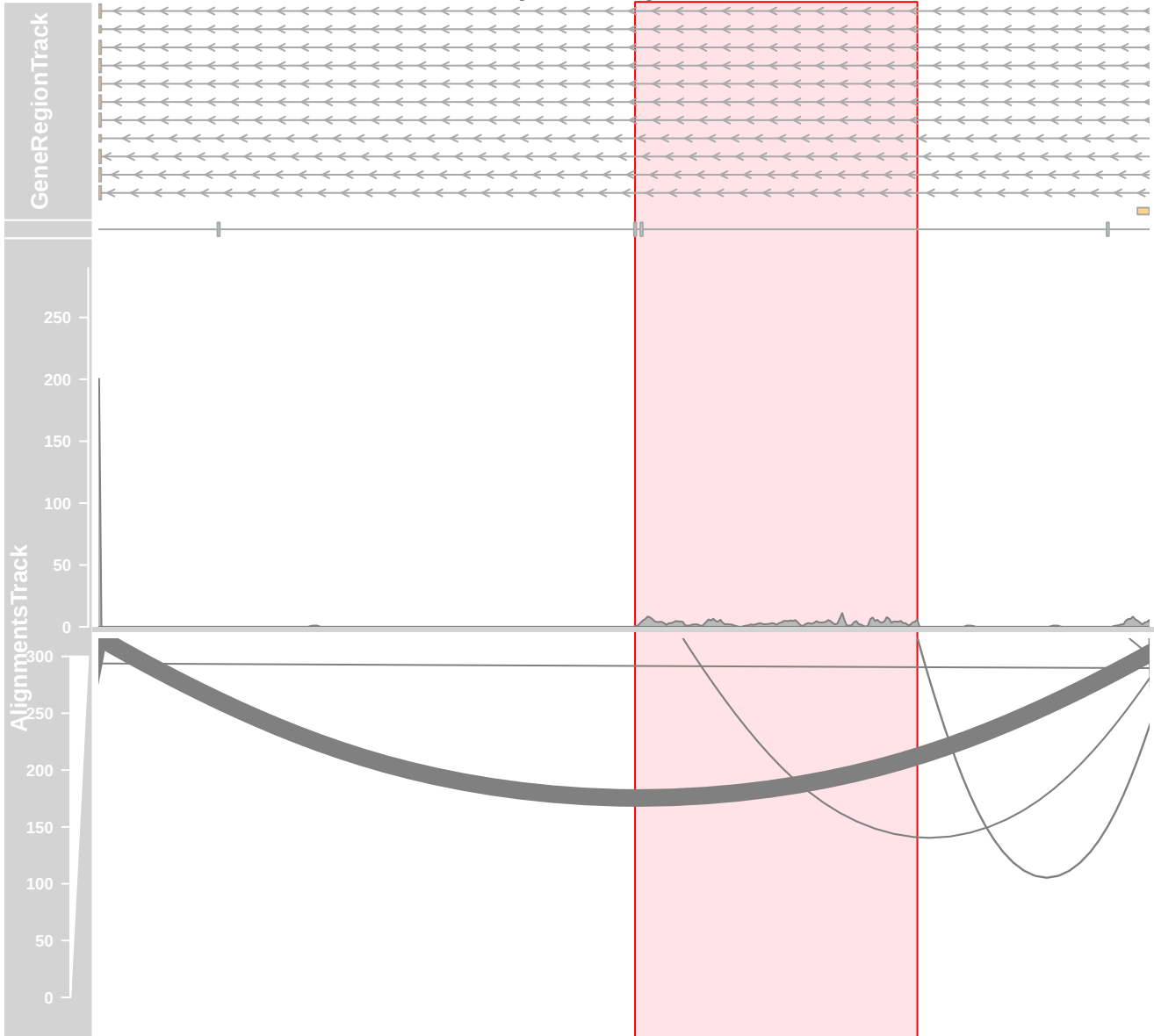


Region: 22:19899049–19911071

Novel exon: 22:19905184–19908418 Strand: -

Terminal probability: 0.9999

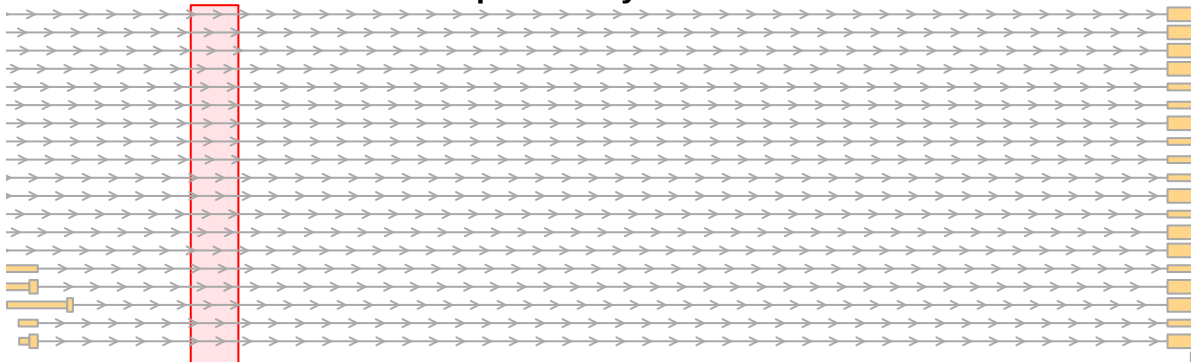


Region: 22:23857919–23862202

Novel exon: 22:23858580–23858751 Strand: +

Terminal probability: 0.8004

GeneRegionTrack



AlignmentsTrack

