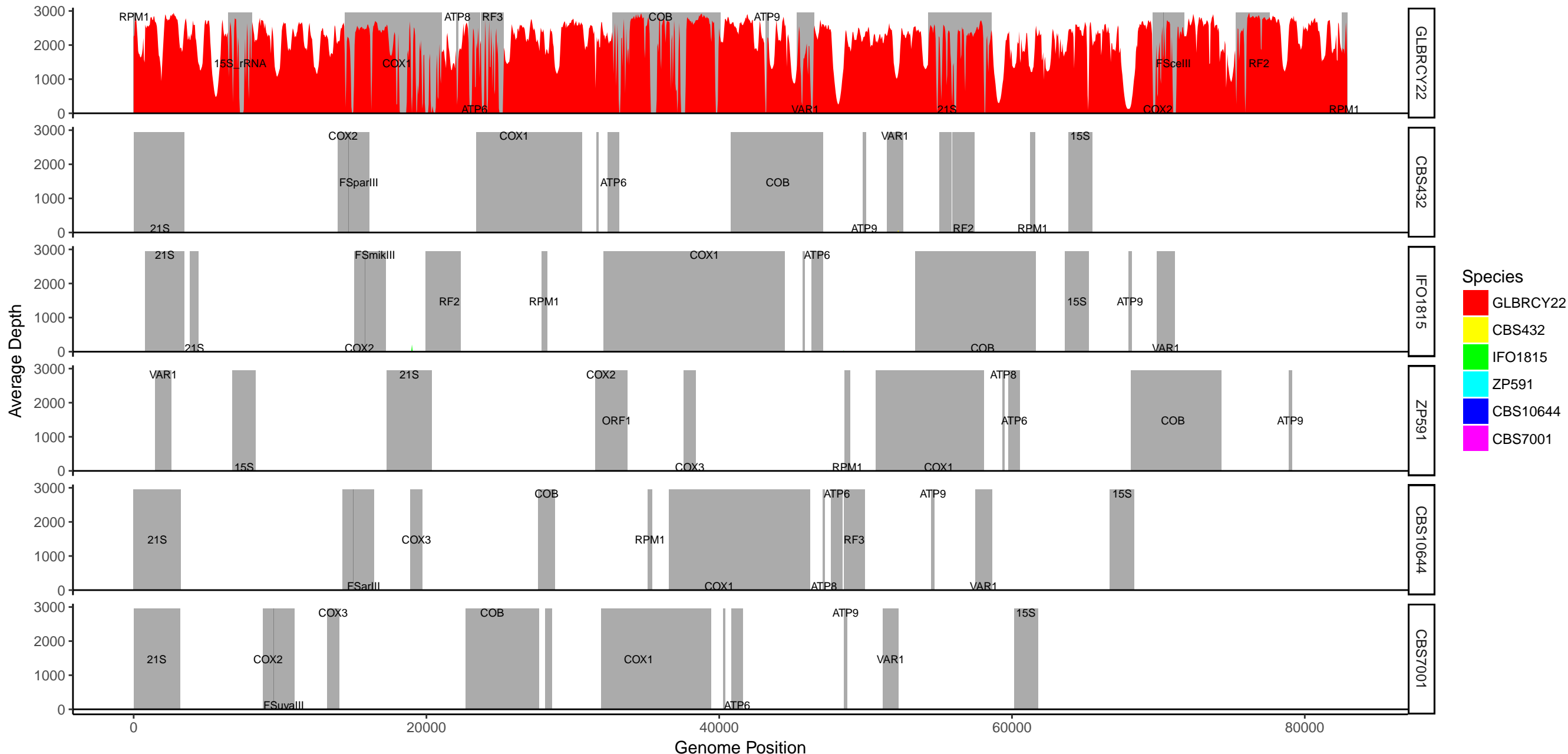
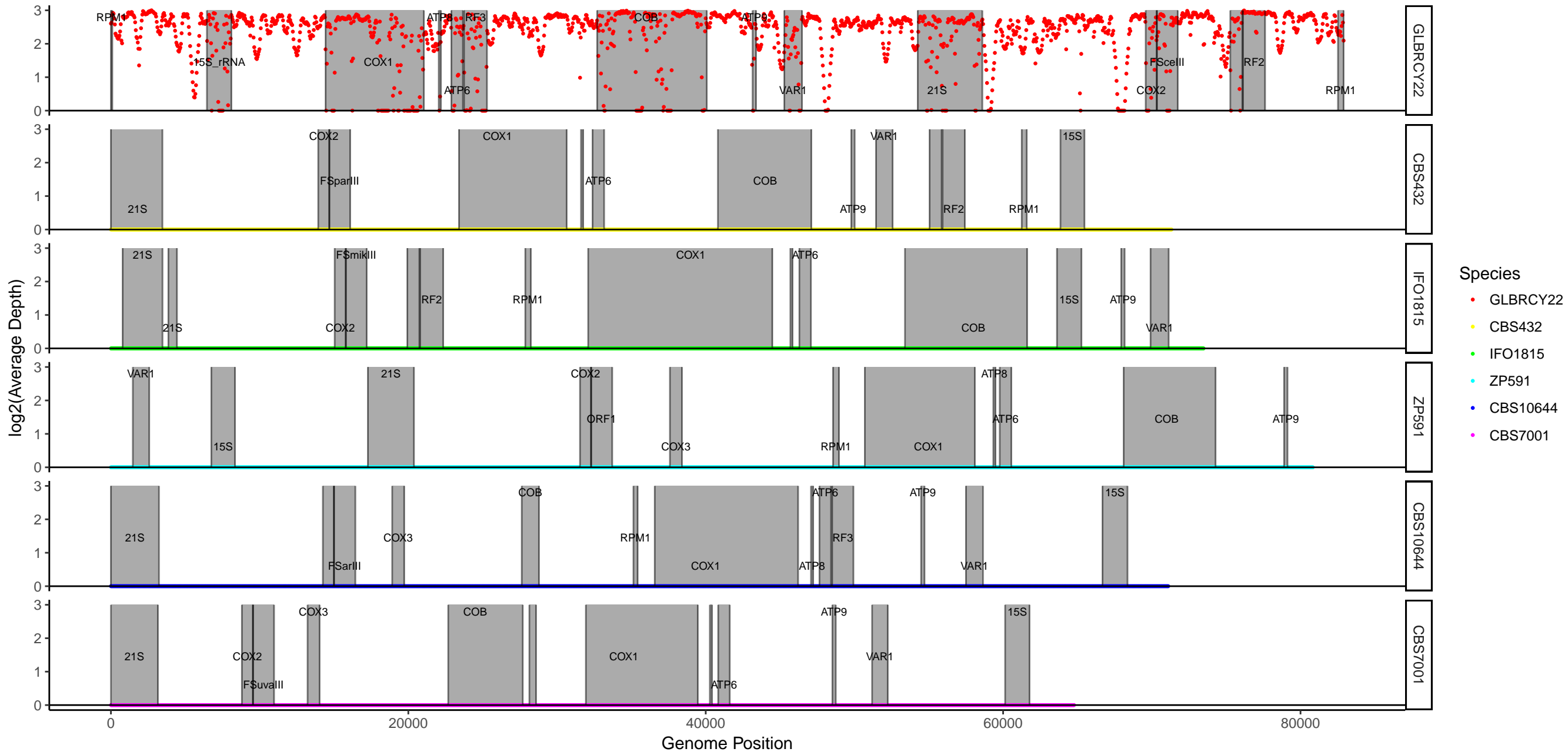


- GLBRCY22
- CBS432
- IFO1815
- ZP591
- CBS10644
- CBS7001

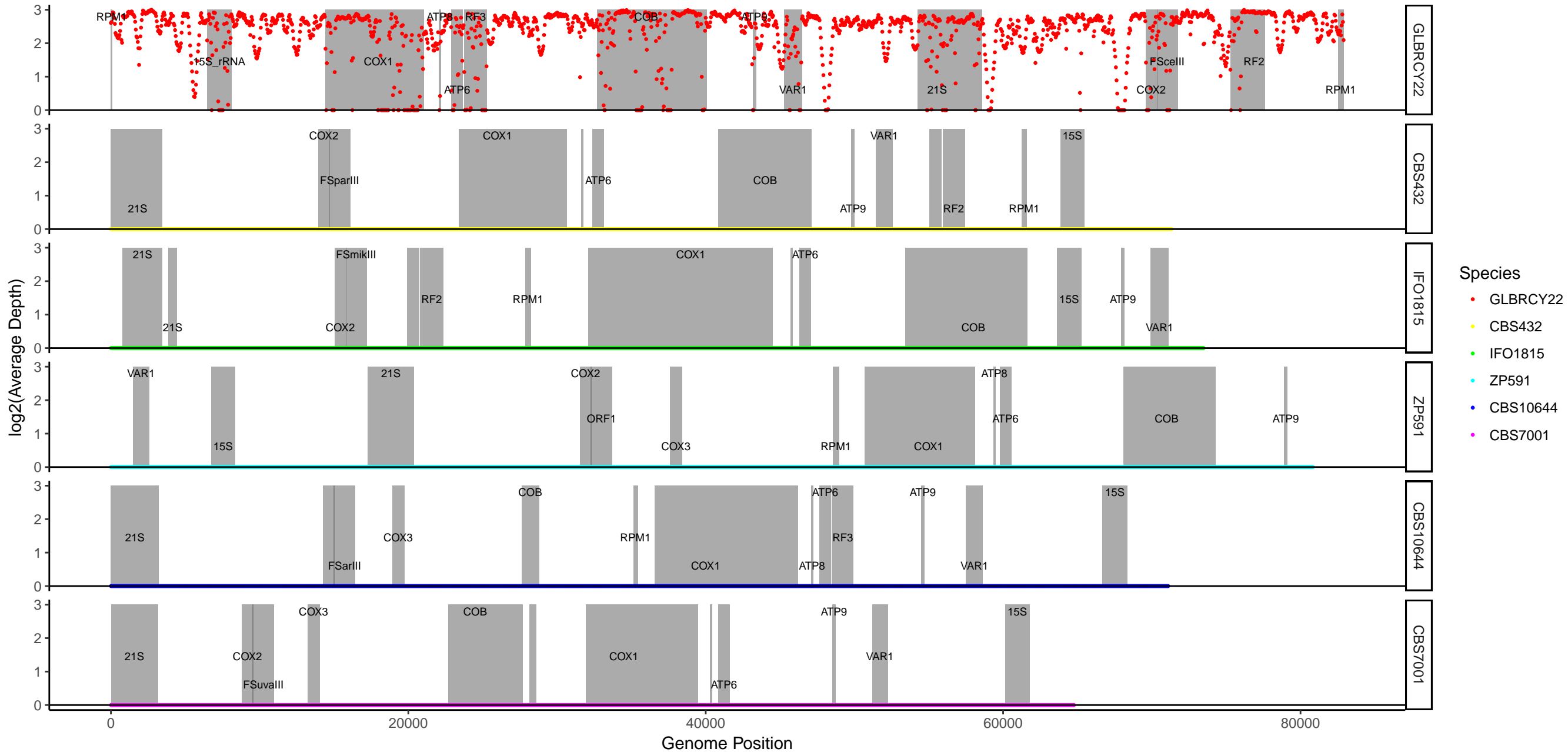
