

The figure displays a genomic map of the mitochondrial genome of *Glomera glaberrima*, showing the distribution and expression of genes across six different strains. The x-axis represents the Genome Position (0 to 80,000). The y-axis represents the expression level (RPM1).

The strains are listed on the right side of the map:

- GLBRCY22
- CBS432
- IFO1815
- ZP591
- CBS10644
- CBS7001

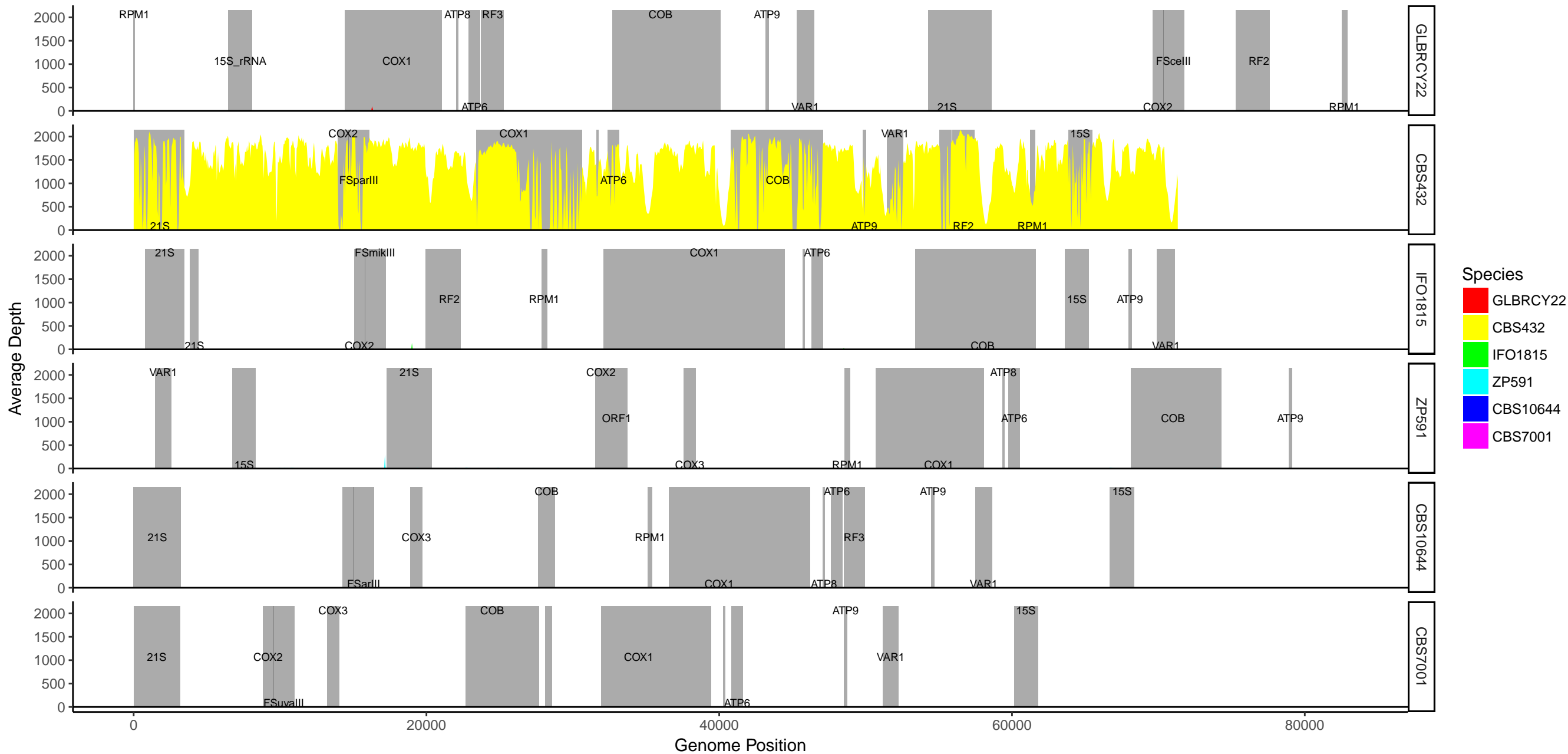
The genes shown in the map include:

- ATP6
- ATP8
- ATP9
- COX1
- COX2
- COX3
- COB
- FSceIII
- FSparIII
- FSuvalIII
- FSmikIII
- RF2
- RF3
- RPM1
- VAR1
- 21S
- 15S
- 15S_rRNA
- ORF1

