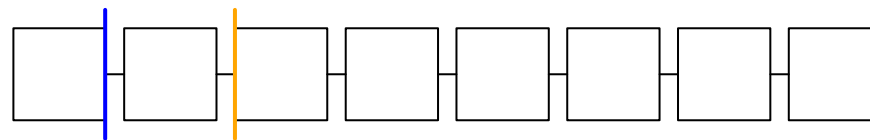
Will add some text to geneB plot later on,
probably related to protein domains



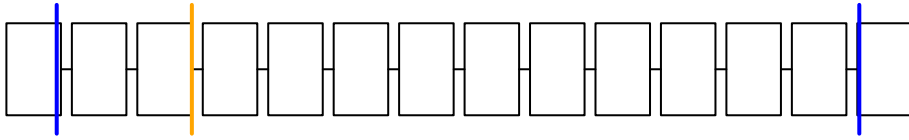
MGA--NUTM1

Spanning.pairs=146
Spanning.unique.reads=4
Predicted.effect=in-frame

GeneA.transcript=ENST00000219905
GeneB.transcript=ENST00000537011



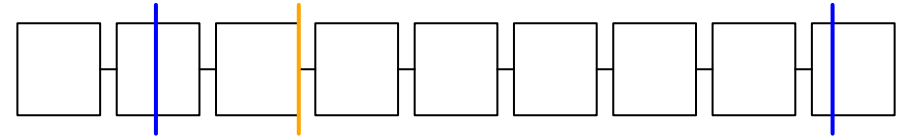
Will add some text to geneB plot later on,
probably related to protein domains



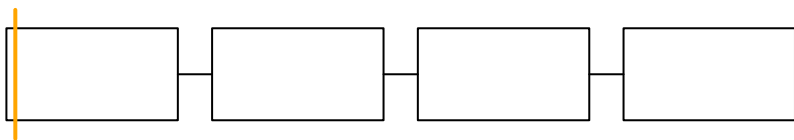
DPYSL2--KRT23

Spanning.pairs=11
Spanning.unique.reads=6
Predicted.effect=out-of-frame

GeneA.transcript=ENST00000311151
GeneB.transcript=ENST00000209718



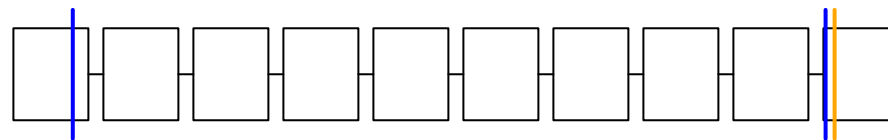
Will add some text to geneB plot later on,
probably related to protein domains



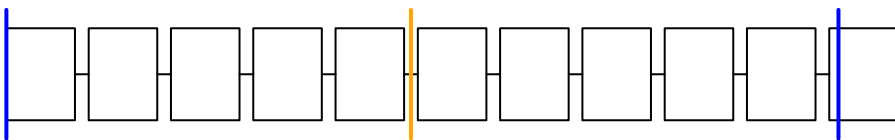
ANAPC7--KDSR

Spanning.pairs=9
Spanning.unique.reads=5
Predicted.effect=UTR/UTR

GeneA.transcript=ENST00000548234
GeneB.transcript=ENST00000406396



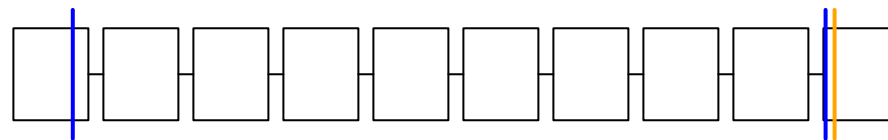
Will add some text to geneB plot later on,
probably related to protein domains



