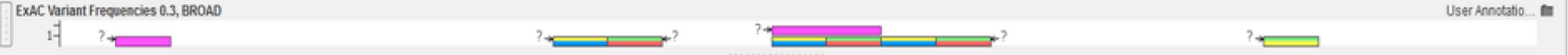
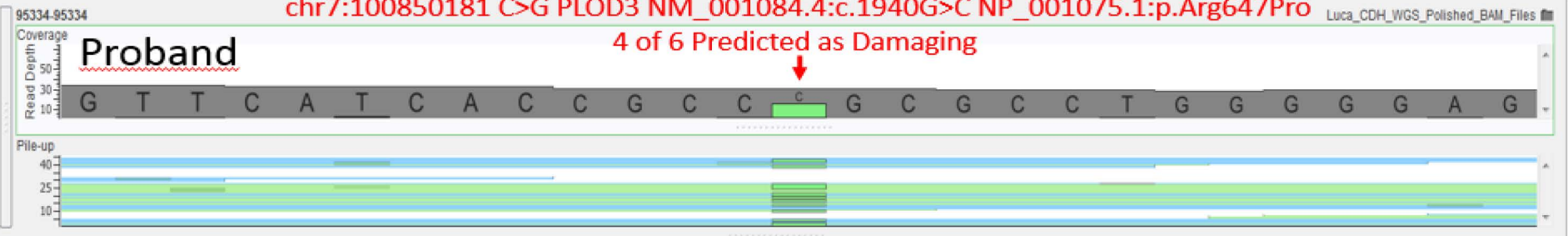
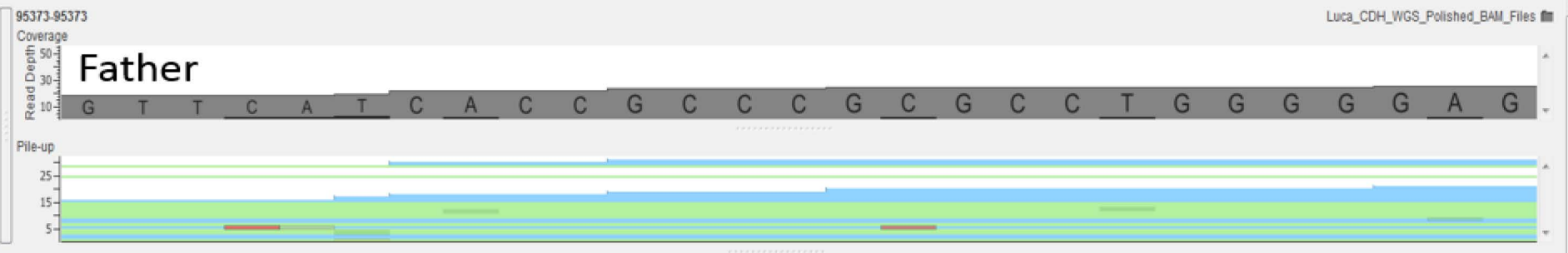
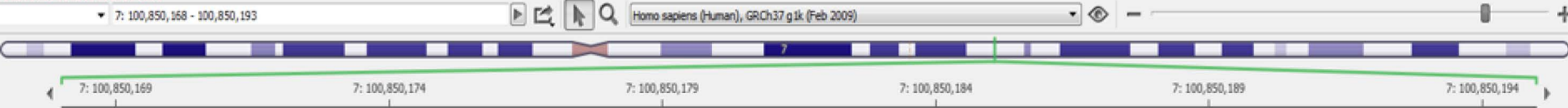
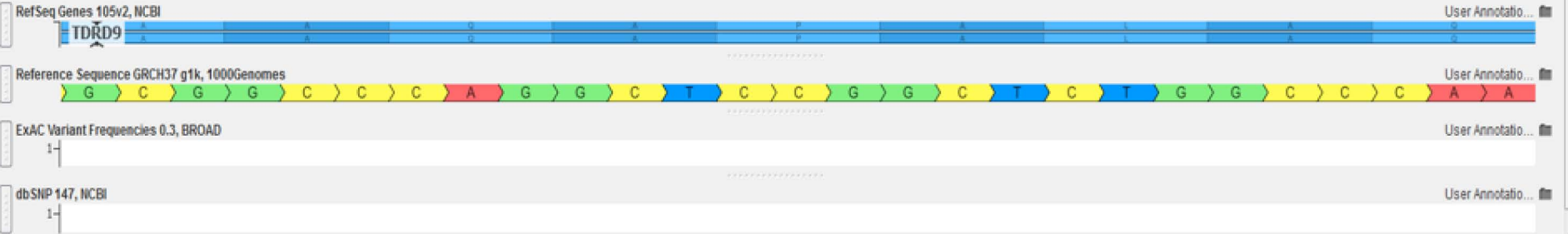
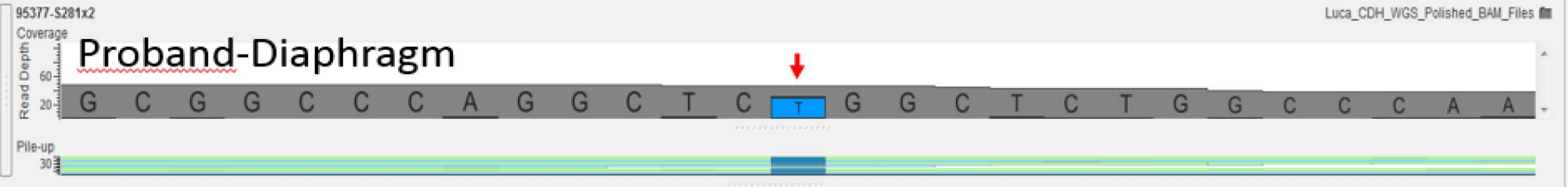
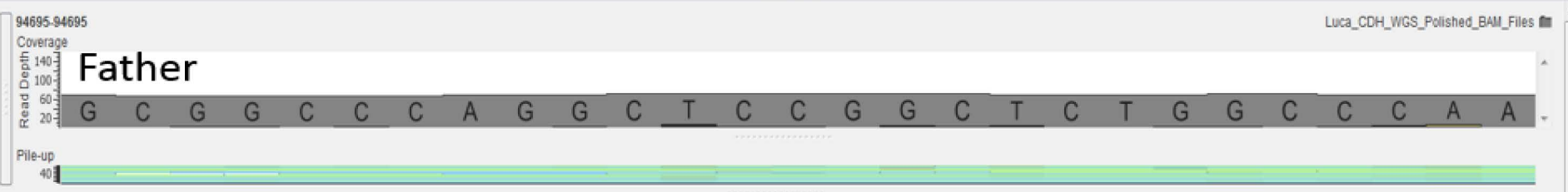
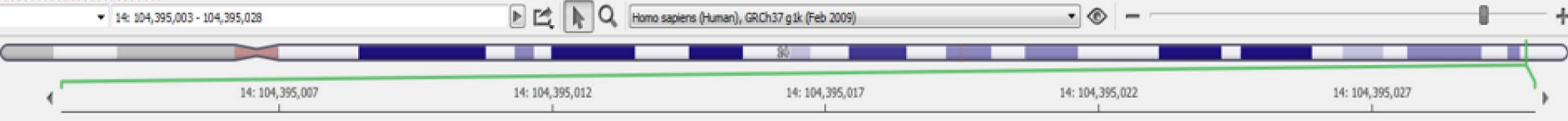


