

yHRWh56 Avg depth of coverage

Average Depth

2000

1500

1000

500

0

*GLBRCY22*

*CBS432*

*IFO1815*

*ZP591*

*CBS10644*

*CBS7001*

*FM1318*

Genome Position

Species

● *GLBRCY22*

● *CBS432*

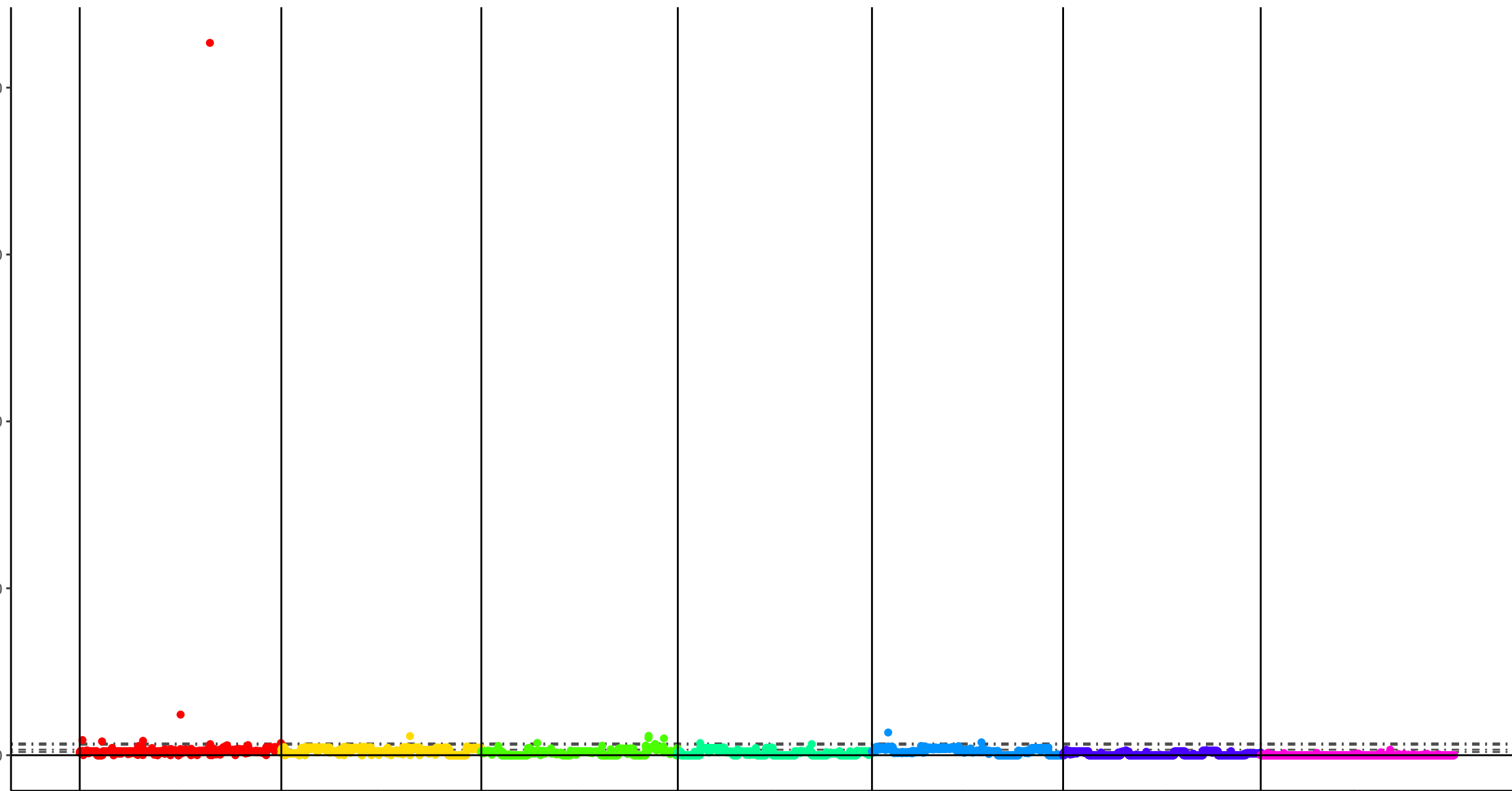
● *IFO1815*

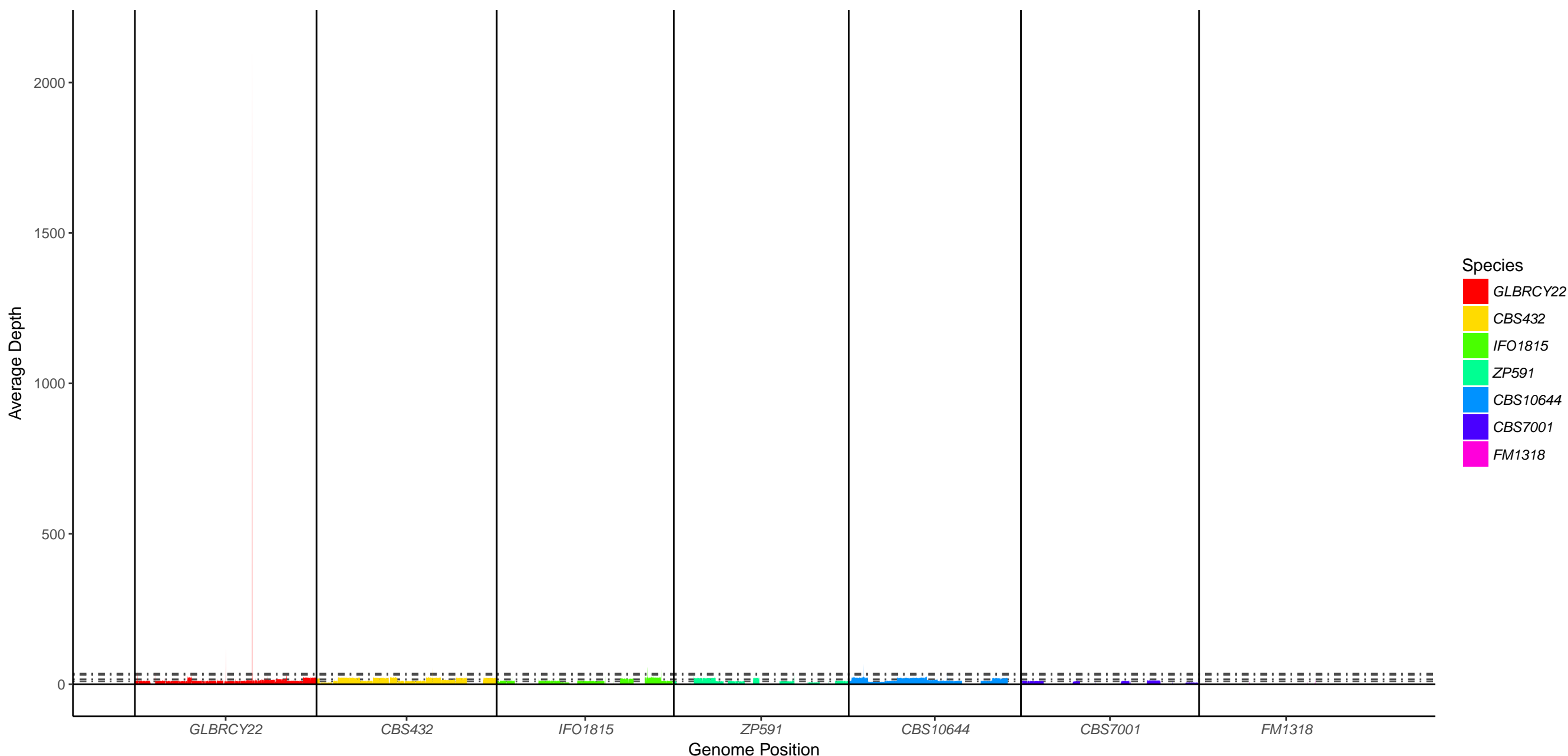
● *ZP591*

● *CBS10644*

● *CBS7001*

● *FM1318*





yHRWh56 Log2(Average Depth)

log2(Average Depth)

