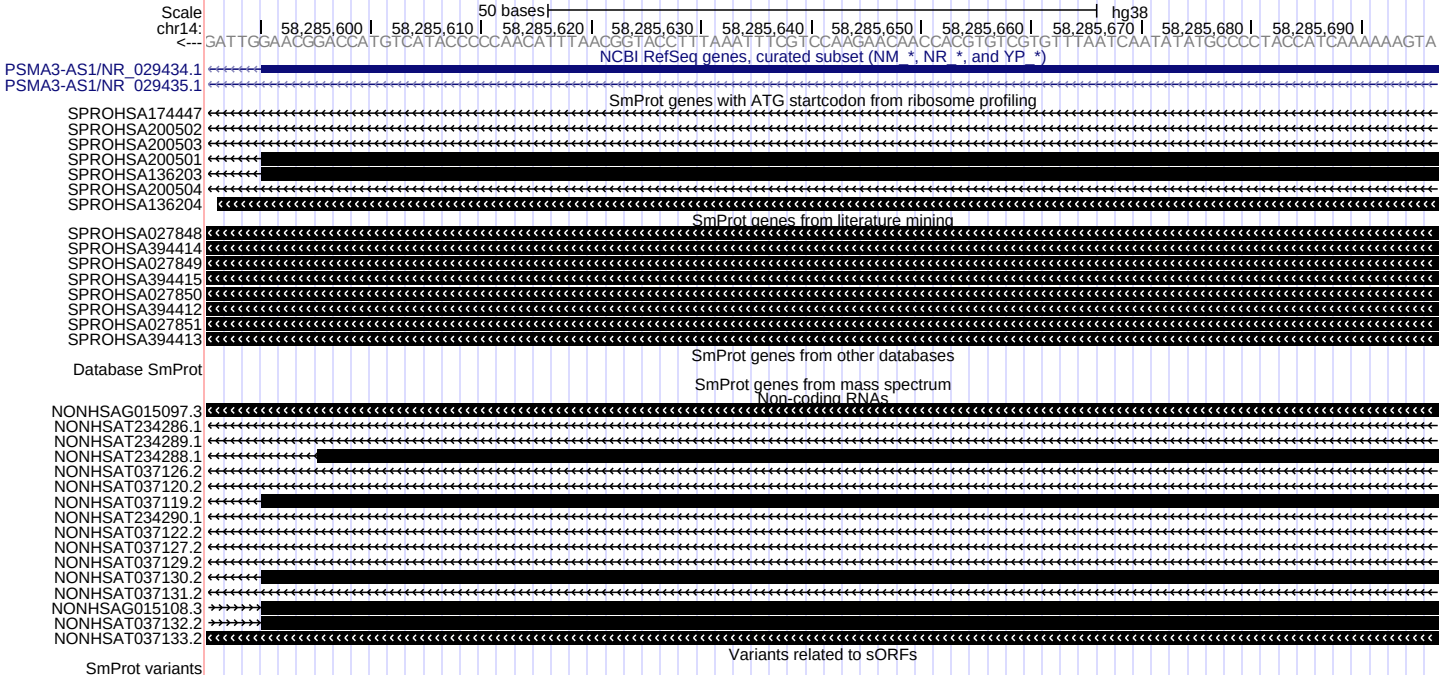
Scale
chr14:
50 bases
hg38
58,285,600
58,285,610
58,285,620
58,285,630
58,285,640
58,285,650
58,285,660
58,285,670
58,285,680
58,285,690
<---GATTGGAACGGACCATGTCATACCCCCAACATTTAACGGTACCTTTAAATTTCGTCCAAGAACAACCACGTGTCGTGTTTAATCAATATATGCCCCTACCATCAAAAAAGTA
NCBI RefSeq genes, curated subset (NM_*, NR_*, and YP_*)
PSMA3-AS1/NR_029434.1
PSMA3-AS1/NR_029435.1
SmProt genes with ATG startcodon from ribosome profiling
SPROHSA174447
SPROHSA200502
SPROHSA200503
SPROHSA200501
SPROHSA136203
SPROHSA200504
SPROHSA136204
SmProt genes from literature mining
SPROHSA027848
SPROHSA394414
SPROHSA027849
SPROHSA394415
SPROHSA027850
SPROHSA394412
SPROHSA027851
SPROHSA394413
Database SmProt
SmProt genes from other databases
SmProt genes from mass spectrum
Non-coding RNAs
NONHSAG015097.3
NONHSAT234286.1
NONHSAT234289.1
NONHSAT234288.1
NONHSAT037126.2
NONHSAT037120.2
NONHSAT037119.2
NONHSAT234290.1
NONHSAT037122.2
NONHSAT037127.2
NONHSAT037129.2
NONHSAT037130.2
NONHSAT037131.2
NONHSAG015108.3
NONHSAT037132.2
NONHSAT037133.2
SmProt variants
Variants related to sORFs