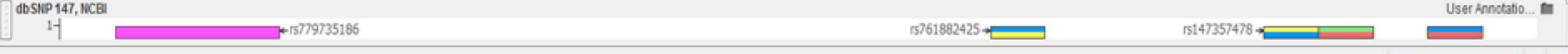
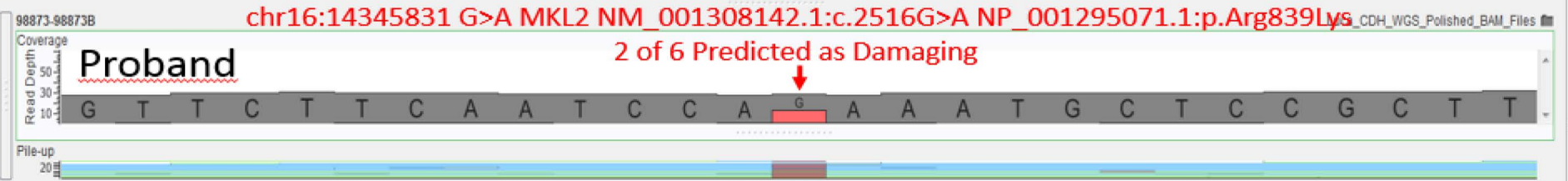
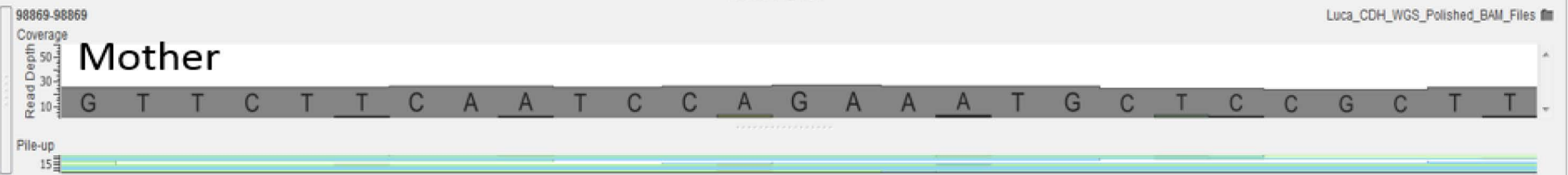
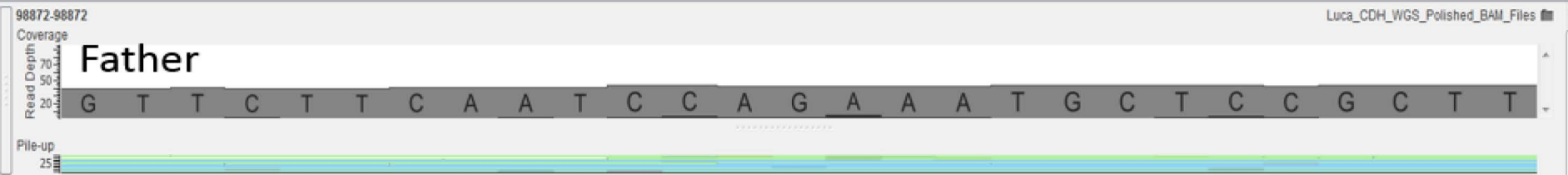
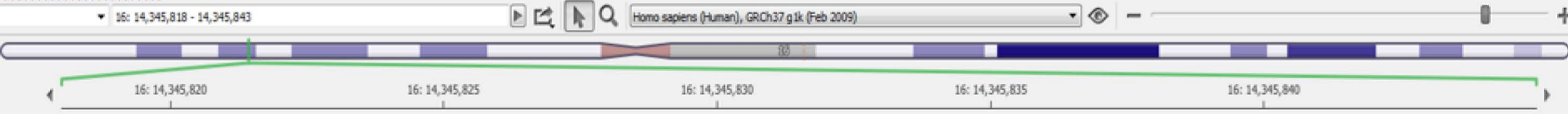
# Proband 95334