Species

● *GLBRCY22*

● *CBS432*

● *IFO1815*

● *ZP591*

● *CBS10644*

● *CBS7001*

● *FM1318*

*GLBRCY22*

*CBS432*

*IFO1815*

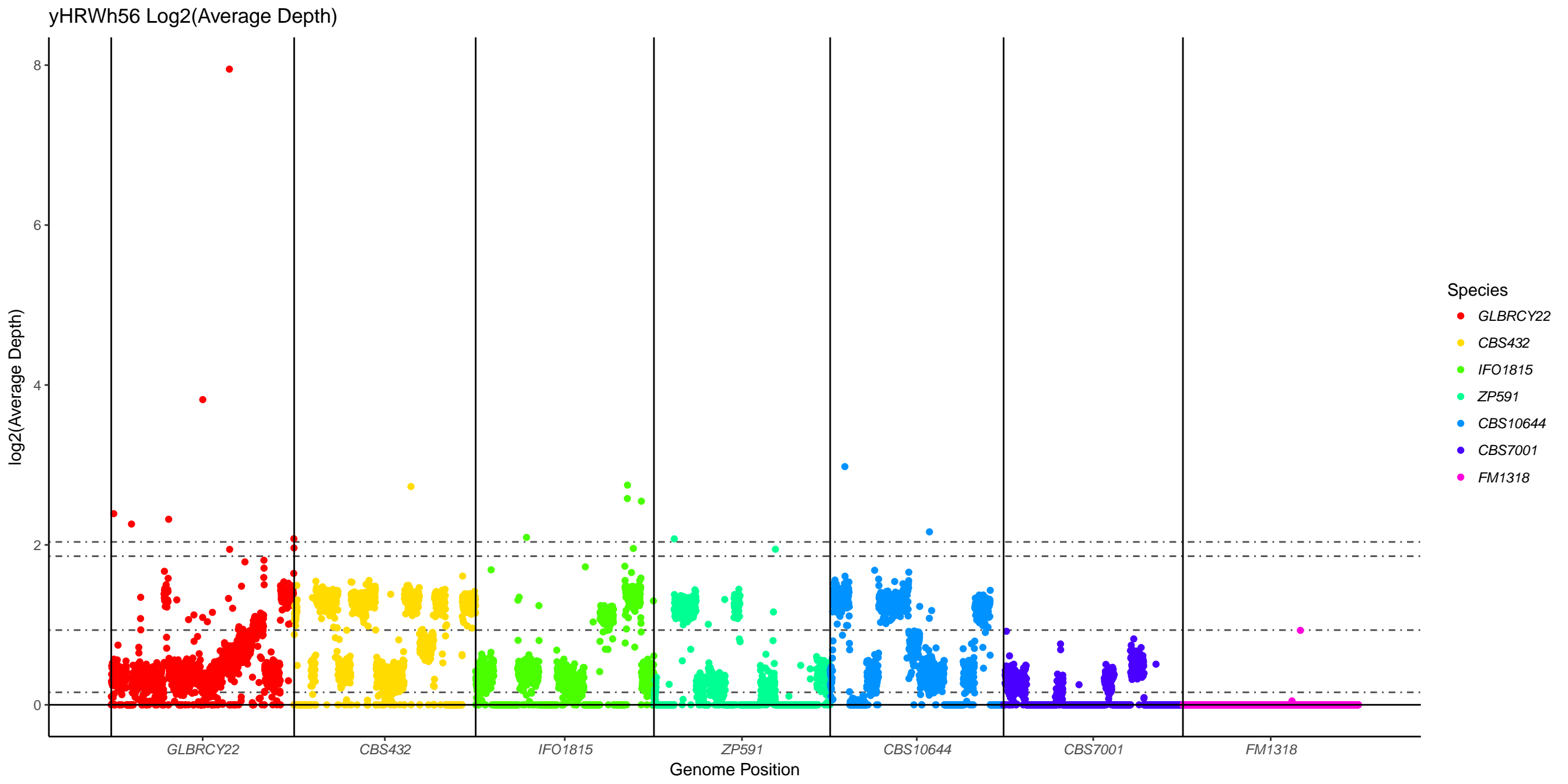
*ZP591*

*CBS10644*

*CBS7001*

*FM1318*

Genome Position



yHRWh56 Log2(Average Depth)

log2(Average Depth)