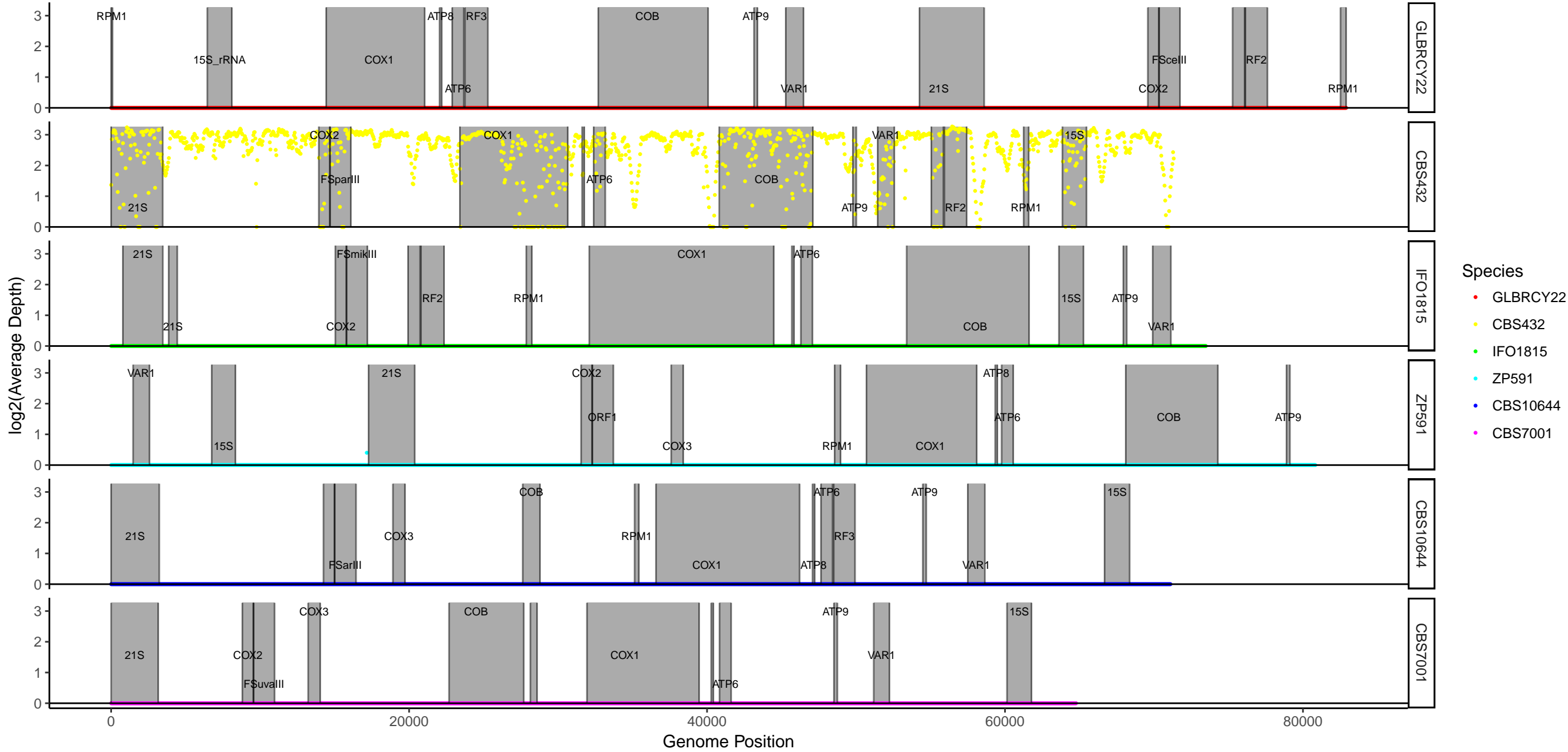
The expression levels are indicated by the height of the bars and the color of the dots. The colors used are red, yellow, green, cyan, blue, and purple. The map shows that the expression of genes varies significantly between strains, with some genes being highly expressed in certain strains and others being less expressed or absent.