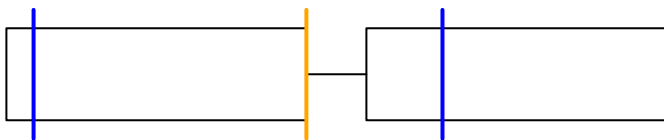
ANAPC7--KDSR

Spanning.pairs=9
Spanning.unique.reads=2
Predicted.effect=intronic/UTR

GeneA.transcript=ENST00000455511
GeneB.transcript=ENST00000406396



Will add some text to geneB plot later on,
probably related to protein domains



HOXC10--FLJ12825

no_protein,lincrna,10K<gap<100K

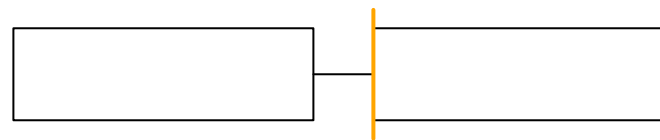
Spanning.pairs=9

Spanning.unique.reads=8

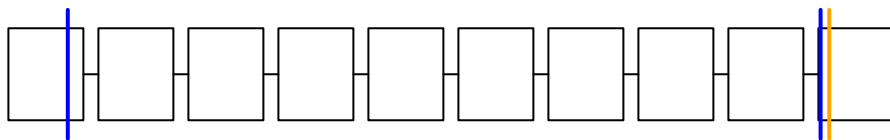
Predicted.effect=CDS(truncated)/exonic(no-known-CDS)

GeneA.transcript=ENST00000303460

GeneB.transcript=ENST00000515617



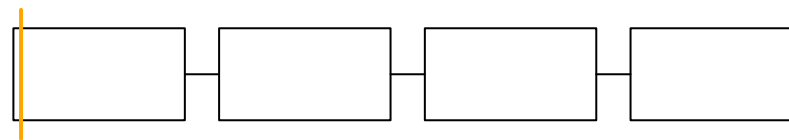
Will add some text to geneB plot later on,
probably related to protein domains