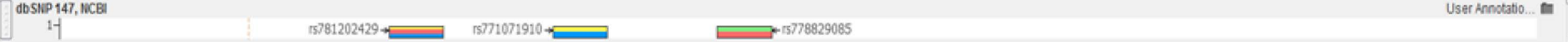
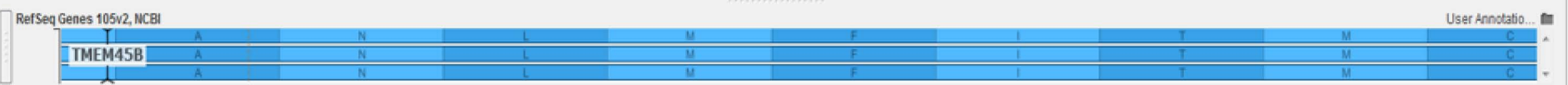
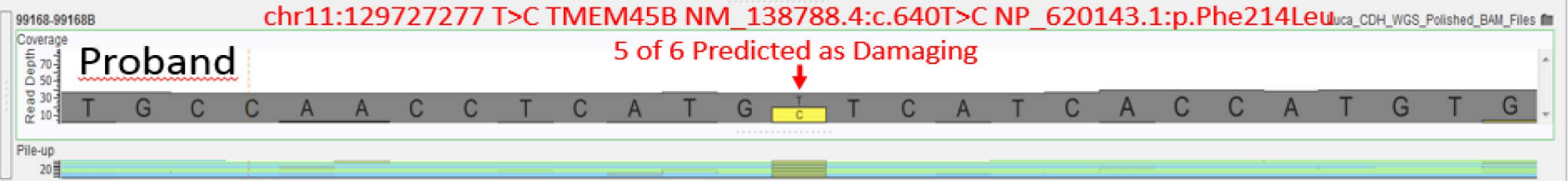
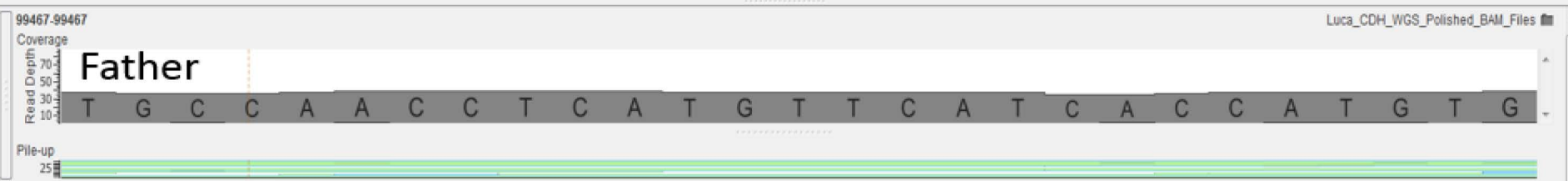
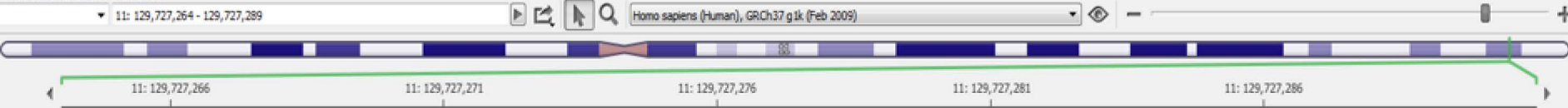
# Proband 95377



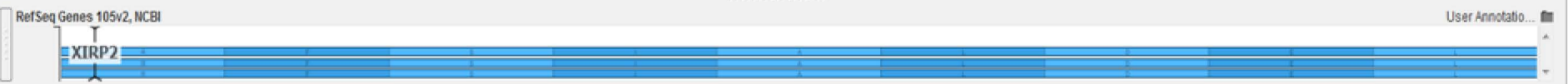
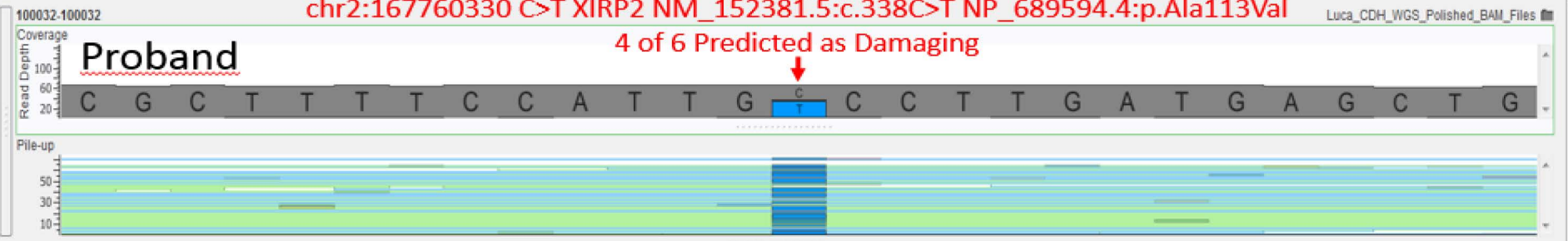
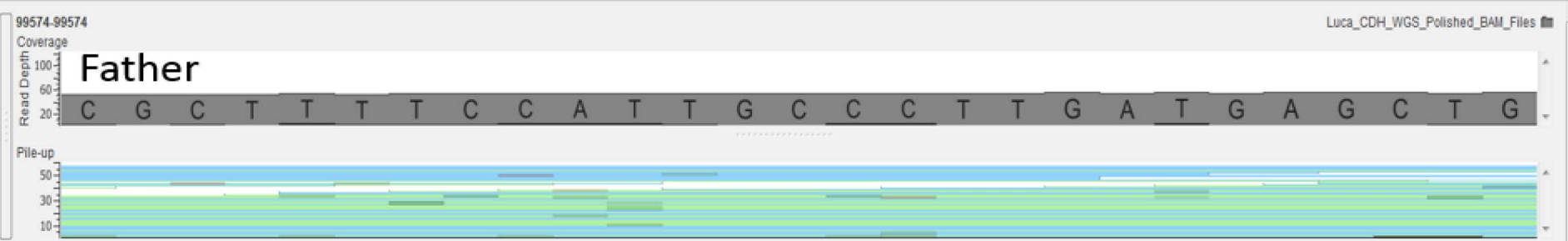
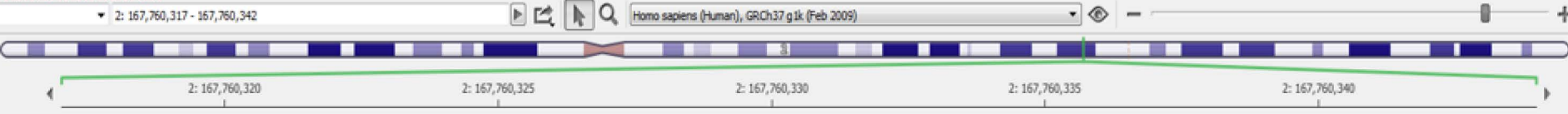
# Proband 98873