- GLBRCY22
- CBS432
- IFO1815
- ZP591
- CBS10644
- CBS7001

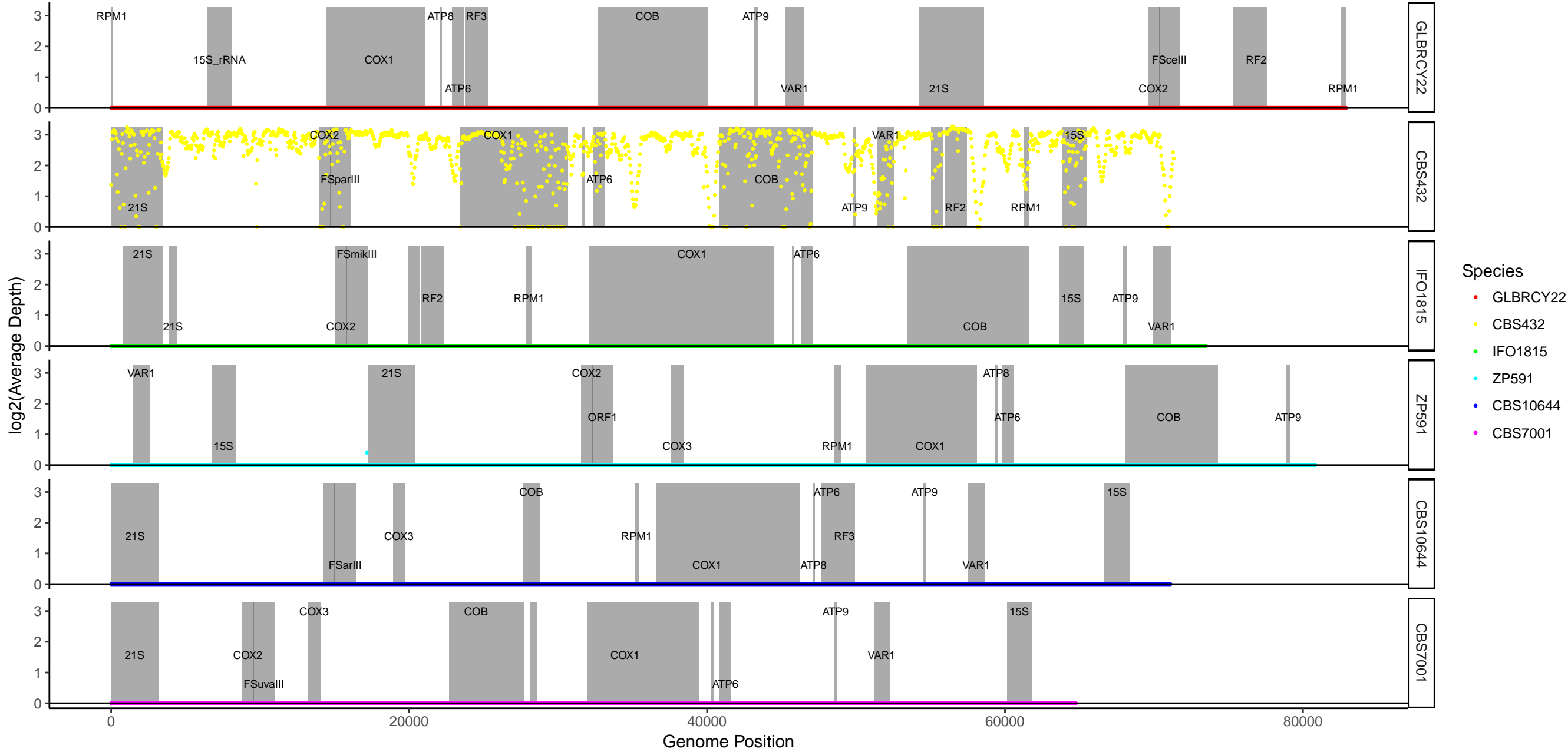
yHRWh18 Avg depth of coverage