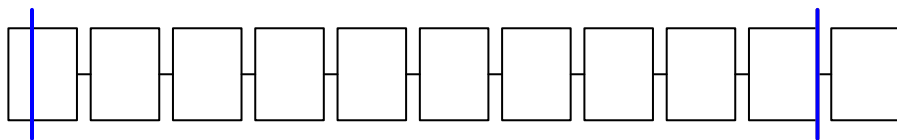
KDSR--ANAPC7

Spanning.pairs=9
Spanning.unique.reads=3
Predicted.effect=UTR/UTR

GeneA.transcript=ENST00000406396
GeneB.transcript=ENST00000548234



Will add some text to geneB plot later on,
probably related to protein domains

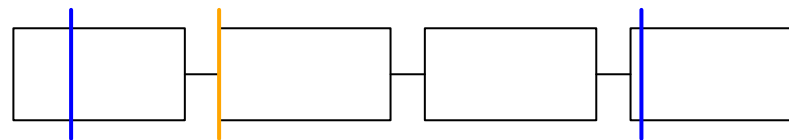


CPSF6--LYZ

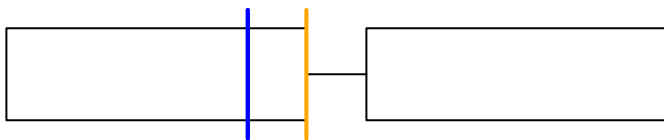
10K<gap<100K

Spanning.pairs=6
Spanning.unique.reads=4
Predicted.effect=out-of-frame

GeneA.transcript=ENST00000266679
GeneB.transcript=ENST00000261267



Will add some text to geneB plot later on,
probably related to protein domains

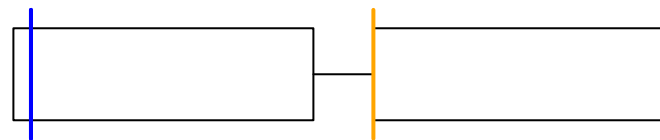


HIST1H2AC--HIST1H2BD

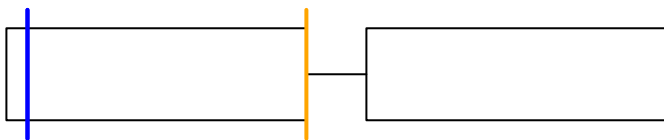
10K<gap<100K

Spanning.pairs=6
Spanning.unique.reads=3
Predicted.effect=UTR/UTR

GeneA.transcript=ENST00000314088
GeneB.transcript=ENST00000289316



Will add some text to geneB plot later on,
probably related to protein domains

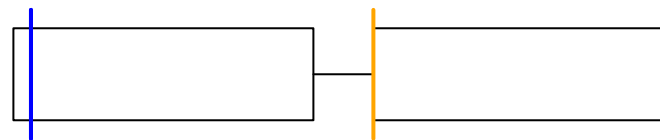


HIST1H2AC--HIST1H2BD

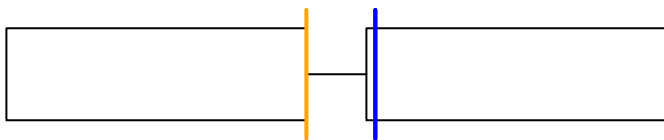
10K<gap<100K

Spanning.pairs=6
Spanning.unique.reads=2
Predicted.effect=UTR/UTR

GeneA.transcript=ENST00000602637
GeneB.transcript=ENST00000289316



Will add some text to geneB plot later on,
probably related to protein domains



MRPS6--AP000318.2

no_protein,lincrna,ribosomal_protein,10K<gap<100K

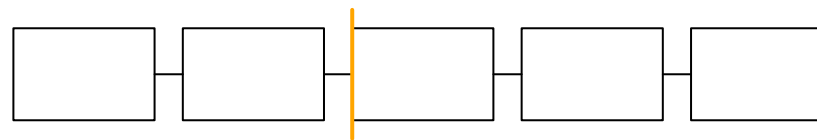
Spanning.pairs=6

Spanning.unique.reads=1

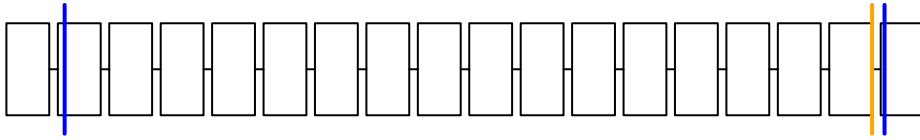
Predicted.effect=CDS(truncated)/exonic(no-known-CDS)

