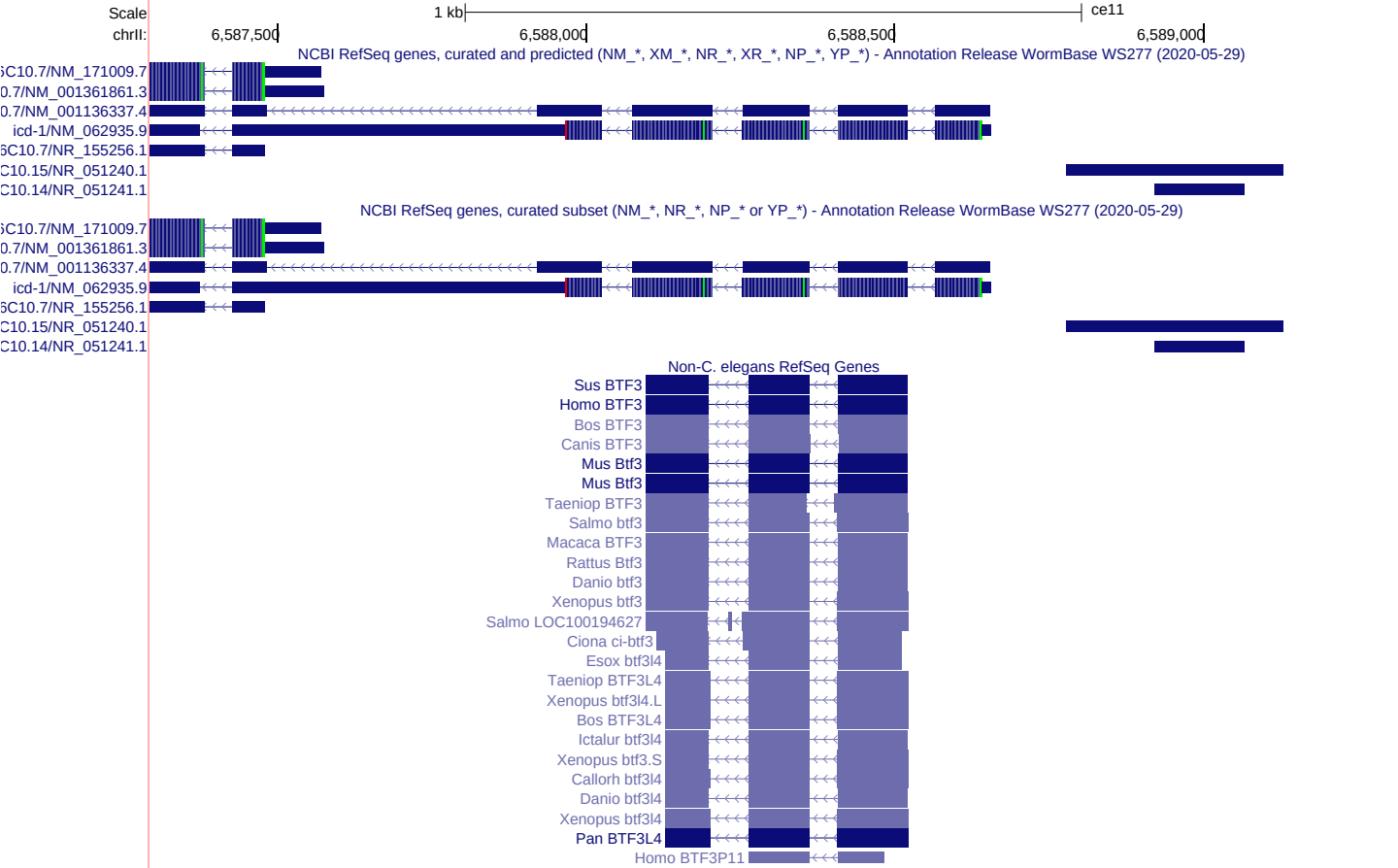
Scale
1 kb
ce11
chrII:
6,587,500
6,588,000
6,588,500
6,589,000
NCBI RefSeq genes, curated and predicted (NM_*, XM_*, NR_*, XR_*, NP_*, YP_*) - Annotation Release WormBase WS277 (2020-05-29)
6C10.7/NM_171009.7
0.7/NM_001361861.3
0.7/NM_001136337.4
icd-1/NM_062935.9
6C10.7/NR_155256.1
C10.15/NR_051240.1
C10.14/NR_051241.1
NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*) - Annotation Release WormBase WS277 (2020-05-29)
6C10.7/NM_171009.7
0.7/NM_001361861.3
0.7/NM_001136337.4
icd-1/NM_062935.9
6C10.7/NR_155256.1
C10.15/NR_051240.1
C10.14/NR_051241.1
Non-C. elegans RefSeq Genes
Sus BTF3
Homo BTF3
Bos BTF3
Canis BTF3
Mus Btf3
Mus Btf3
Taeniop BTF3
Salmo btf3
Macaca BTF3
Rattus Btf3
Danio btf3
Xenopus btf3
Salmo LOC100194627
Ciona ci-btf3
Esox btf3l4
Taeniop BTF3L4
Xenopus btf3l4.L
Bos BTF3L4
Ictalur btf3l4
Xenopus btf3.S
Callorh btf3l4
Danio btf3l4
Xenopus btf3l4
Pan BTF3L4
Homo BTF3P11