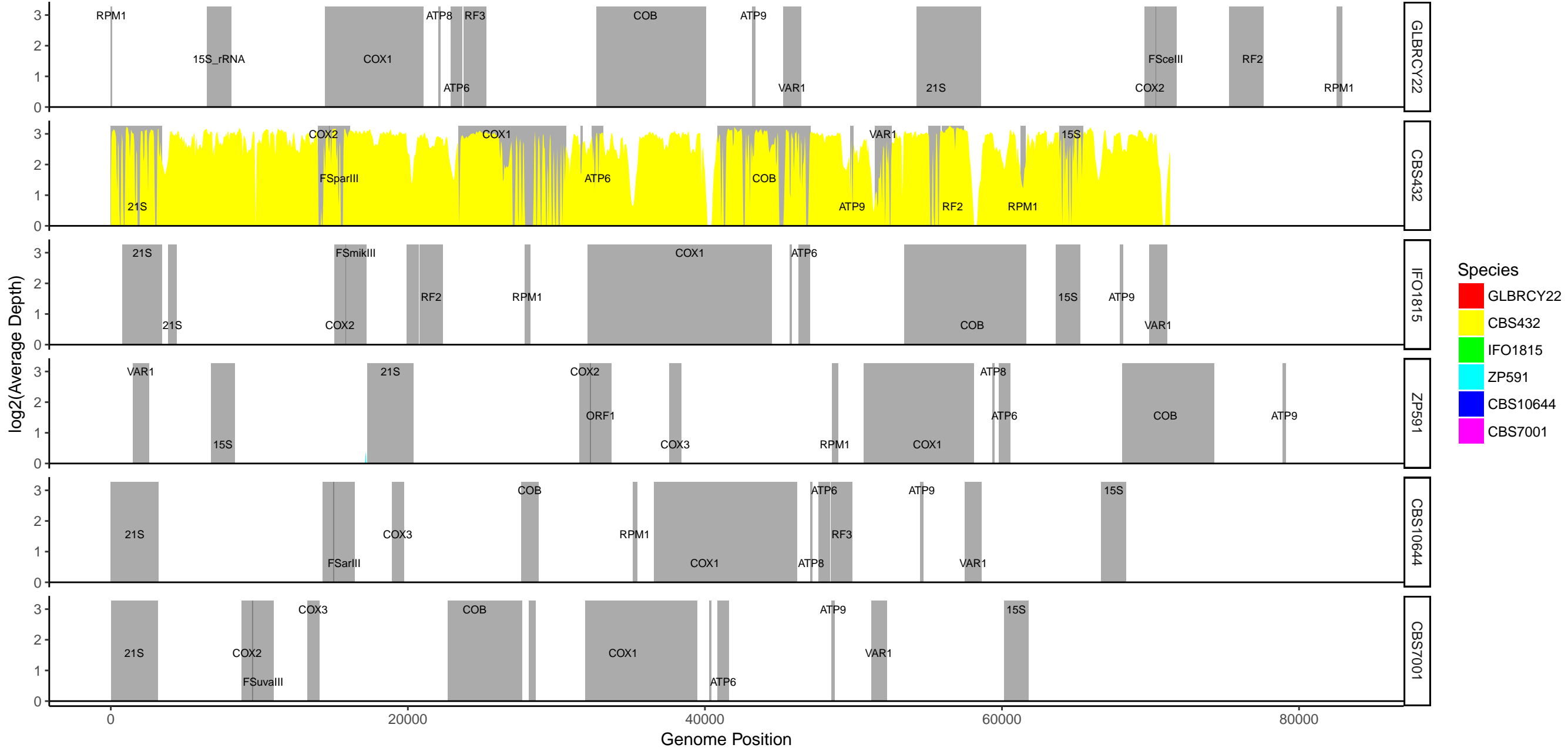
yHRWh18 Log2 Avg depth of coverage



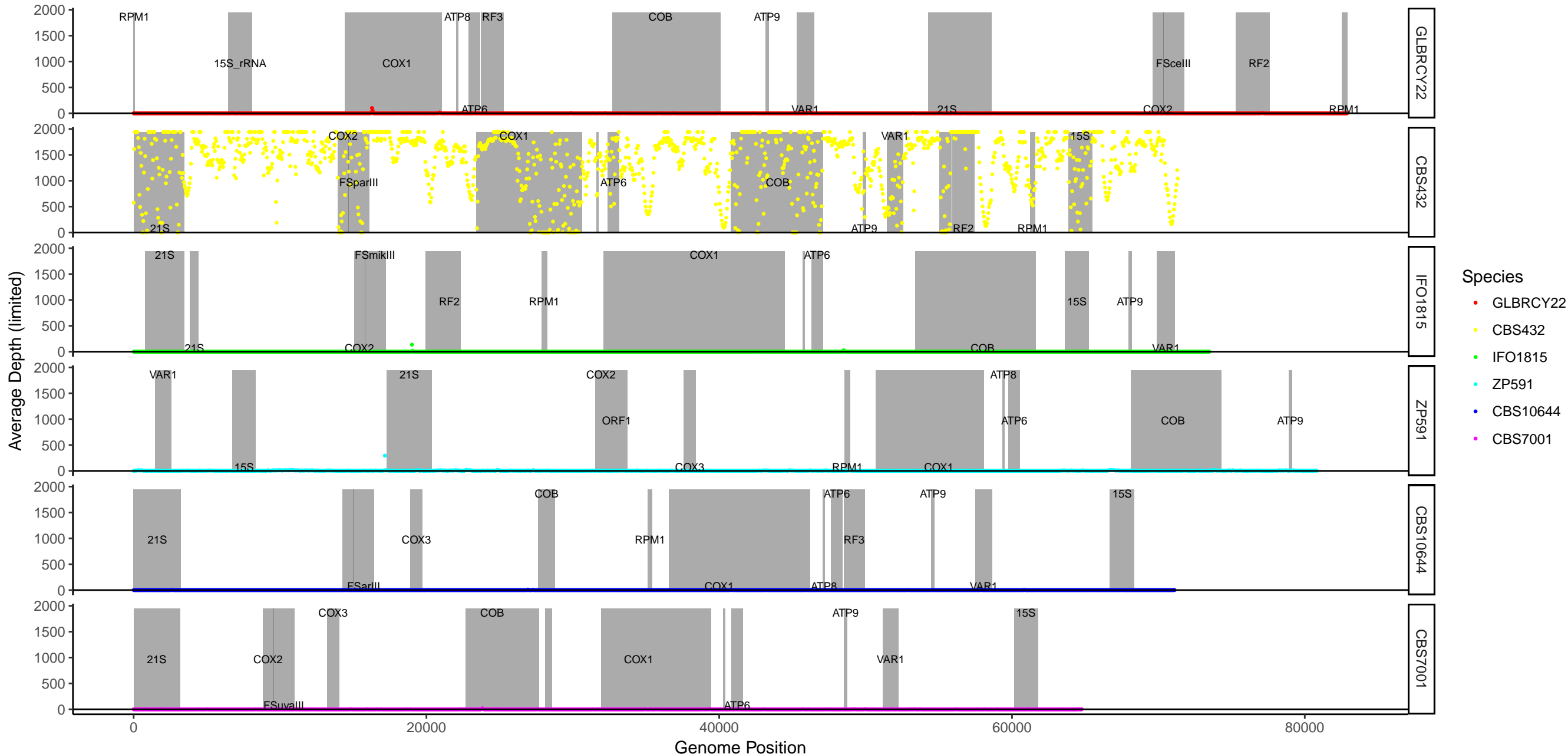