8  
6  
4  
2  
0

Species

█ *GLBRCY22*  
█ *CBS432*  
█ *IFO1815*  
█ *ZP591*  
█ *CBS10644*  
█ *CBS7001*  
█ *FM1318*

*GLBRCY22*

*CBS432*

*IFO1815*

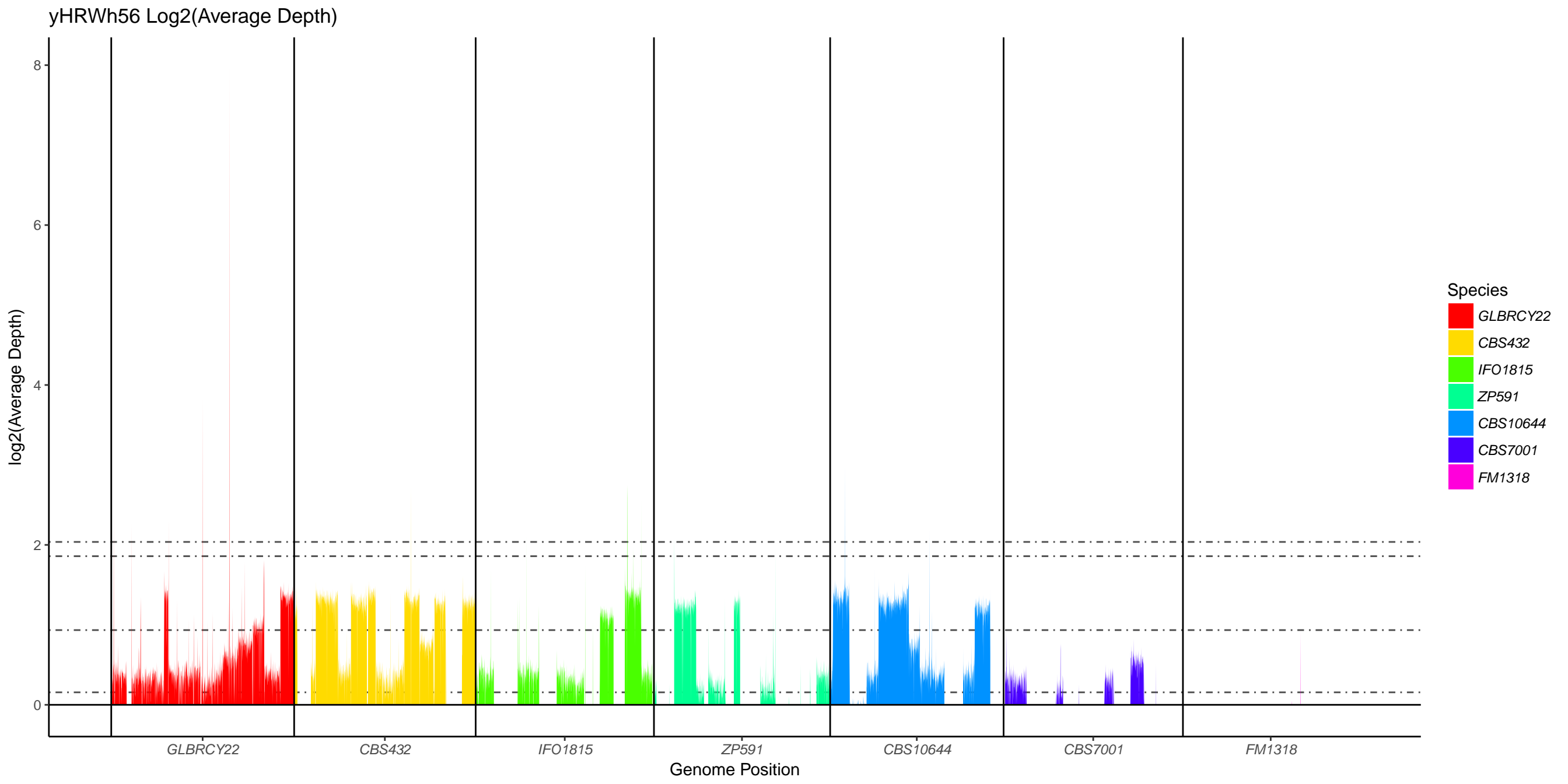
*ZP591*

*CBS10644*

*CBS7001*

*FM1318*