yHRWh87 Avg depth of coverage



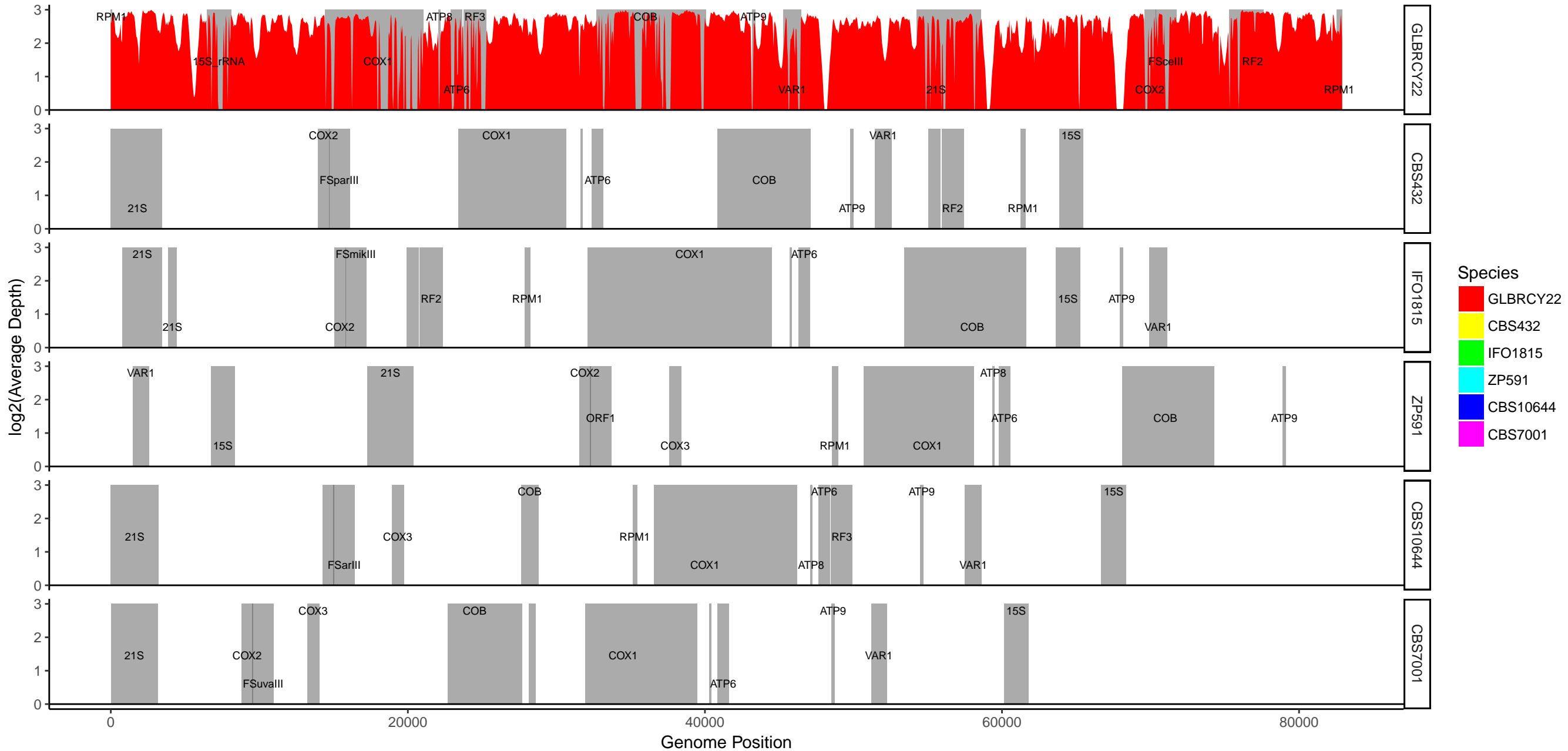
yHRWh87 Log2 Avg depth of coverage



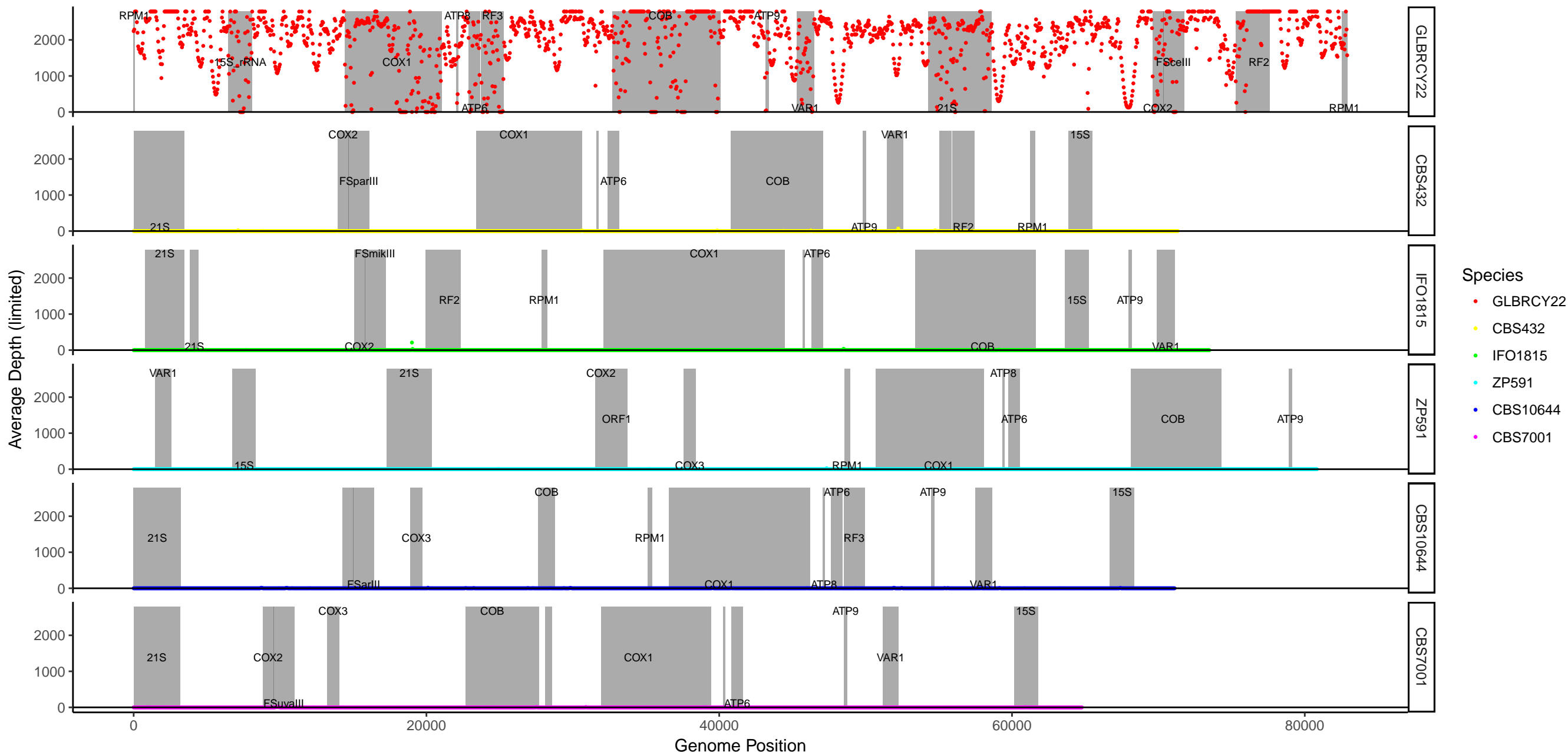
yHRWh87 Log2 Avg depth of coverage



yHRWh87 Avg depth of coverage