yHRWh18 Log2 Avg depth of coverage



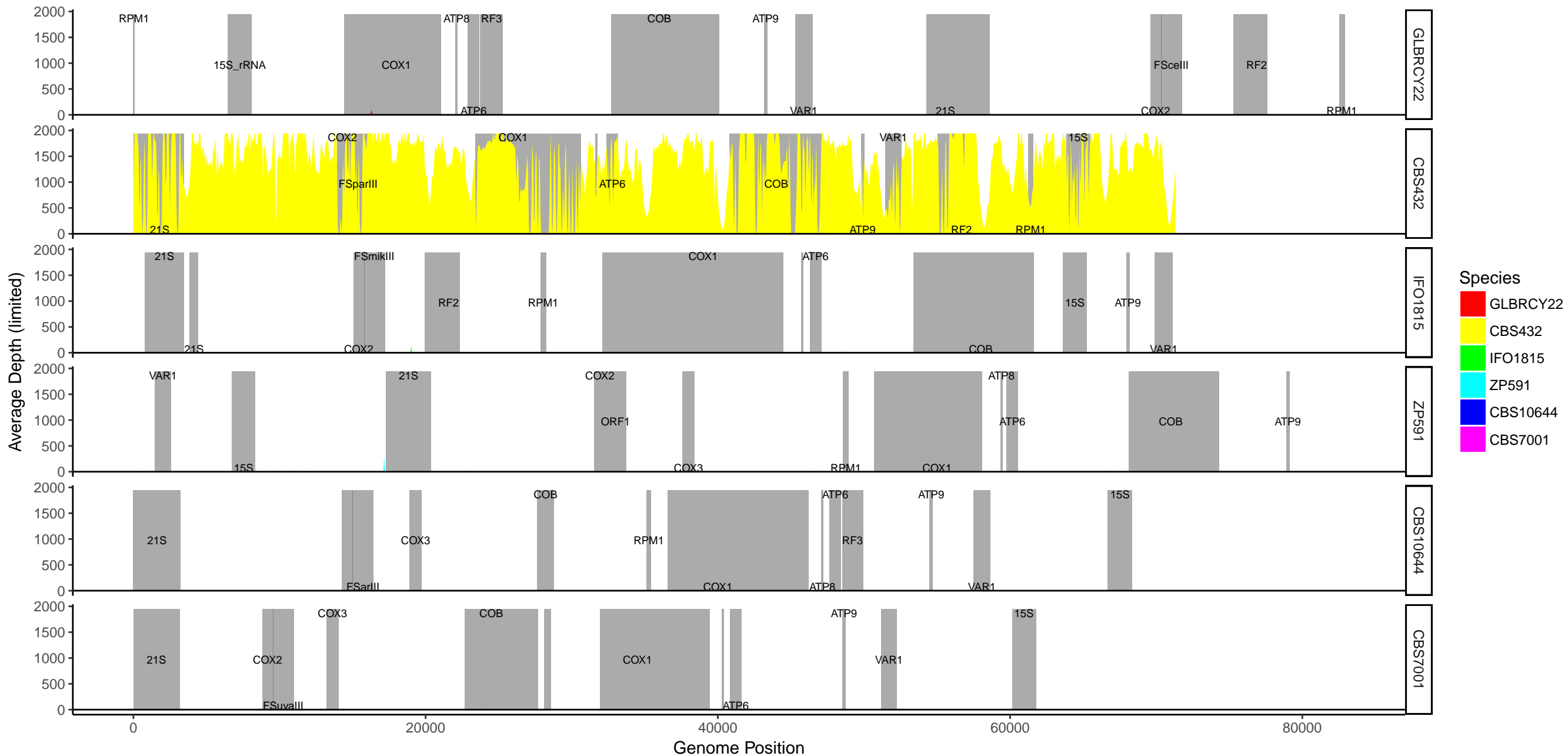
yHRWh18 Avg depth of coverage