GeneA.transcript=ENST00000608209

GeneB.transcript=ENST00000427022



Will add some text to geneB plot later on,
probably related to protein domains

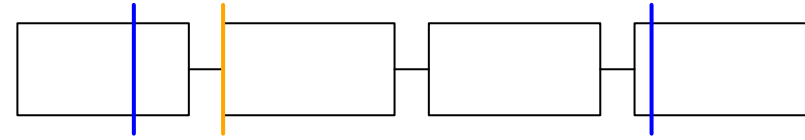


PIBF1--KLF5

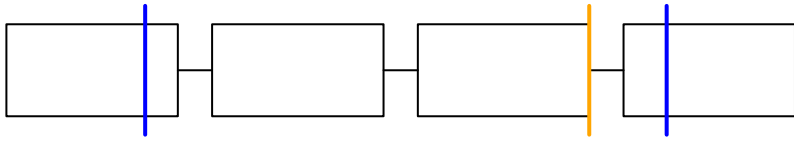
10K<gap<100K

Spanning.pairs=6
Spanning.unique.reads=4
Predicted.effect=in-frame

GeneA.transcript=ENST00000326291
GeneB.transcript=ENST00000377687



Will add some text to geneB plot later on,
probably related to protein domains



CLN5--MYCBP2-AS1

no_protein,antisense,10K<gap<100K

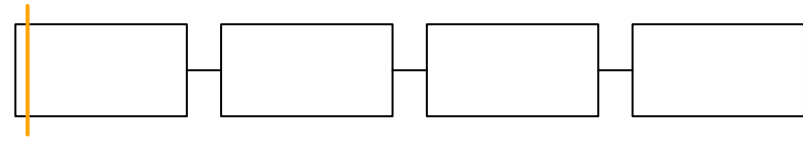
Spanning.pairs=4

Spanning.unique.reads=2

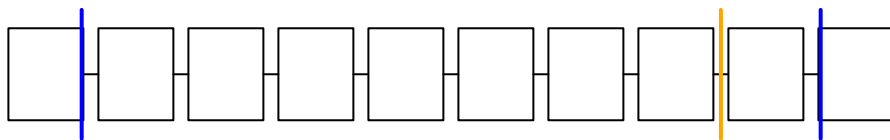
Predicted.effect=CDS(truncated)/exonic(no-known-CDS)

GeneA.transcript=ENST00000377453

GeneB.transcript=ENST00000422231



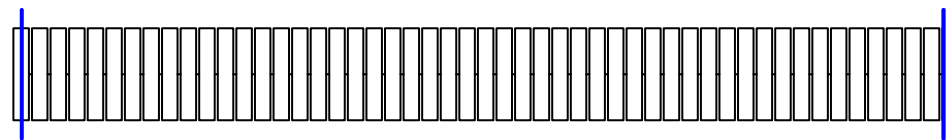
Will add some text to geneB plot later on,
probably related to protein domains



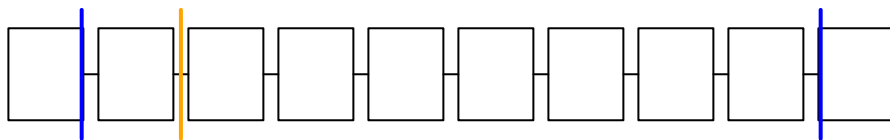
COLEC12--COL1A1

Spanning.pairs=4
Spanning.unique.reads=2
Predicted.effect=intronic/UTR

GeneA.transcript=ENST00000400256
GeneB.transcript=ENST00000225964



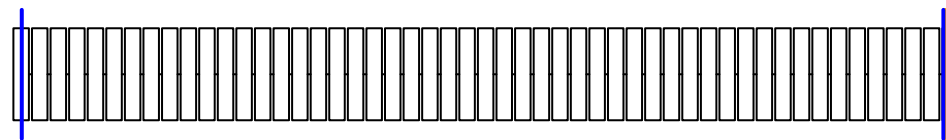
Will add some text to geneB plot later on,
probably related to protein domains



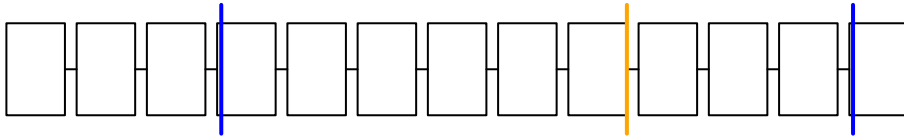
COLEC12--COL1A1

Spanning.pairs=4
Spanning.unique.reads=2
Predicted.effect=intronic/UTR

GeneA.transcript=ENST00000400256
GeneB.transcript=ENST00000225964



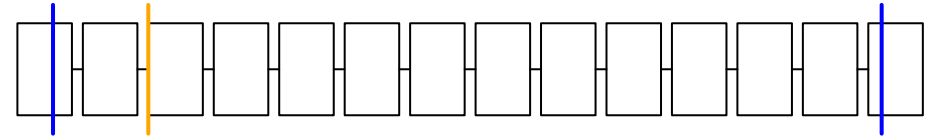
Will add some text to geneB plot later on,
probably related to protein domains



