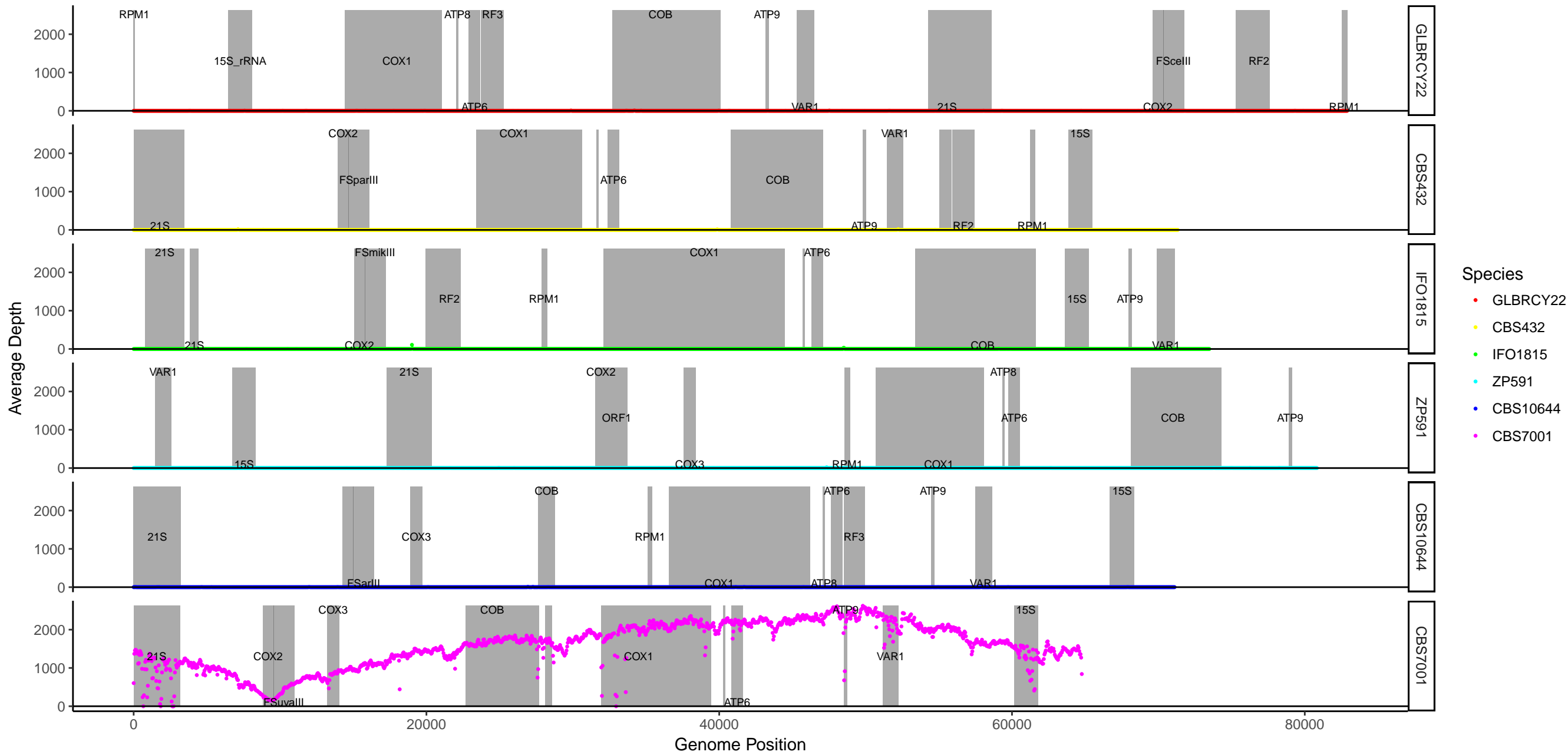