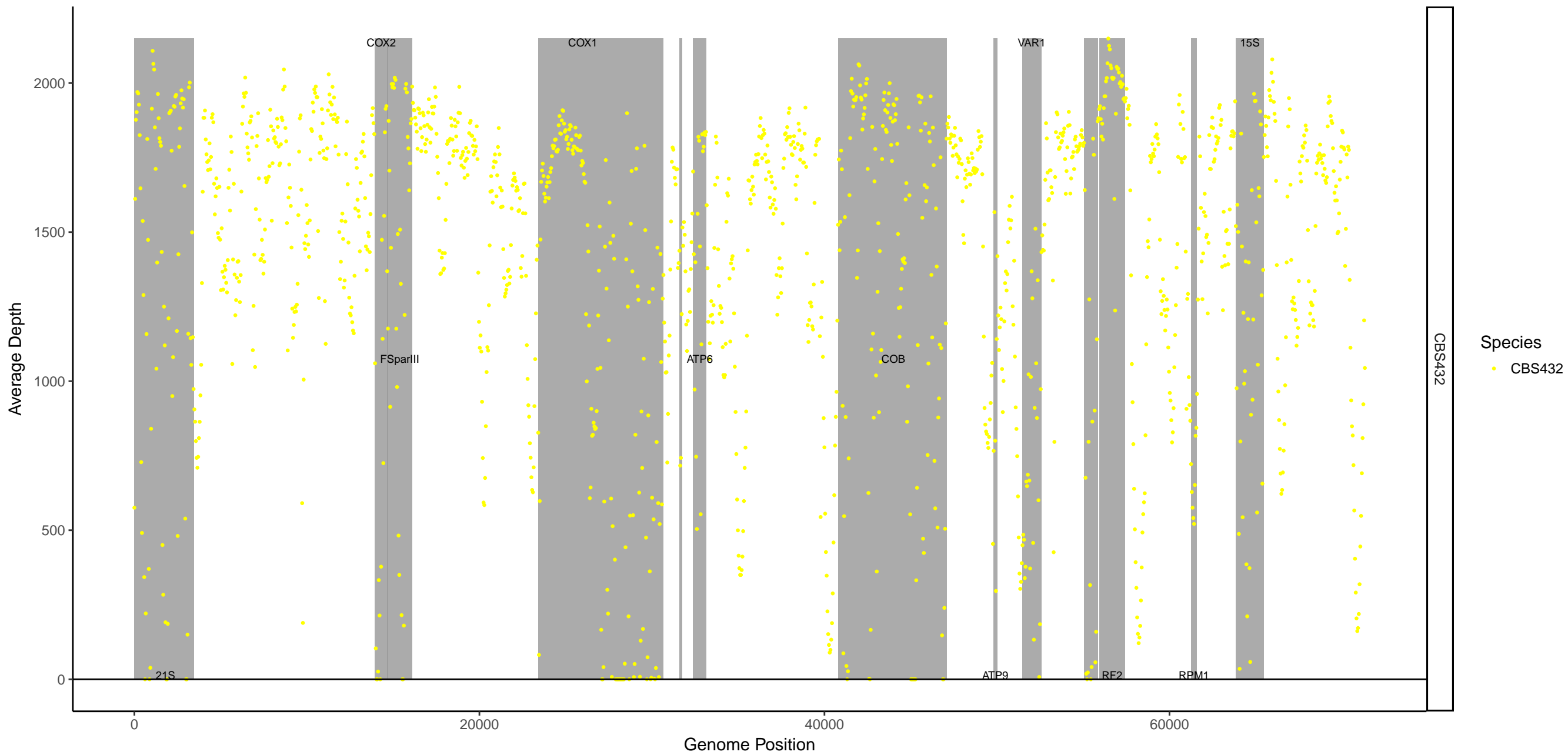
yHRWh18 Avg depth of coverage

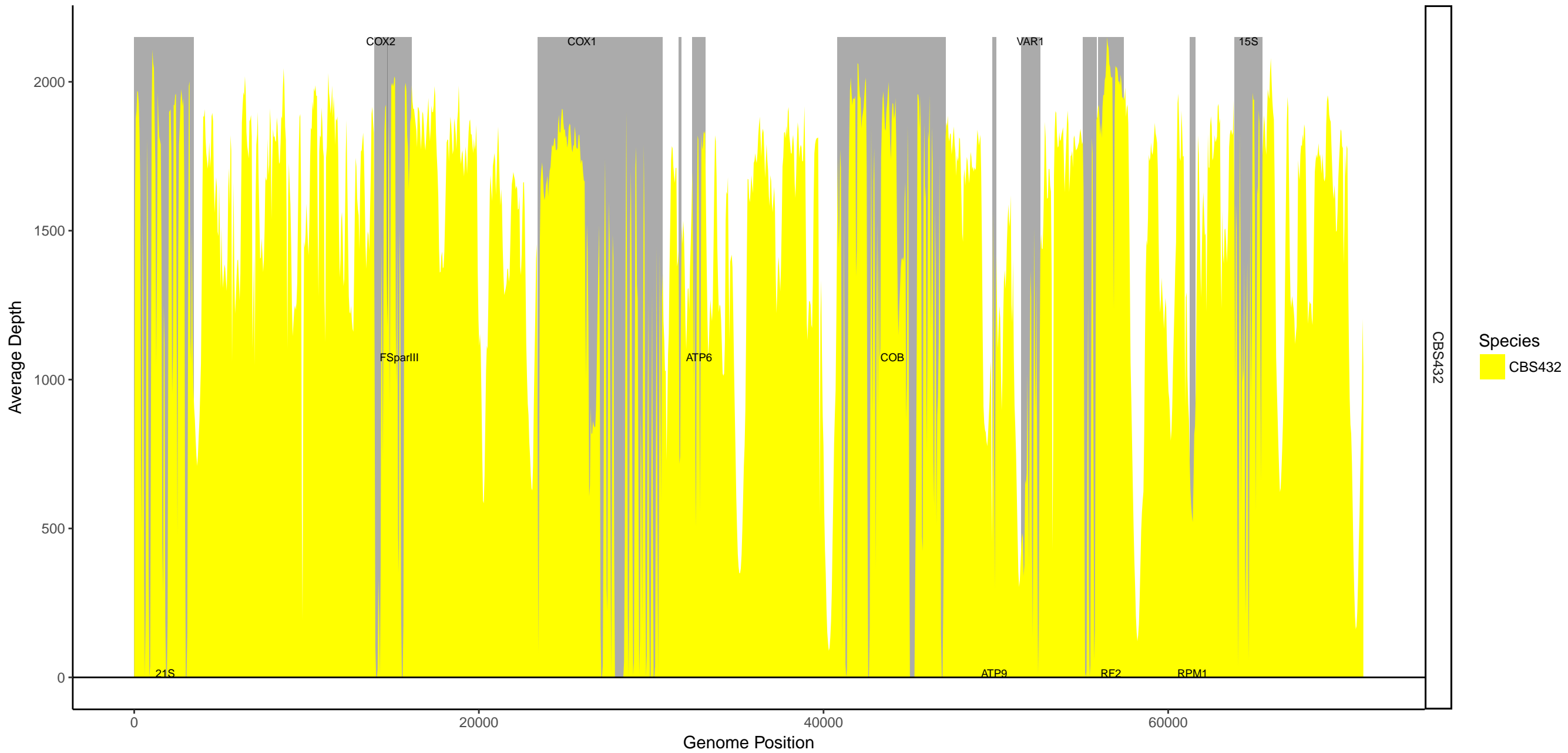