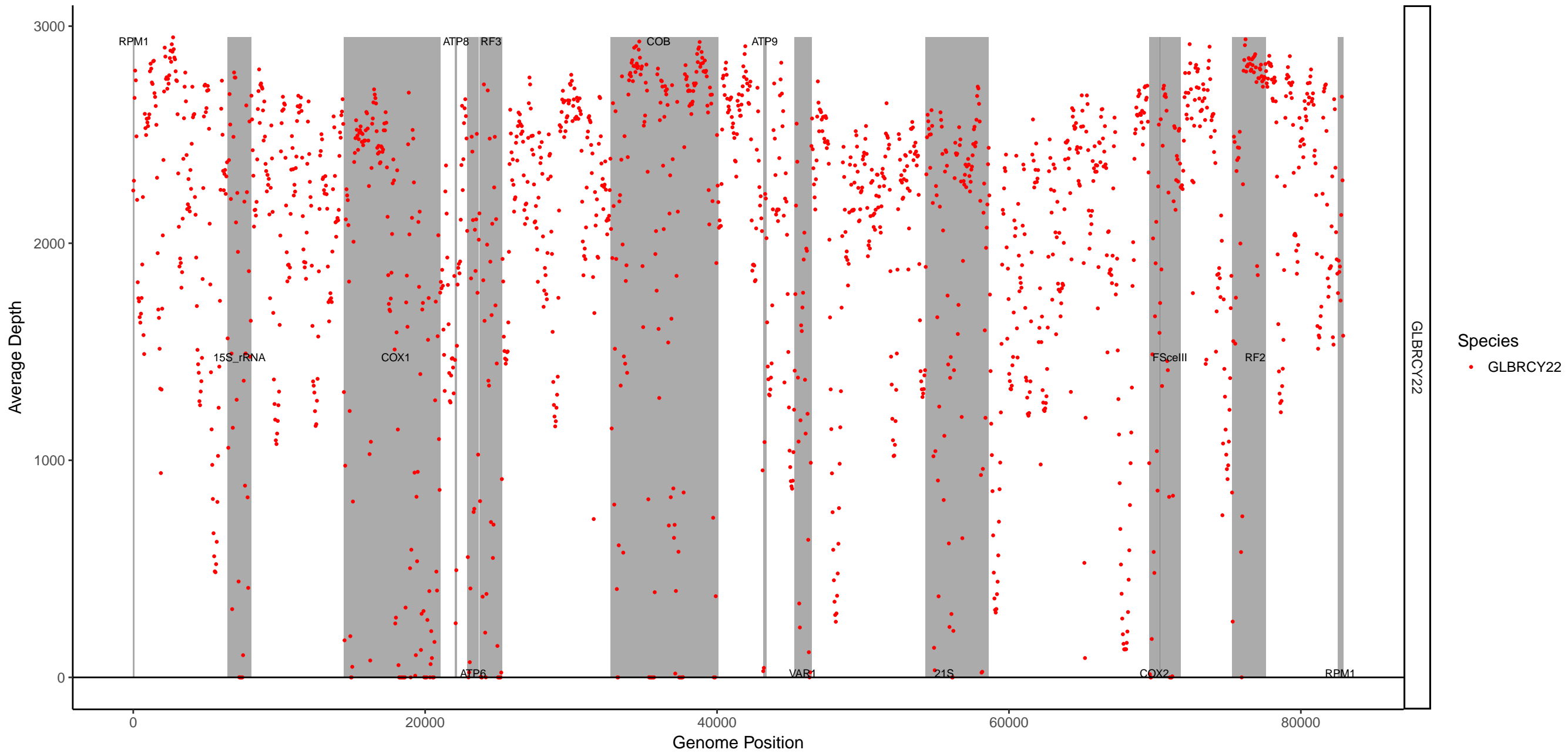


yHRWh87 Avg depth of coverage

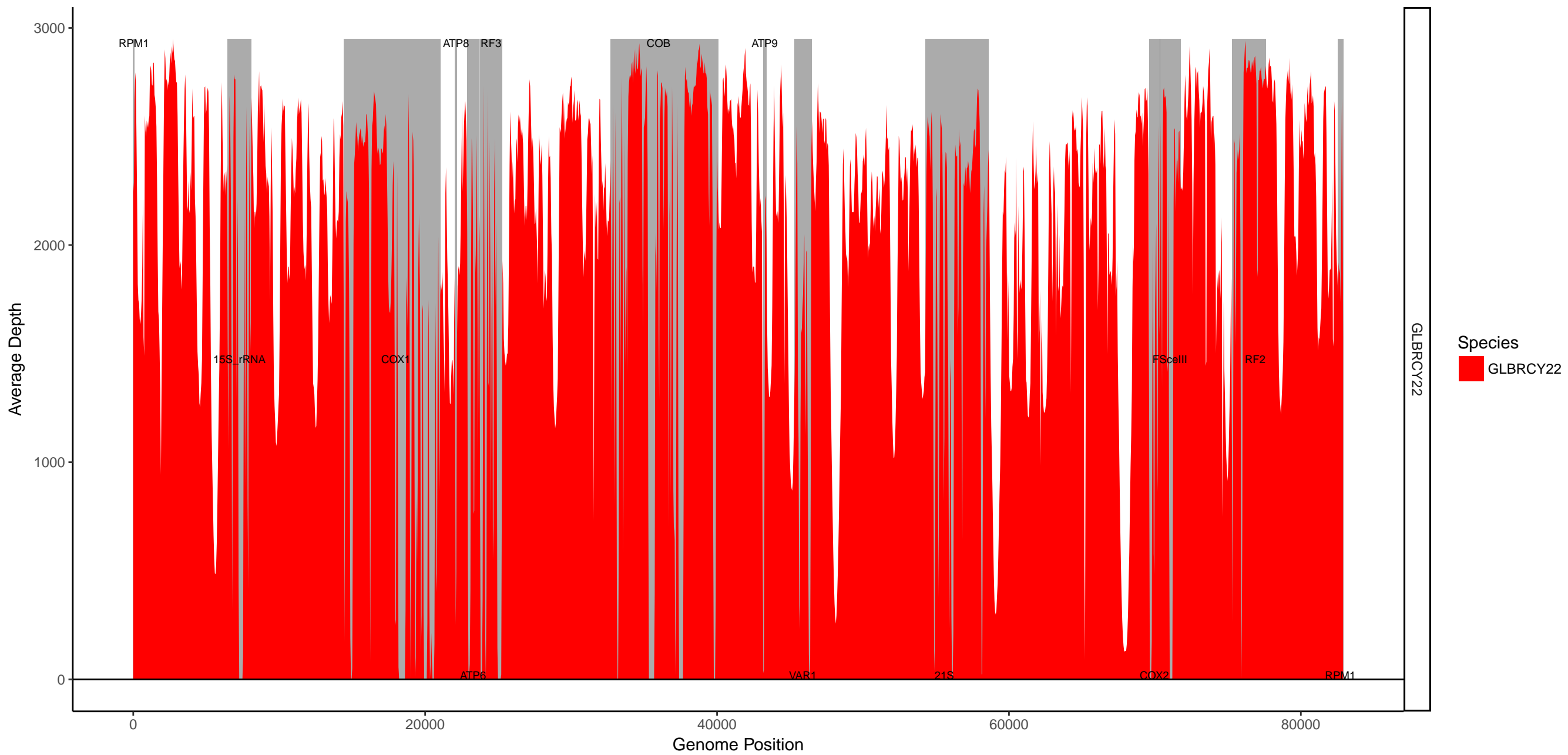


The figure displays six horizontal tracks representing the genomes of different *Aspergillus nidulans* strains. The x-axis at the bottom indicates the 'Genome Position' from 0 to 80,000. Each track shows the presence (grey bar) or absence (white space) of specific genes. The genes are labeled within the tracks, often with their corresponding gene names in a smaller font above or below the bar. The strains are identified on the right side of each track: GLBRCY22, CBS432, IFO1815, ZP591, CBS10644, and CBS7001. The top track (GLBRCY22) is a red line plot, likely representing a signal such as coverage or expression. The other tracks are grey bar plots. The genes shown include: RPM1, 15S_rRNA, COX1, ATP8, RF3, COB, ATP9, VAR1, 21S, FScellIII, RF2, COX2, and RPM1. The bottom track (CBS7001) shows the presence of 21S, COX2, FSuvallI, COX3, COB, COX1, ATP6, ATP9, VAR1, and 15S.