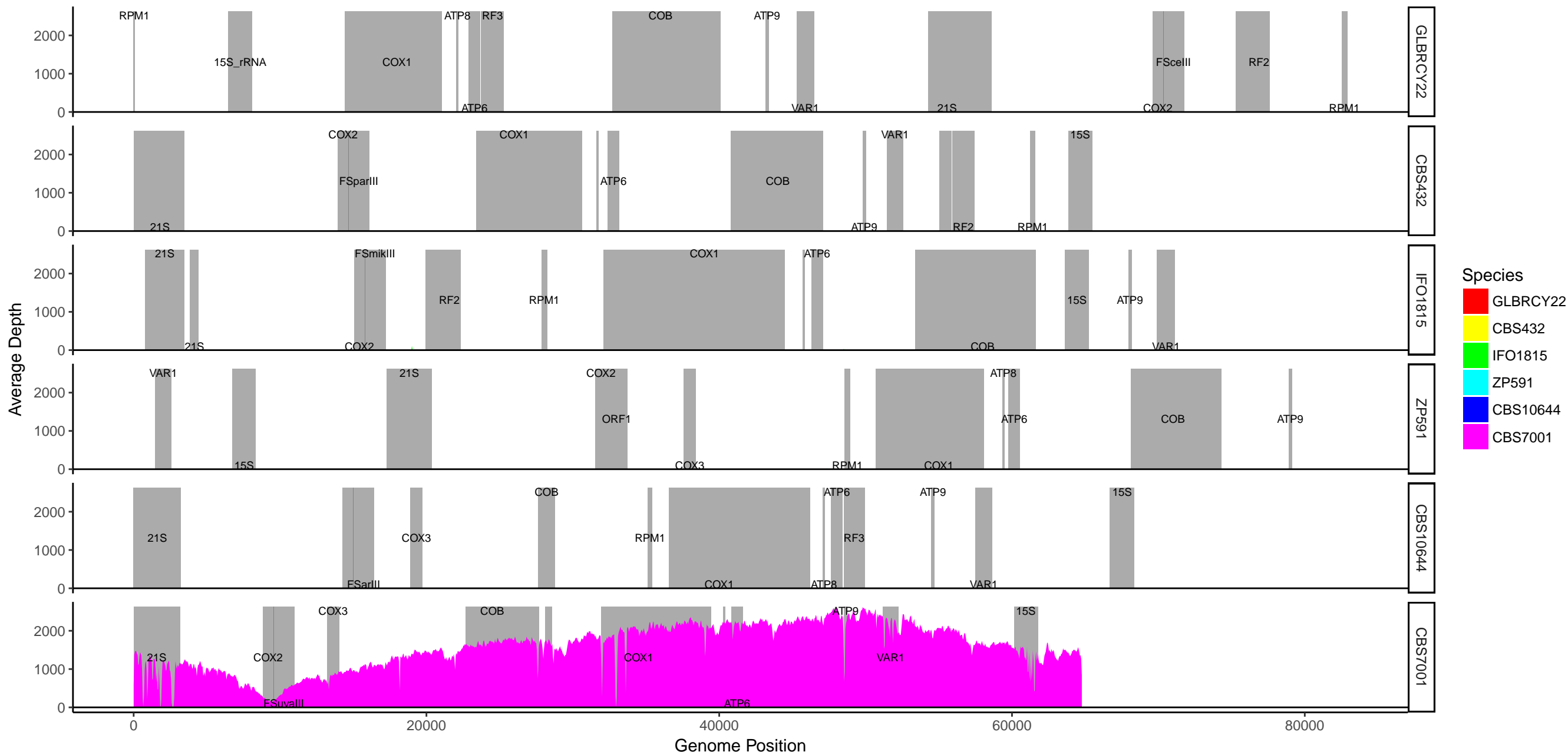