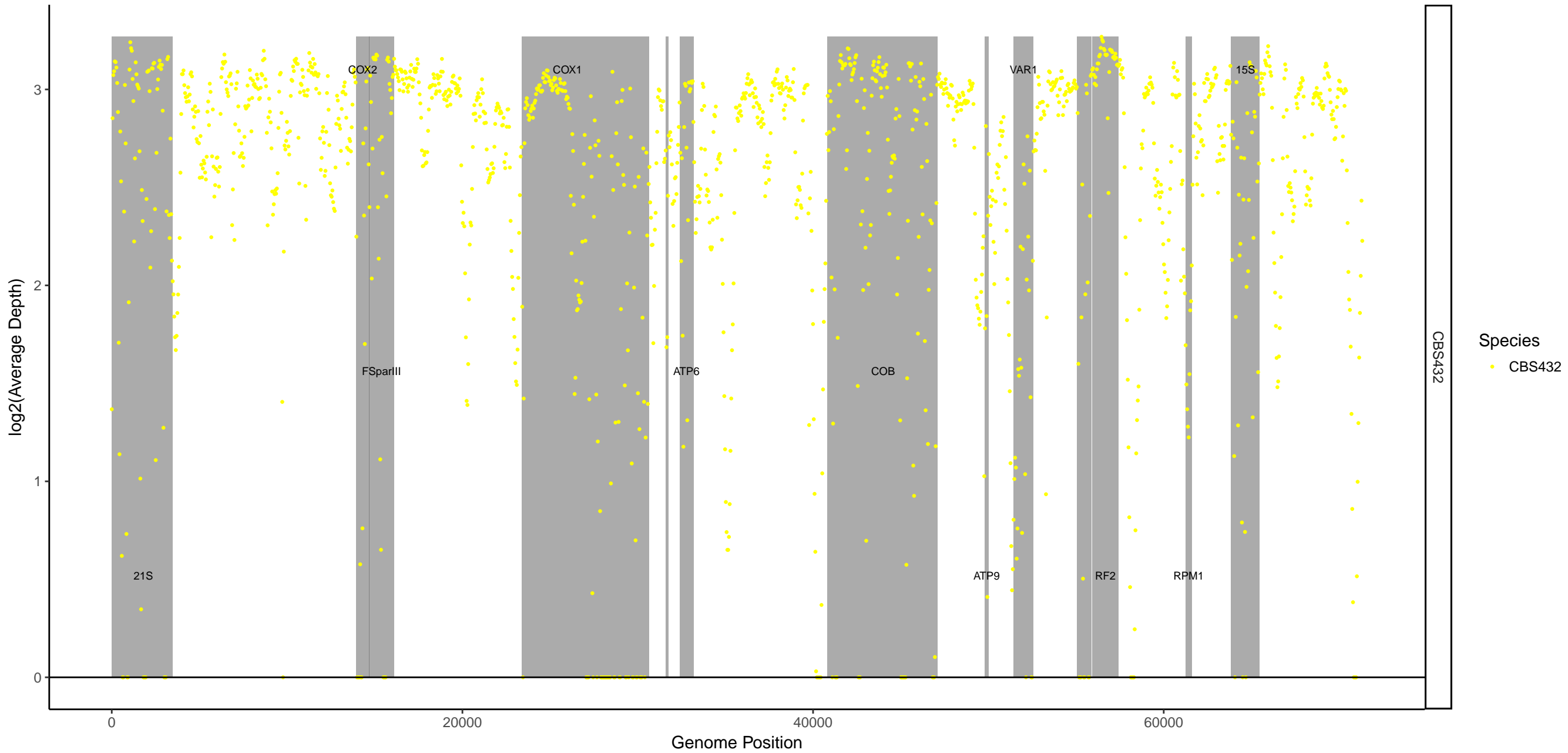
yHRWh18 Avg depth of coverage

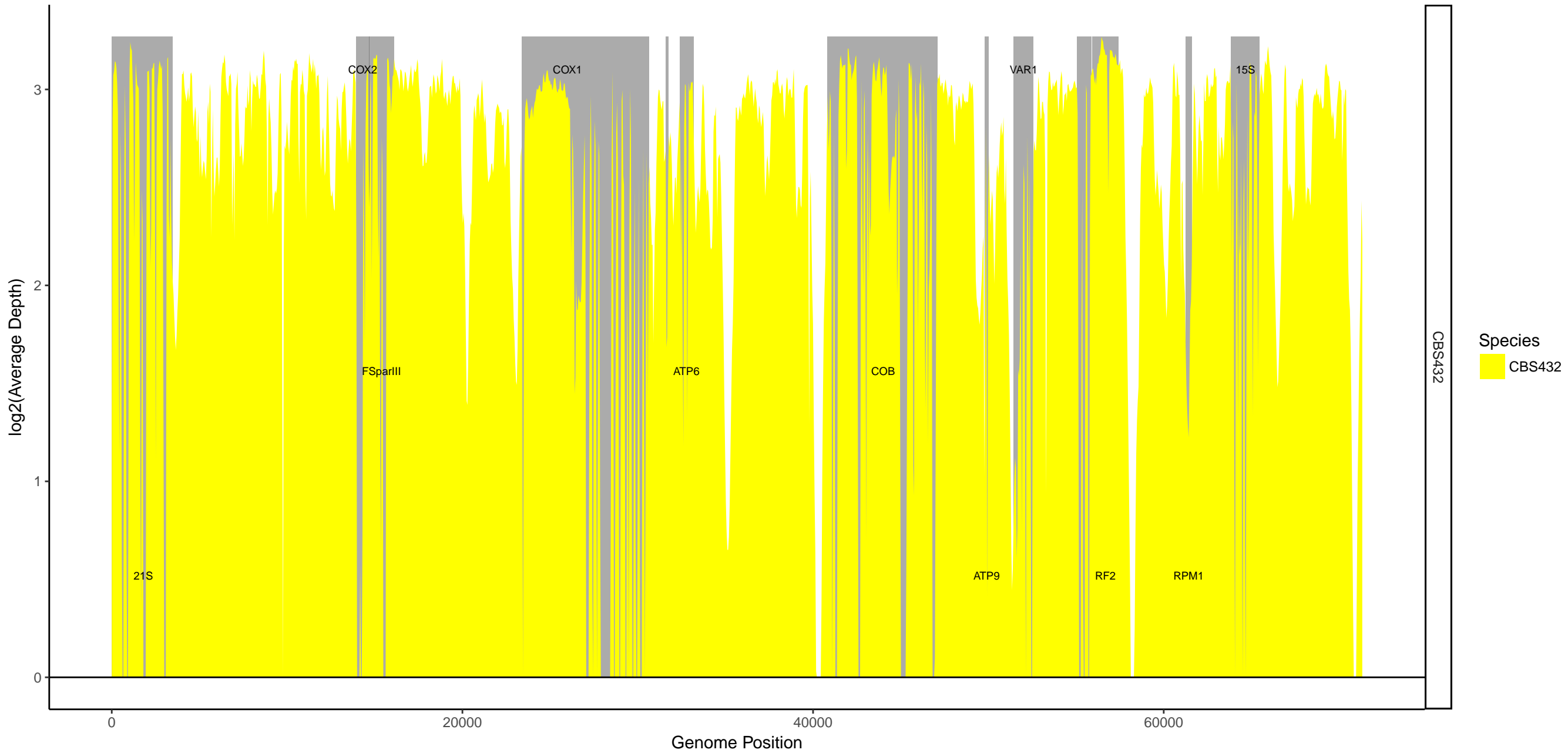