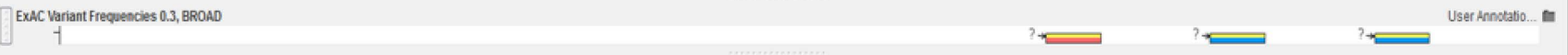
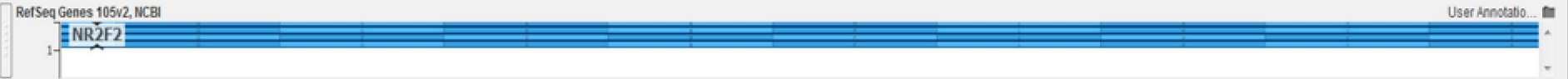
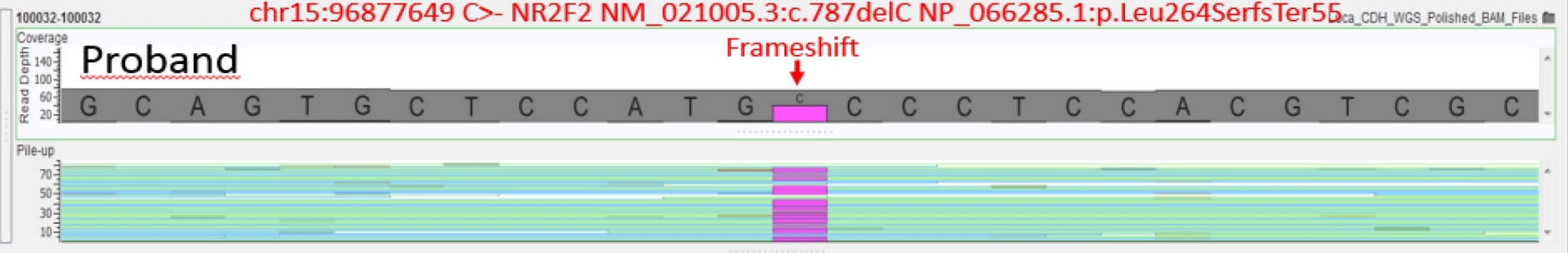
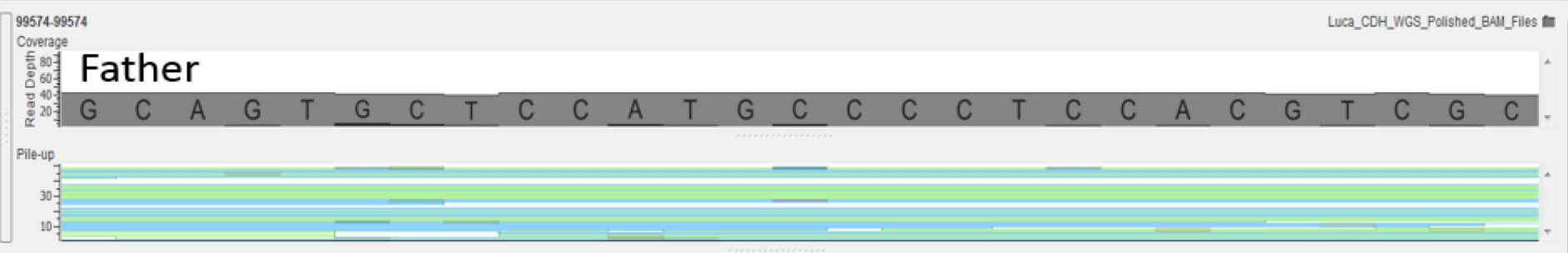
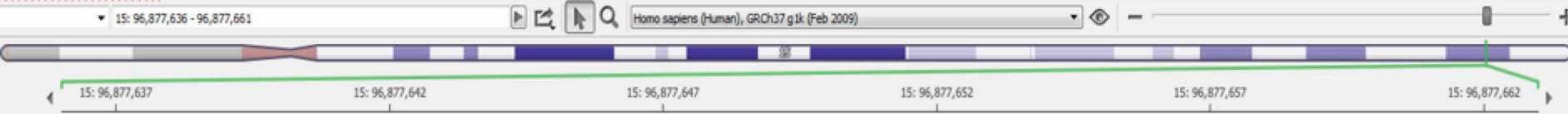
# Proband 99168