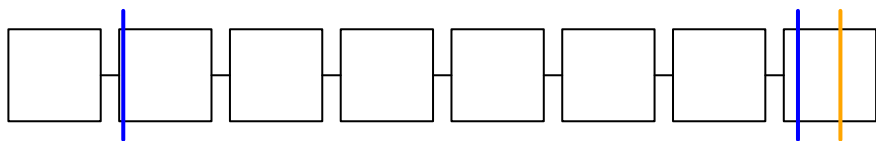
DLGAP1--PARVB

Spanning.pairs=4
Spanning.unique.reads=2
Predicted.effect=out-of-frame

GeneA.transcript=ENST00000315677
GeneB.transcript=ENST00000406477



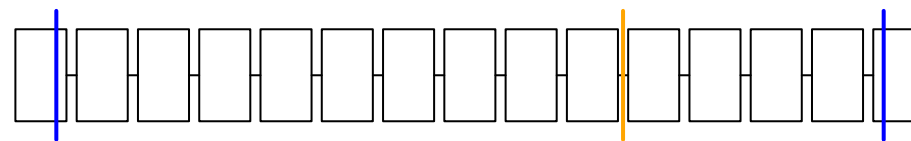
Will add some text to geneB plot later on,
probably related to protein domains



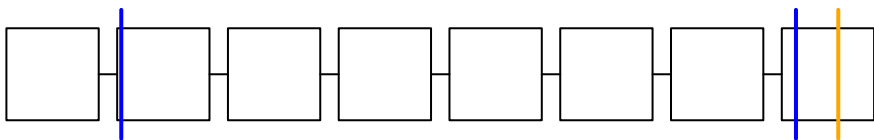
BGN--SH3PXD2A

Spanning.pairs=3
Spanning.unique.reads=2
Predicted.effect=UTR/intronic

GeneA.transcript=ENST00000331595
GeneB.transcript=ENST00000369774



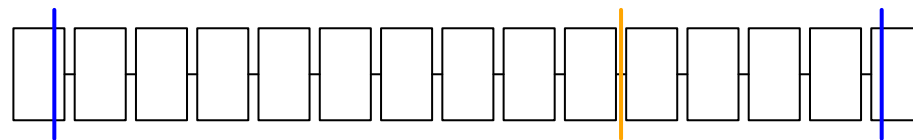
Will add some text to geneB plot later on,
probably related to protein domains



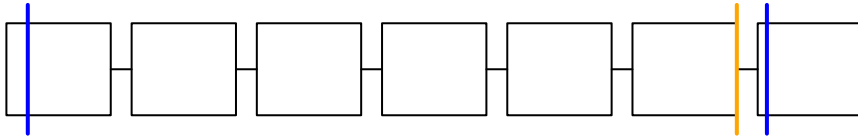
BGN--SH3PXD2A

Spanning.pairs=3
Spanning.unique.reads=2
Predicted.effect=UTR/intronic

GeneA.transcript=ENST00000331595
GeneB.transcript=ENST00000369774



Will add some text to geneB plot later on,
probably related to protein domains

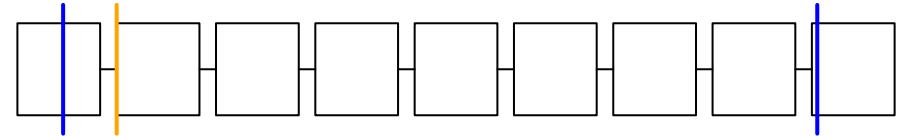


C11ORF70--YAP1

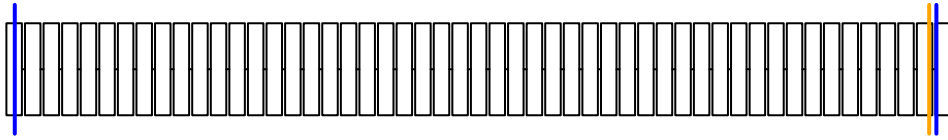
tcga, cell_lines, prostates, 10K < gap < 100K