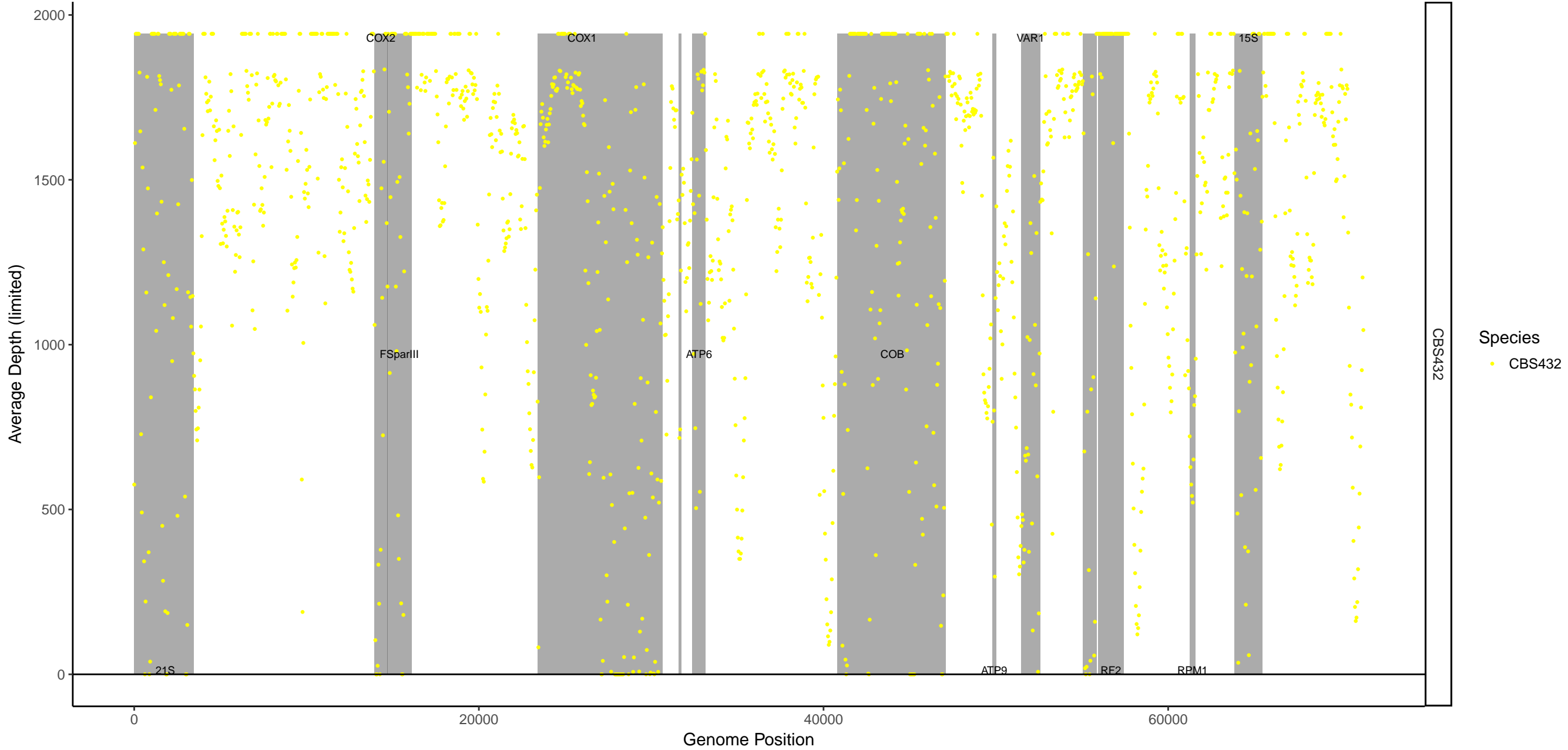
yHRWh18 Avg depth of coverage











yHRWh18 Avg depth of coverage