# Proband 100032



# Proband 100032



# Proband 101260

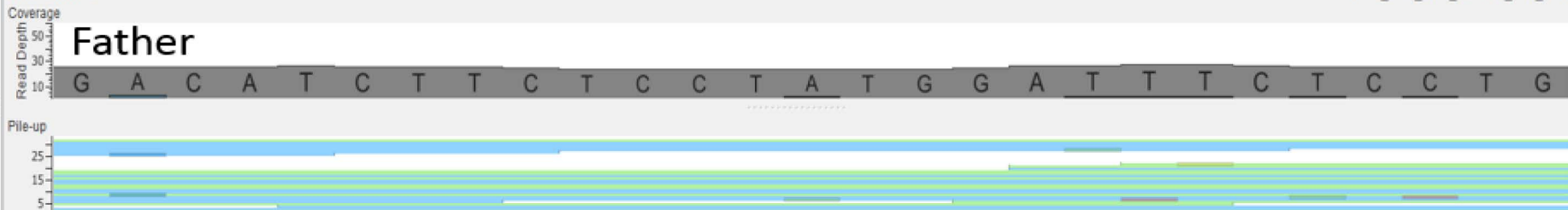
6: 76,072,481 - 76,072,506

Homo sapiens (Human), GRCh37 g1k (Feb 2009)

6: 76,072,486      6: 76,072,491      6: 76,072,496      6: 76,072,501      6: 76,072,506

101189-101189

Luca\_CDH\_WGS\_Polished\_BAM\_Files



101188-101188

Luca\_CDH\_WGS\_Polished\_BAM\_Files

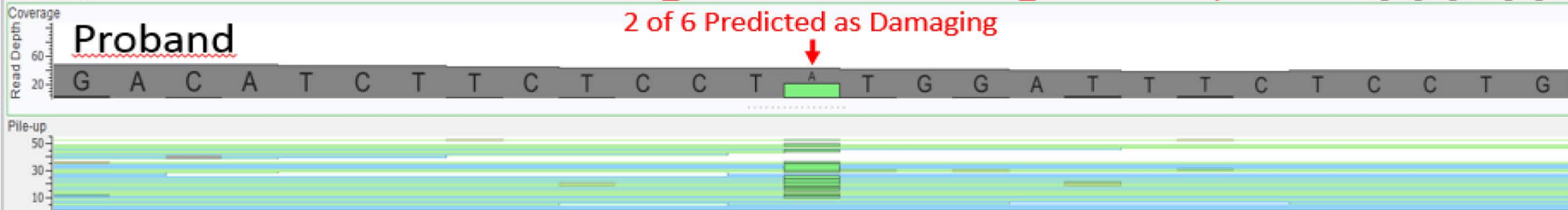


chr6:76072494 A>G FILIP1 NM\_001289987.2:c.425T>C NP\_001276916.1:p.Ile142Thr

2 of 6 Predicted as Damaging

101260-101260

Luca\_CDH\_WGS\_Polished\_BAM\_Files



RefSeq Genes 105v2, NCBI

User Annotatio...



Reference Sequence GRCh37 g1k, 1000Genomes

User Annotatio...



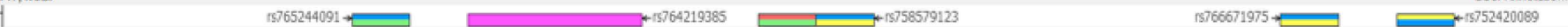
ExAC Variant Frequencies 0.3, BROAD

User Annotatio...

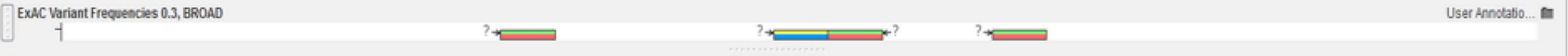
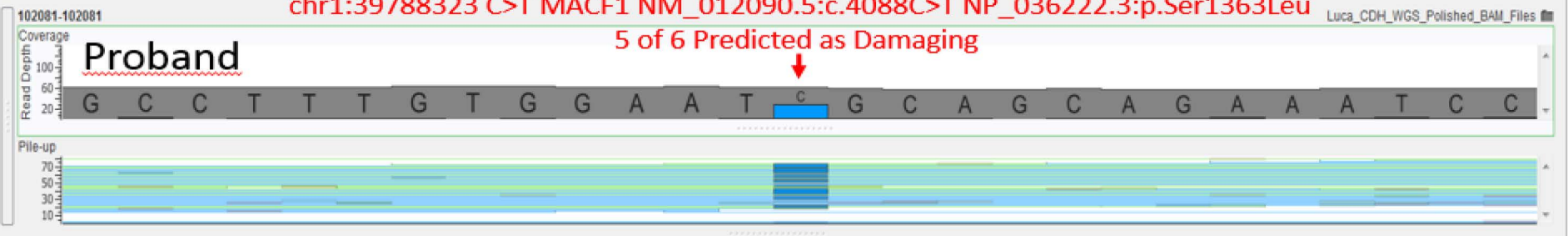
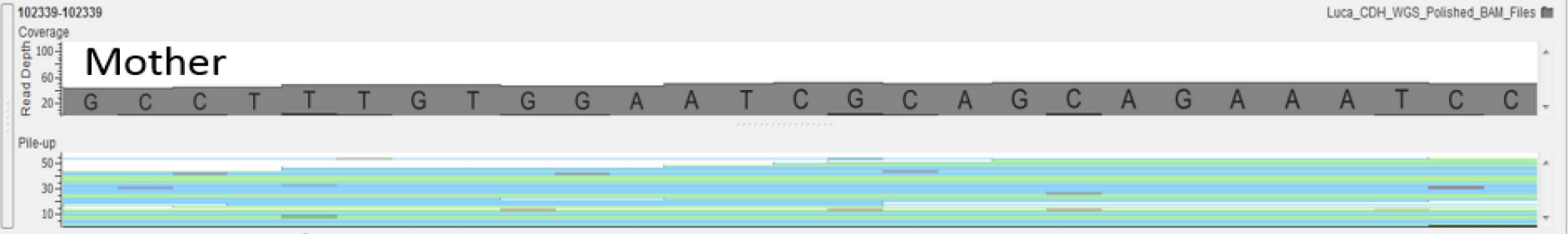
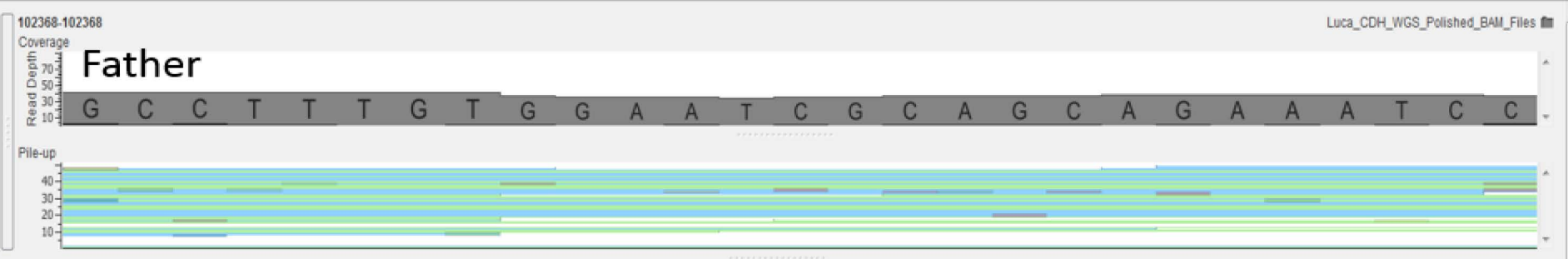
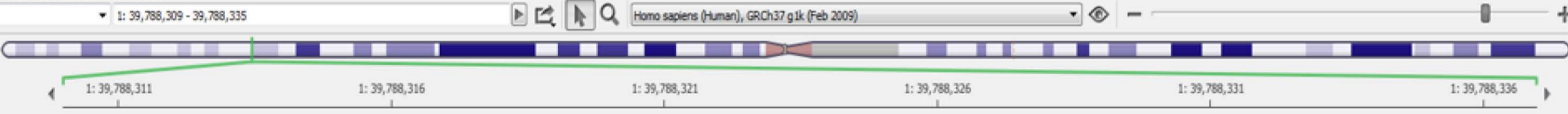