Genome Position

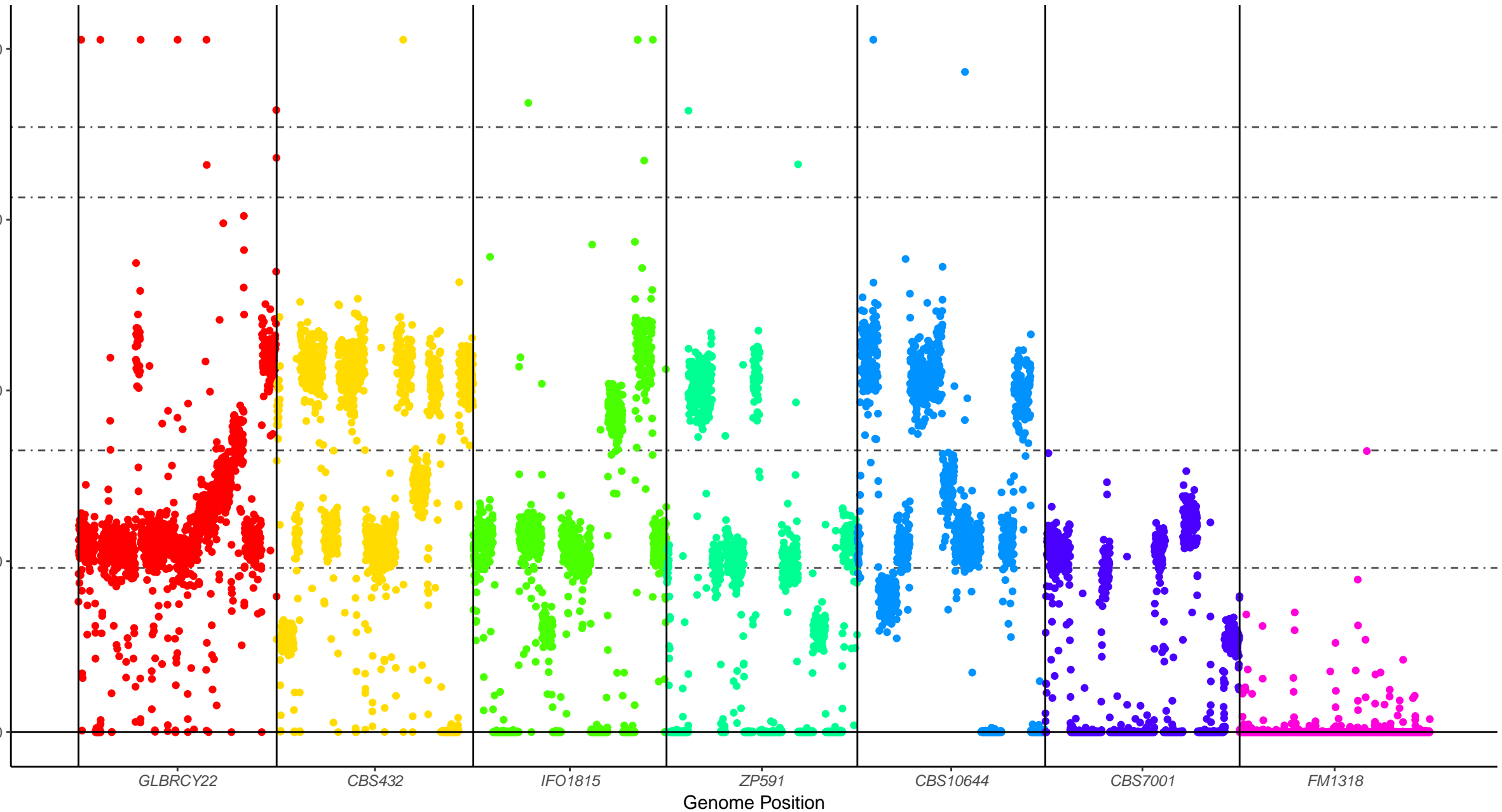


yHRWh56 Avg depth of coverage

Average Depth (limited)

Species

- GLBRCY22
- CBS432
- IFO1815
- ZP591
- CBS10644
- CBS7001
- FM1318



yHRWh56 Avg depth of coverage

Average Depth (limited)

