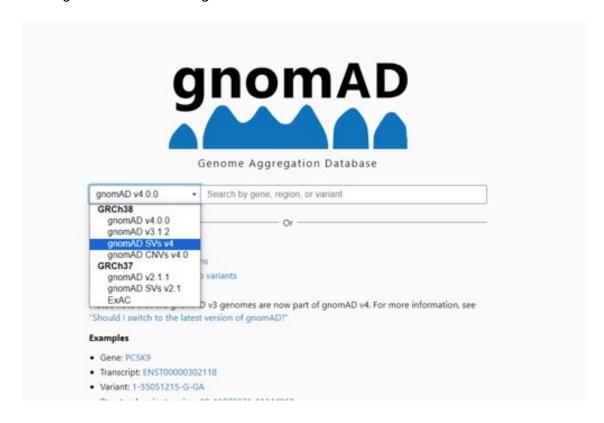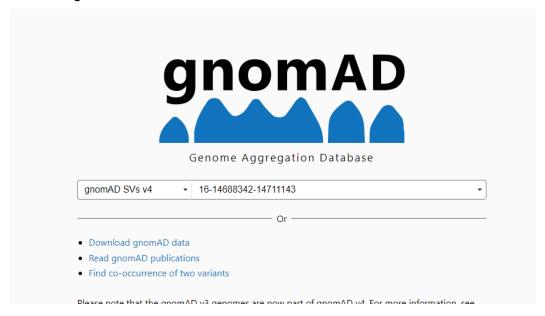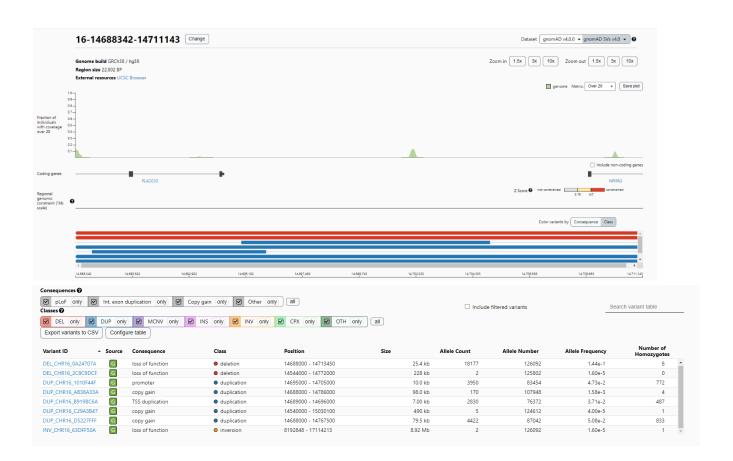1. Go to https://gnomad.broadinstitute.org/

2. Click "gnomAD v4.0.0"

3. Click "gnomAD SVs v4" or "gnomAD SVs v2.1"

4. Enter the genomic coordinates

5. Structural variants (SVs) within this region are displayed including gains, losses and rearrangements of at least 50 nucleotides.

Structural variants in gnomAD can be displayed in UCSC browser. For more information, go to ==XXX (link to Casey's document).==

Author Information and Affiliations:

Aiko Otsubo, PhD, FACMG

University of Michigan

Diagnostic Genetics and Genomics

Created: 3/13/24