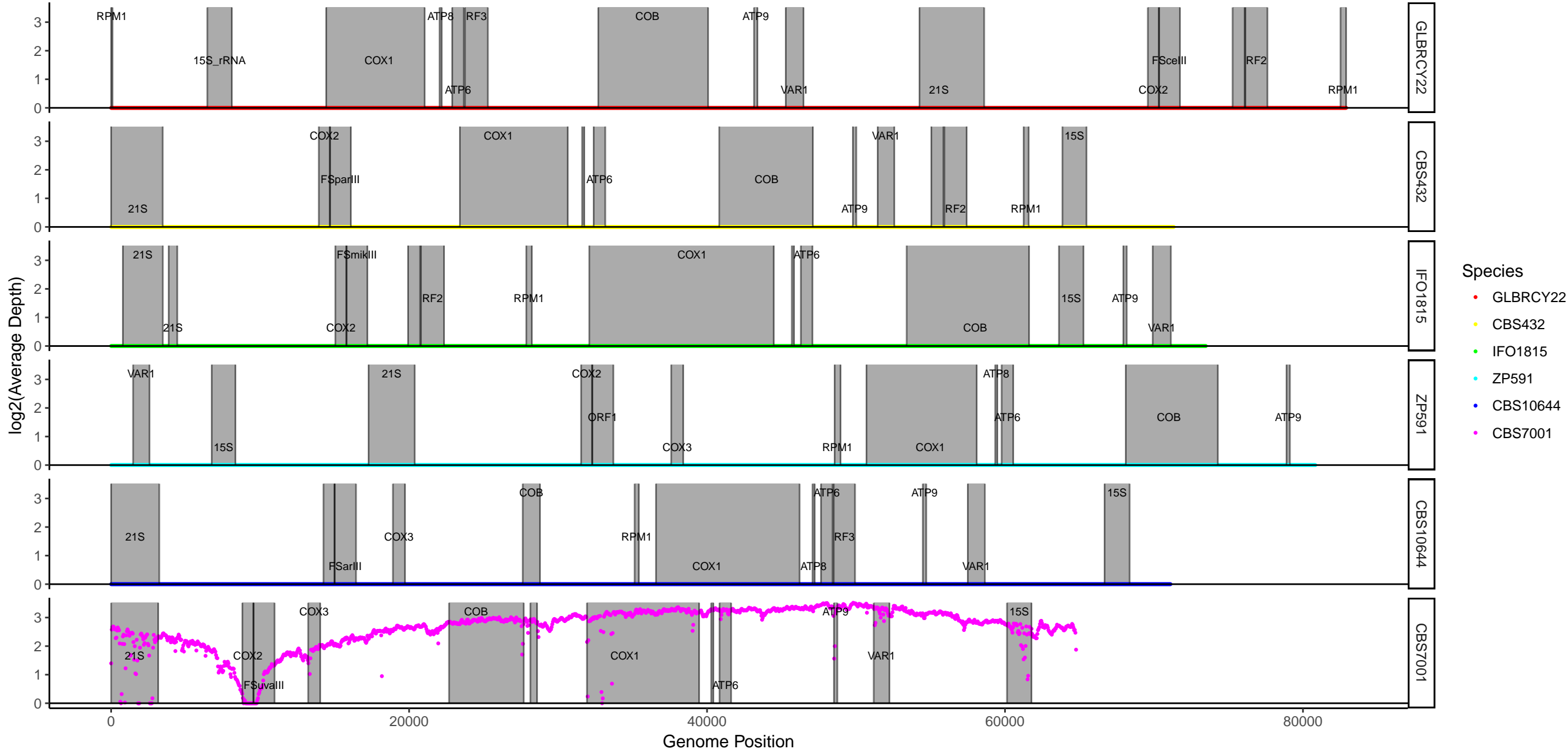
yHRWh51 Avg depth of coverage



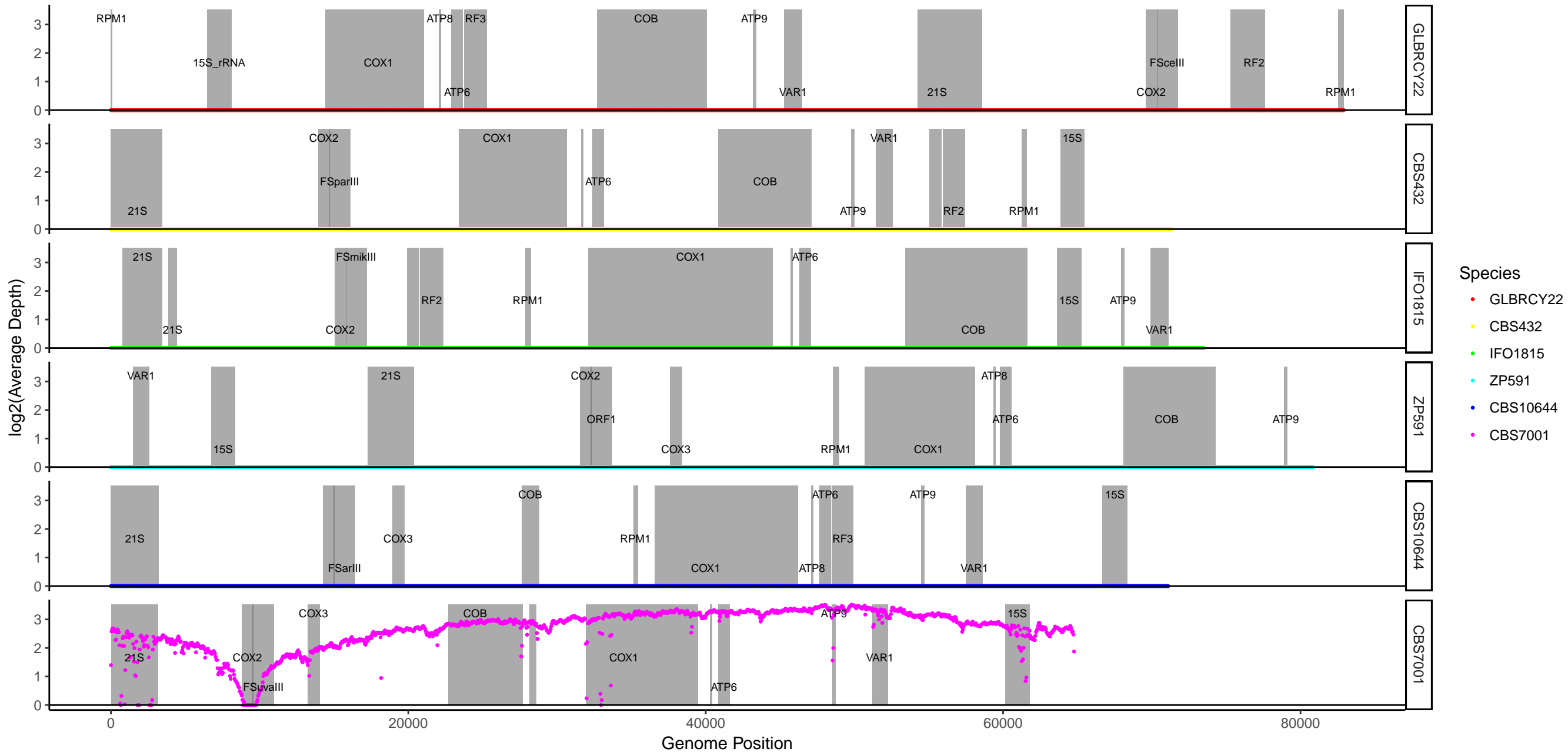
yHRWh51 Avg depth of coverage



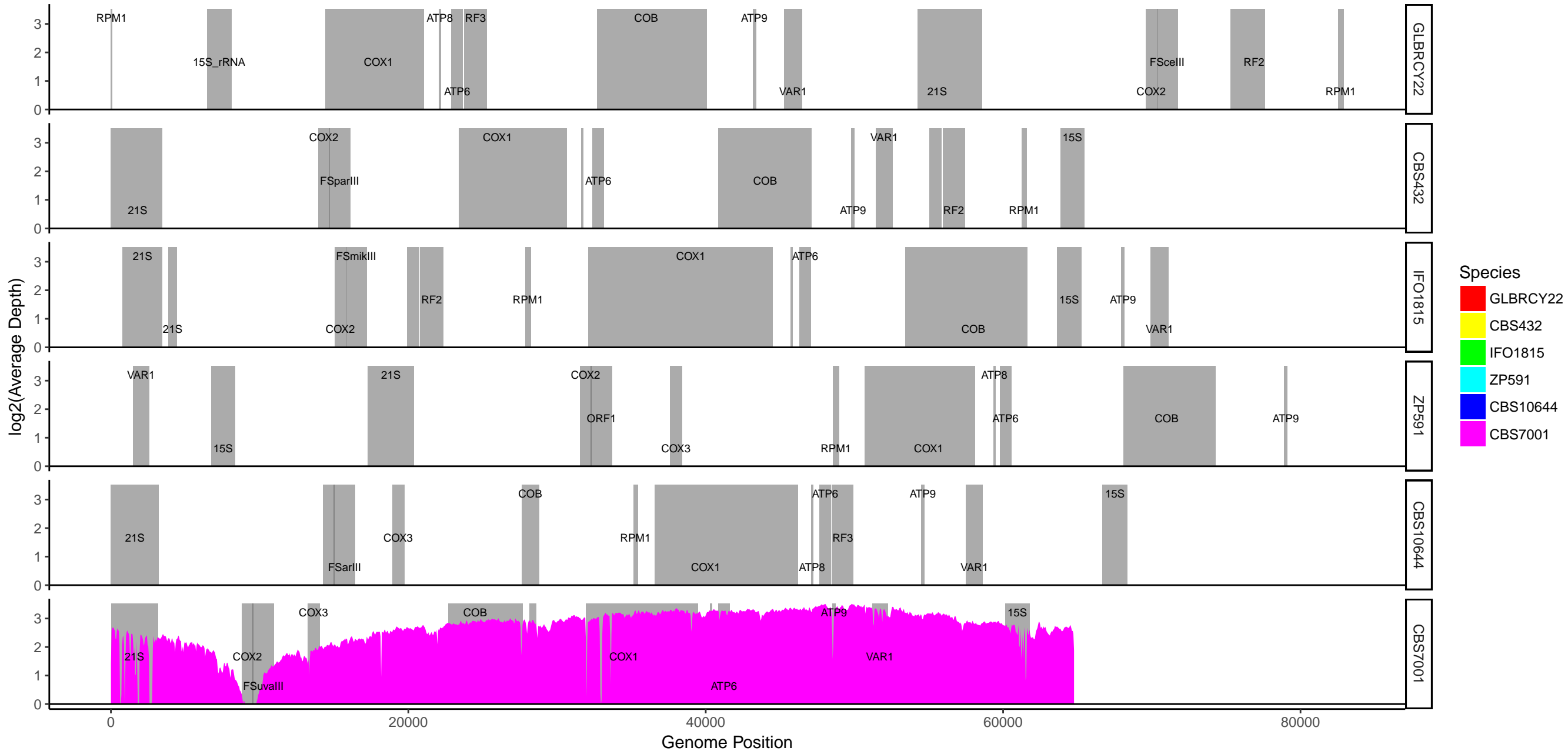
yHRWh51 Log2 Avg depth of coverage



yHRWh51 Log2 Avg depth of coverage



yHRWh51 Avg depth of coverage



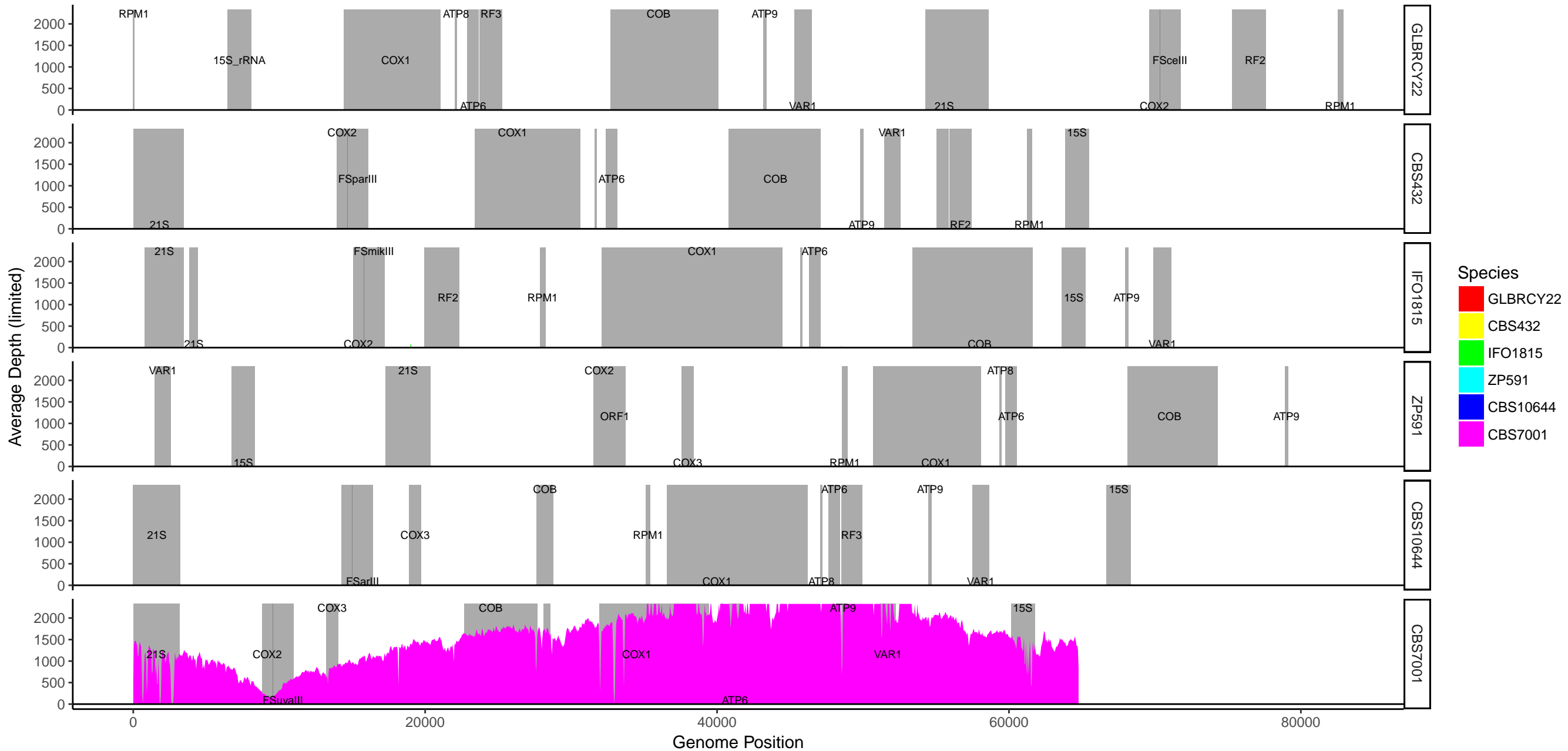
The figure displays a genomic map of the mitochondrial genome of *Glomeribacter geophilum*, showing the