Spanning.pairs=3
Spanning.unique.reads=1
Predicted.effect=in-frame

GeneA.transcript=ENST00000434758
GeneB.transcript=ENST00000282441



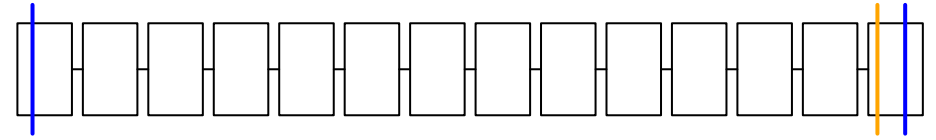
Will add some text to geneB plot later on,
probably related to protein domains



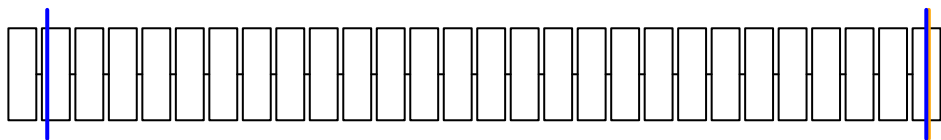
COL1A1--CPXM1

Spanning.pairs=3
Spanning.unique.reads=1
Predicted.effect=in-frame

GeneA.transcript=ENST00000225964
GeneB.transcript=ENST00000380605



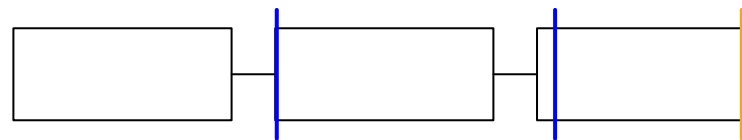
Will add some text to geneB plot later on,
probably related to protein domains



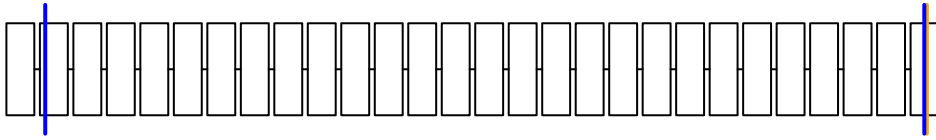
COL6A2--FMOD

Spanning.pairs=3
Spanning.unique.reads=1
Predicted.effect=UTR/UTR

GeneA.transcript=ENST00000357838
GeneB.transcript=ENST00000354955



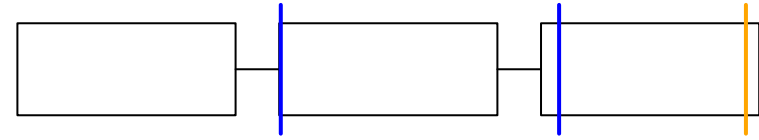
Will add some text to geneB plot later on,
probably related to protein domains



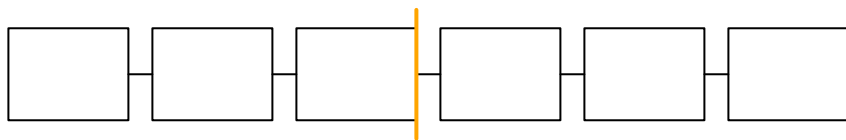
COL6A2--FMOD

Spanning.pairs=3
Spanning.unique.reads=1
Predicted.effect=UTR/UTR

GeneA.transcript=ENST00000357838
GeneB.transcript=ENST00000354955



Will add some text to geneB plot later on,
probably related to protein domains



MIR4435-2HG--AC093616.4

no_protein,pseudogene,lincrna

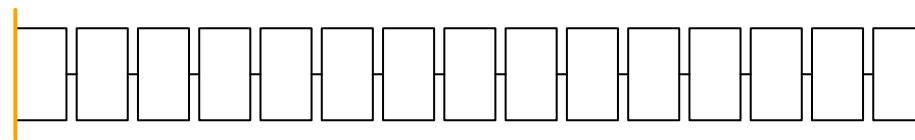
Spanning.pairs=3

Spanning.unique.reads=5

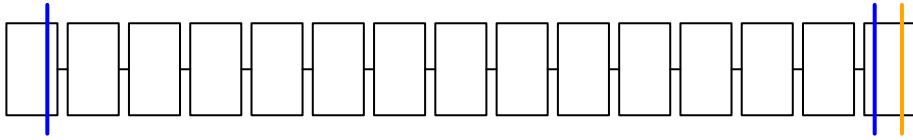
Predicted.effect=exonic(no-known-CDS)/exonic(no-known-CDS)