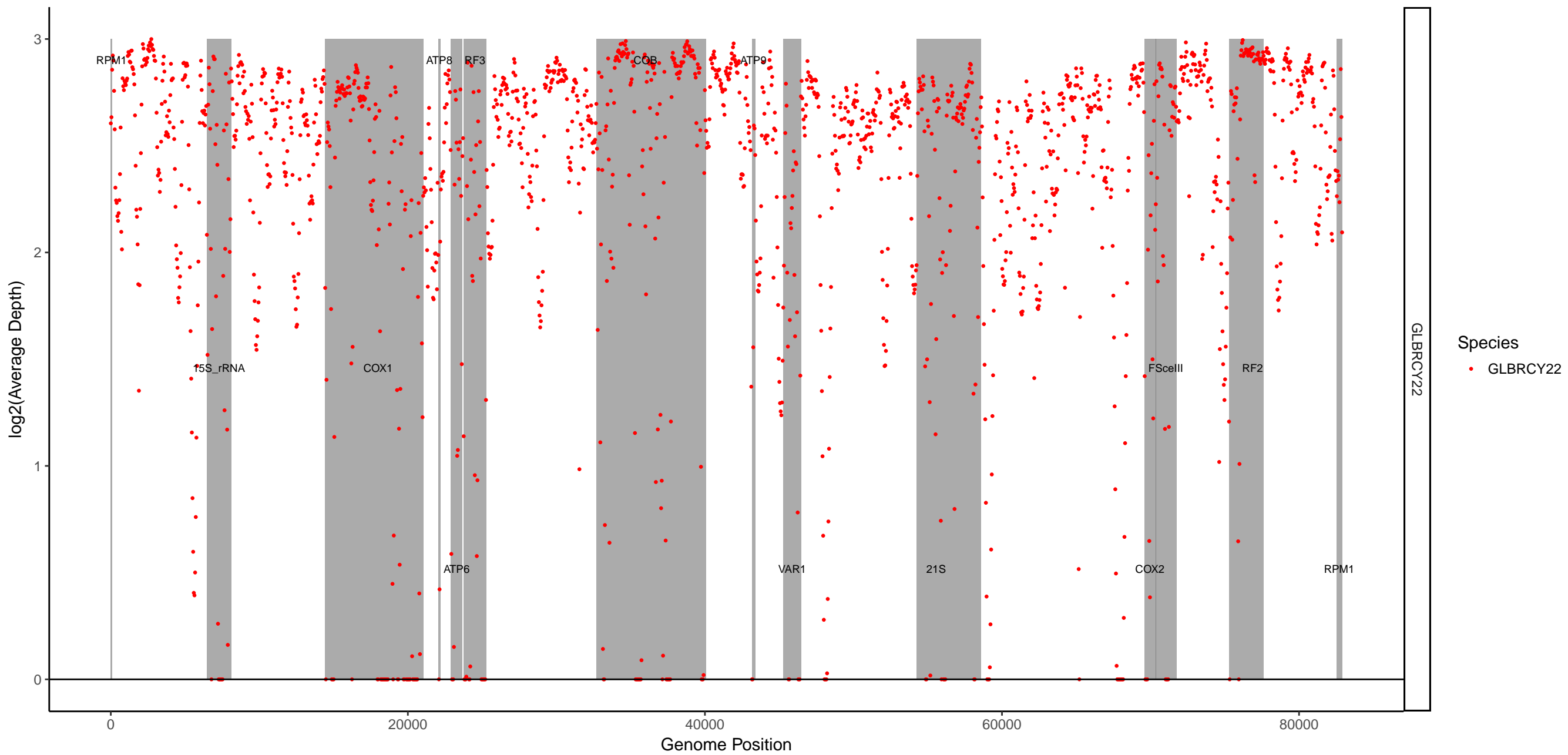
GLBRCY22
CBS432
IFO1815
ZP591
CBS10644
CBS7001



yHRWh87 Avg depth of coverage



yHRWh87 Avg depth of coverage



yHRWh87 Avg depth of coverage