dbSNP 147, NCBI

User Annotatio...



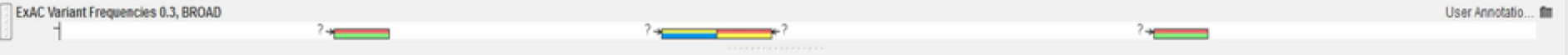
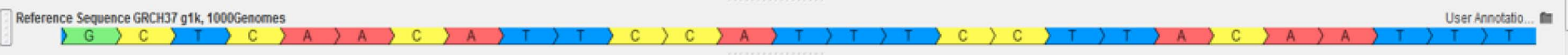
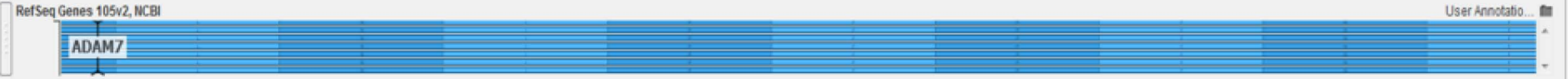
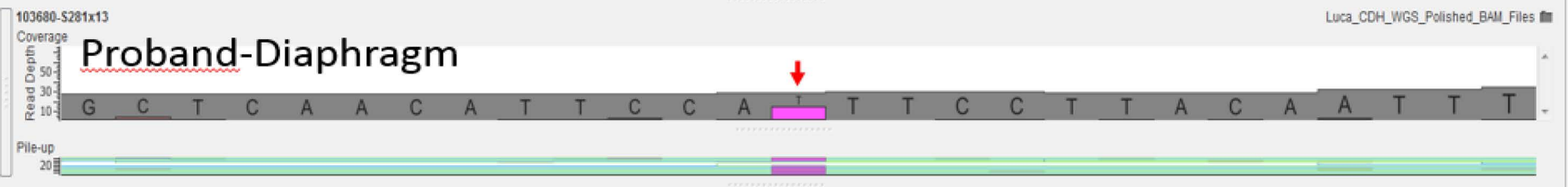
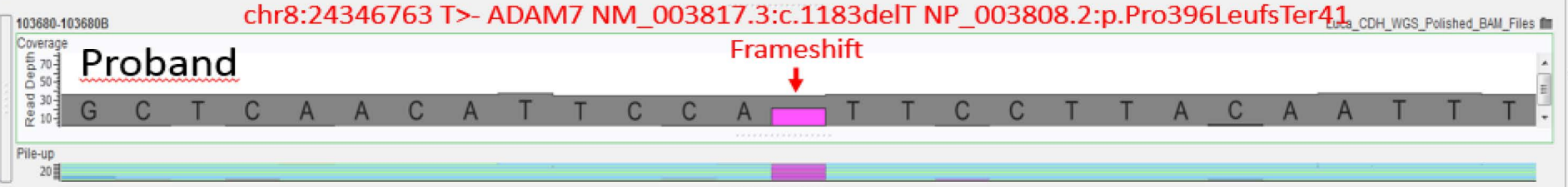
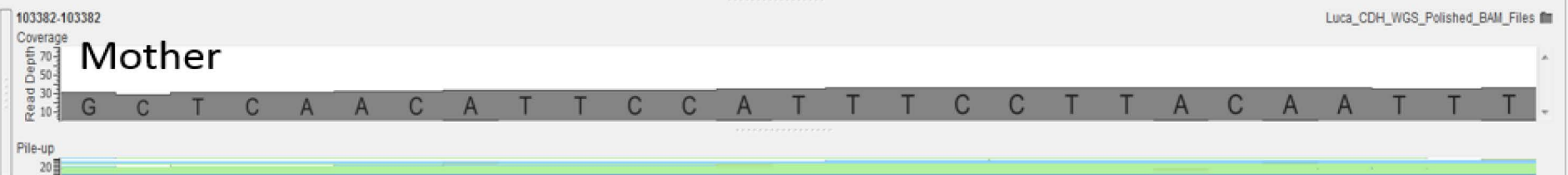
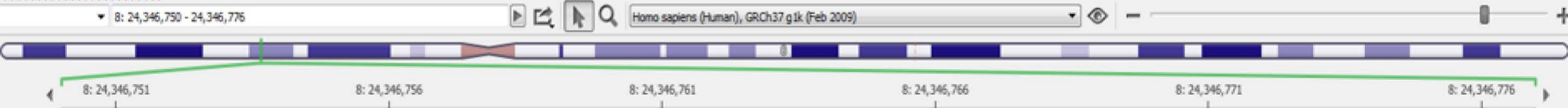
# Proband 102081