GeneA.transcript=ENST00000439362

GeneB.transcript=ENST00000427434



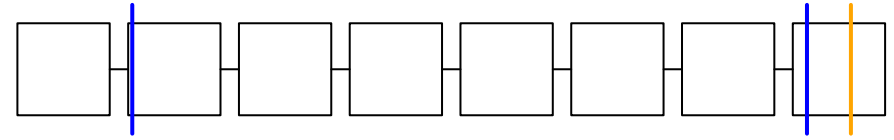
Will add some text to geneB plot later on,
probably related to protein domains



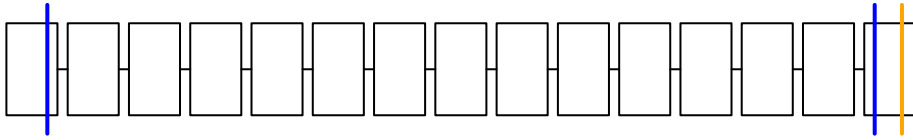
SH3PXD2A--BGN

Spanning.pairs=3
Spanning.unique.reads=2
Predicted.effect=UTR/UTR

GeneA.transcript=ENST00000369774
GeneB.transcript=ENST00000331595



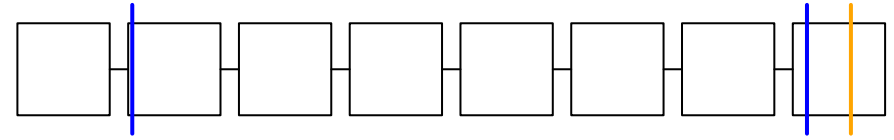
Will add some text to geneB plot later on,
probably related to protein domains



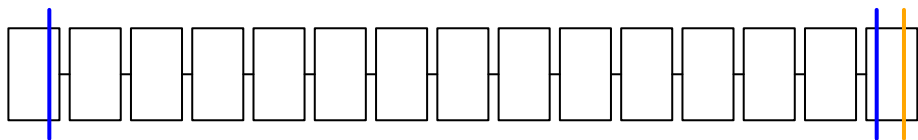
SH3PXD2A--BGN

Spanning.pairs=3
Spanning.unique.reads=2
Predicted.effect=UTR/UTR

GeneA.transcript=ENST00000369774
GeneB.transcript=ENST00000331595



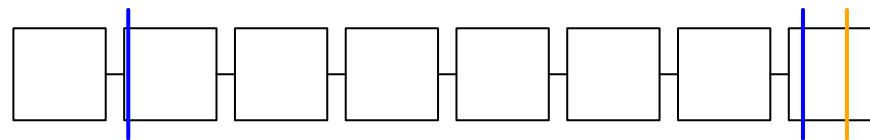
Will add some text to geneB plot later on,
probably related to protein domains