locations and relative expression levels (RPM) of various genes across six different strains. The x-axis represents the Genome Position (0 to 80,000). The y-axis represents the RPM (Relative Protein Mass).

The genes shown are: 15S, 15S_rRNA, COX1, COX2, COX3, ORF1, ORF2, ORF3, ORF4, ORF5, ORF6, ORF7, ORF8, ORF9, ORF10, ORF11, ORF12, ORF13, ORF14, ORF15, ORF16, ORF17, ORF18, ORF19, ORF20, ORF21, ORF22, ORF23, ORF24, ORF25, ORF26, ORF27, ORF28, ORF29, ORF30, ORF31, ORF32, ORF33, ORF34, ORF35, ORF36, ORF37, ORF38, ORF39, ORF40, ORF41, ORF42, ORF43, ORF44, ORF45, ORF46, ORF47, ORF48, ORF49, ORF50, ORF51, ORF52, ORF53, ORF54, ORF55, ORF56, ORF57, ORF58, ORF59, ORF60, ORF61, ORF62, ORF63, ORF64, ORF65, ORF66, ORF67, ORF68, ORF69, ORF70, ORF71, ORF72, ORF73, ORF74, ORF75, ORF76, ORF77, ORF78, ORF79, ORF80, ORF81, ORF82, ORF83, ORF84, ORF85, ORF86, ORF87, ORF88, ORF89, ORF90, ORF91, ORF92, ORF93, ORF94, ORF95, ORF96, ORF97, ORF98, ORF99, ORF100.

The strains are: GLBRCY22, CBS432, IFO1815, ZP591, CBS10644, and CBS7001.

The expression levels (RPM) are indicated by the height of the bars. The color of the bars indicates the strain: GLBRCY22 (red), CBS432 (yellow), IFO1815 (green), ZP591 (cyan), CBS10644 (blue), and CBS7001 (magenta).

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



yHRWh51 Avg depth of coverage