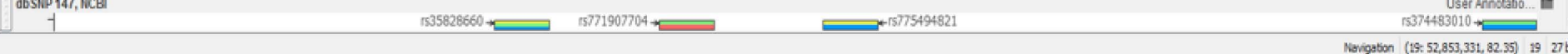
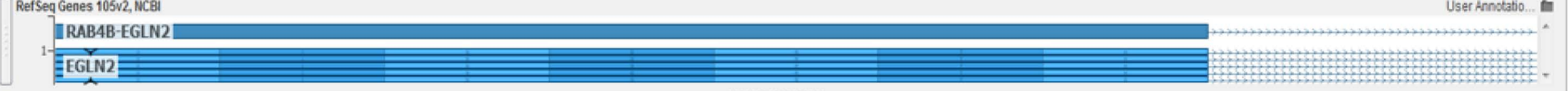
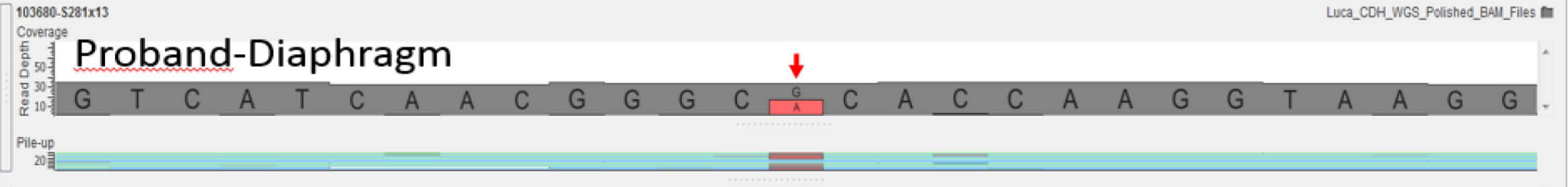
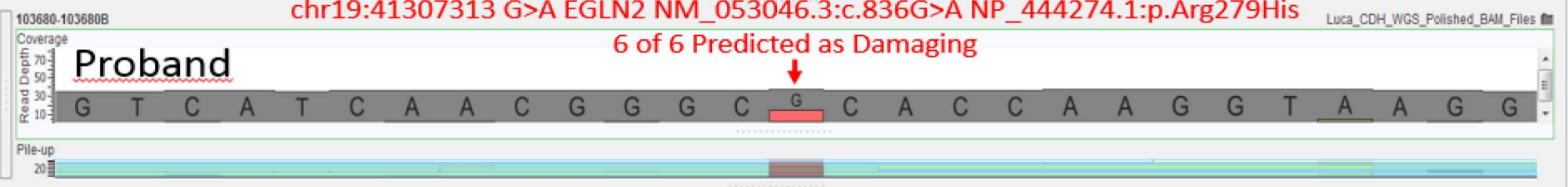
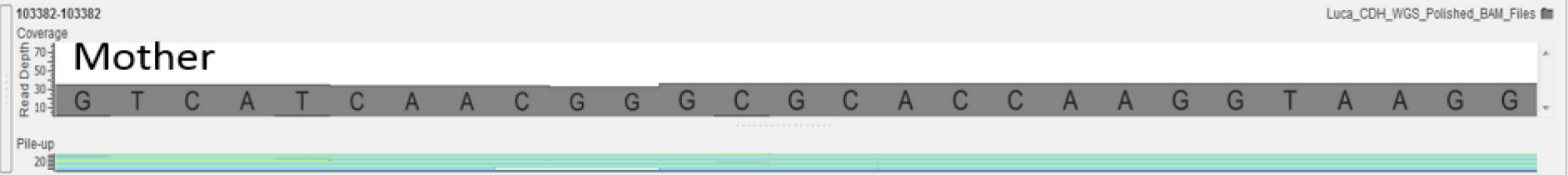
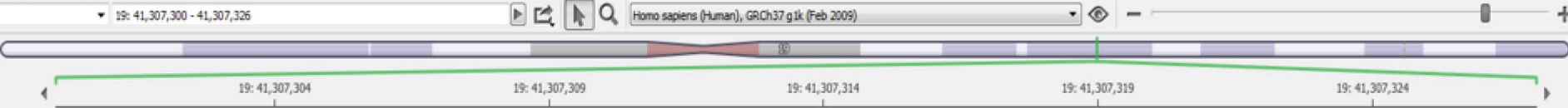