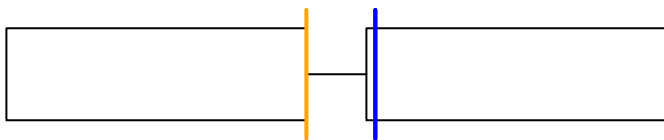
SH3PXD2A--BGN

Spanning.pairs=3
Spanning.unique.reads=2
Predicted.effect=UTR/UTR

GeneA.transcript=ENST00000369774
GeneB.transcript=ENST00000331595



Will add some text to geneB plot later on,
probably related to protein domains



MRPS6--AP000318.2

no_protein,lincrna,ribosomal_protein,10K<gap<100K

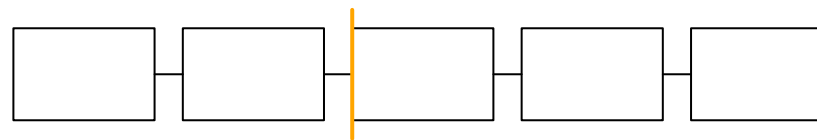
Spanning.pairs=6

Spanning.unique.reads=1

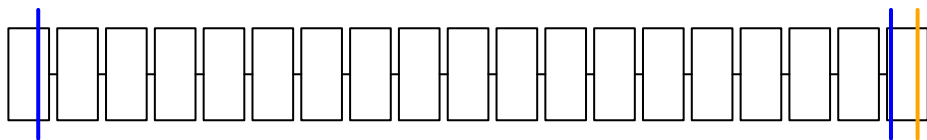
Predicted.effect=CDS(truncated)/exonic(no-known-CDS)

GeneA.transcript=ENST00000608209

GeneB.transcript=ENST00000427022



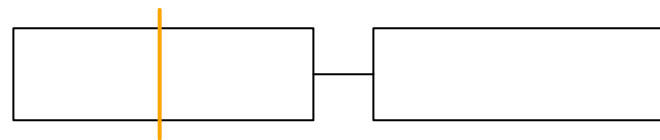
Will add some text to geneB plot later on,
probably related to protein domains



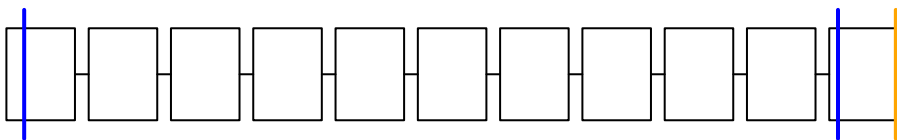
SPEF2--LTBR

Spanning.pairs=3
Spanning.unique.reads=2
Predicted.effect=UTR/UTR

GeneA.transcript=ENST00000509059
GeneB.transcript=ENST00000441074



Will add some text to geneB plot later on,
probably related to protein domains



TNFSF12–TNFSF13--MPDU1

10K<gap<100K

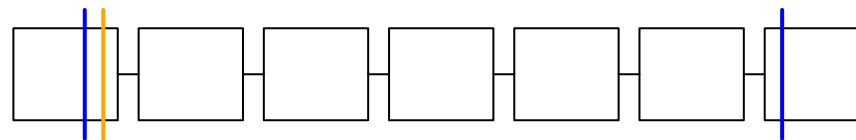
Spanning.pairs=3

Spanning.unique.reads=2

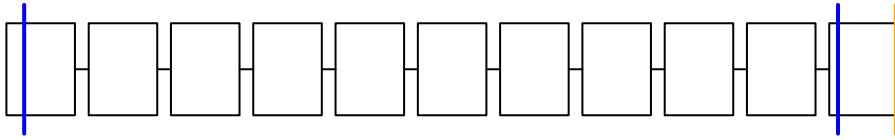
Predicted.effect=UTR/CDS(truncated)

GeneA.transcript=ENST00000293826

GeneB.transcript=ENST00000250124



Will add some text to geneB plot later on,
probably related to protein domains



TNFSF12–TNFSF13--MPDU1

10K<gap<100K

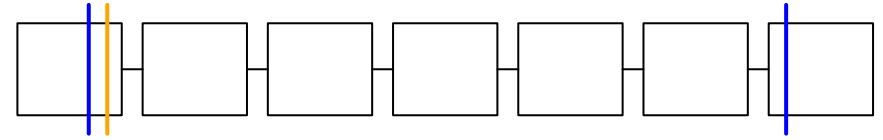
Spanning.pairs=3

Spanning.unique.reads=2

Predicted.effect=UTR/CDS(truncated)

GeneA.transcript=ENST00000293826

GeneB.transcript=ENST00000250124



Will add some text to geneB plot later on,
probably related to protein domains