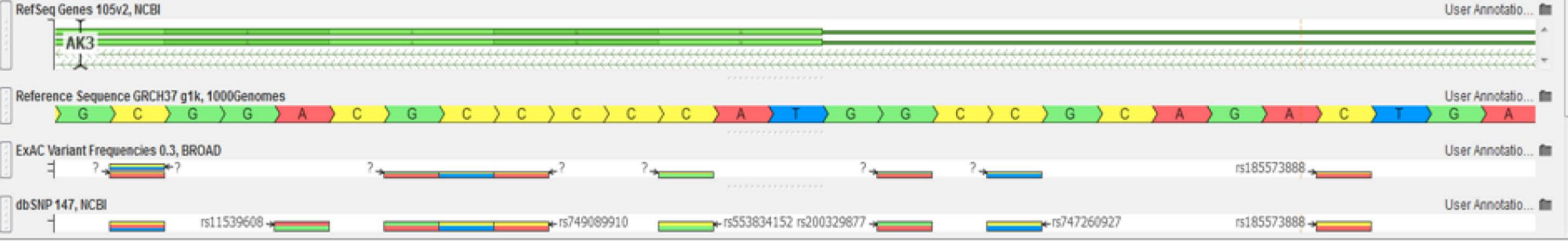
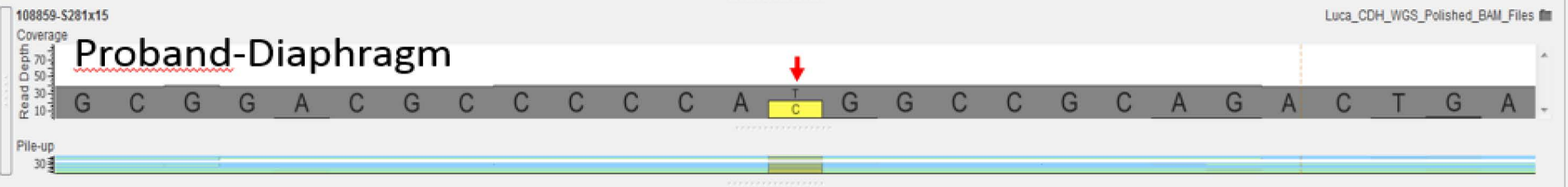
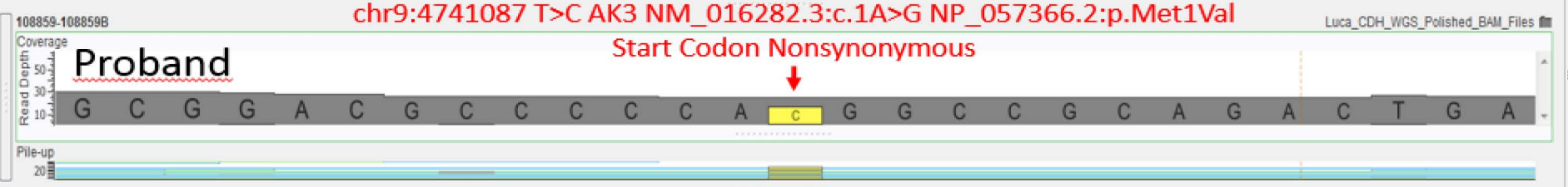
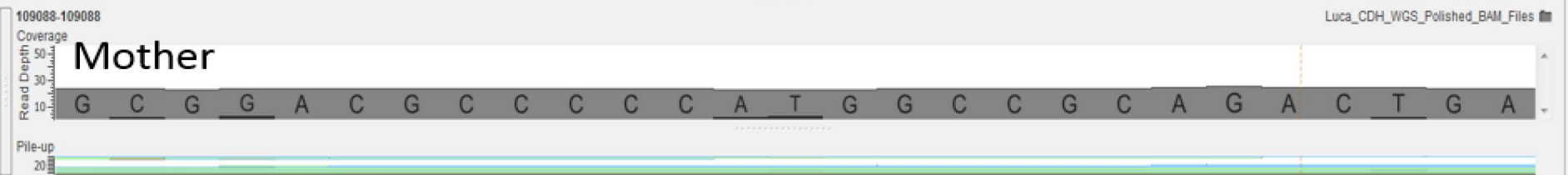
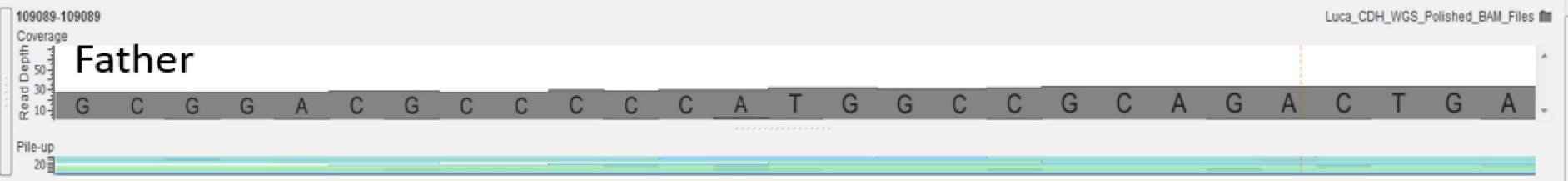
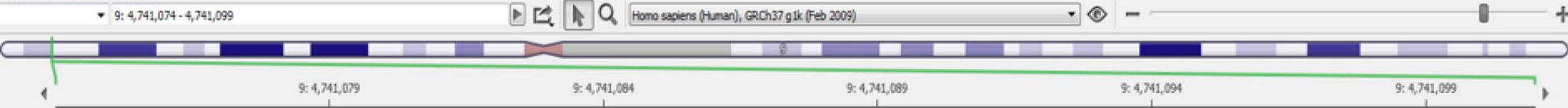
# Proband 103680



# Proband 103680



# Proband 108859



# Proband 108859