40  
30  
20  
10  
0

Species

█ *GLBRCY22*  
█ *CBS432*  
█ *IFO1815*  
█ *ZP591*  
█ *CBS10644*  
█ *CBS7001*  
█ *FM1318*

*GLBRCY22*

*CBS432*

*IFO1815*

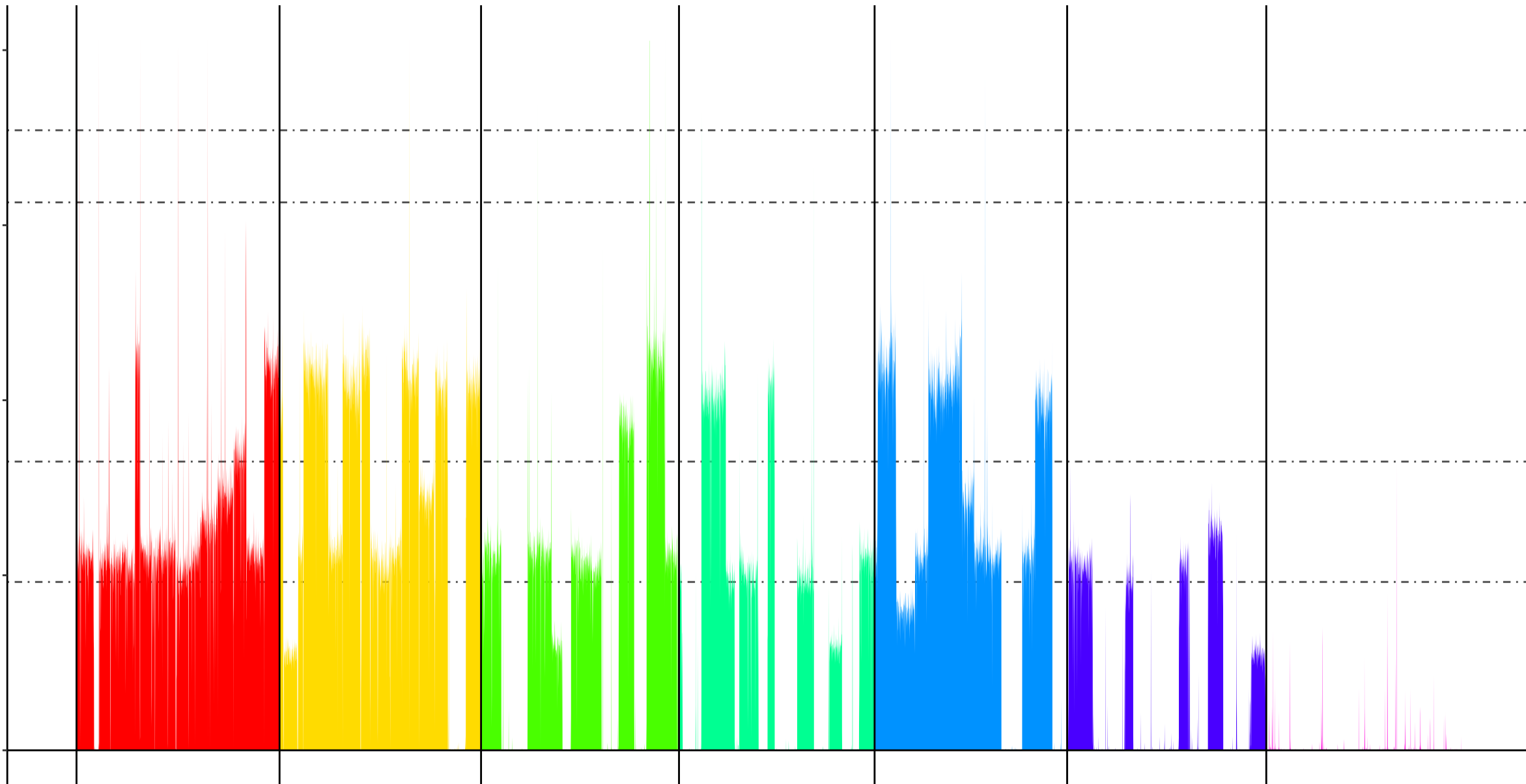
*ZP591*

*CBS10644*

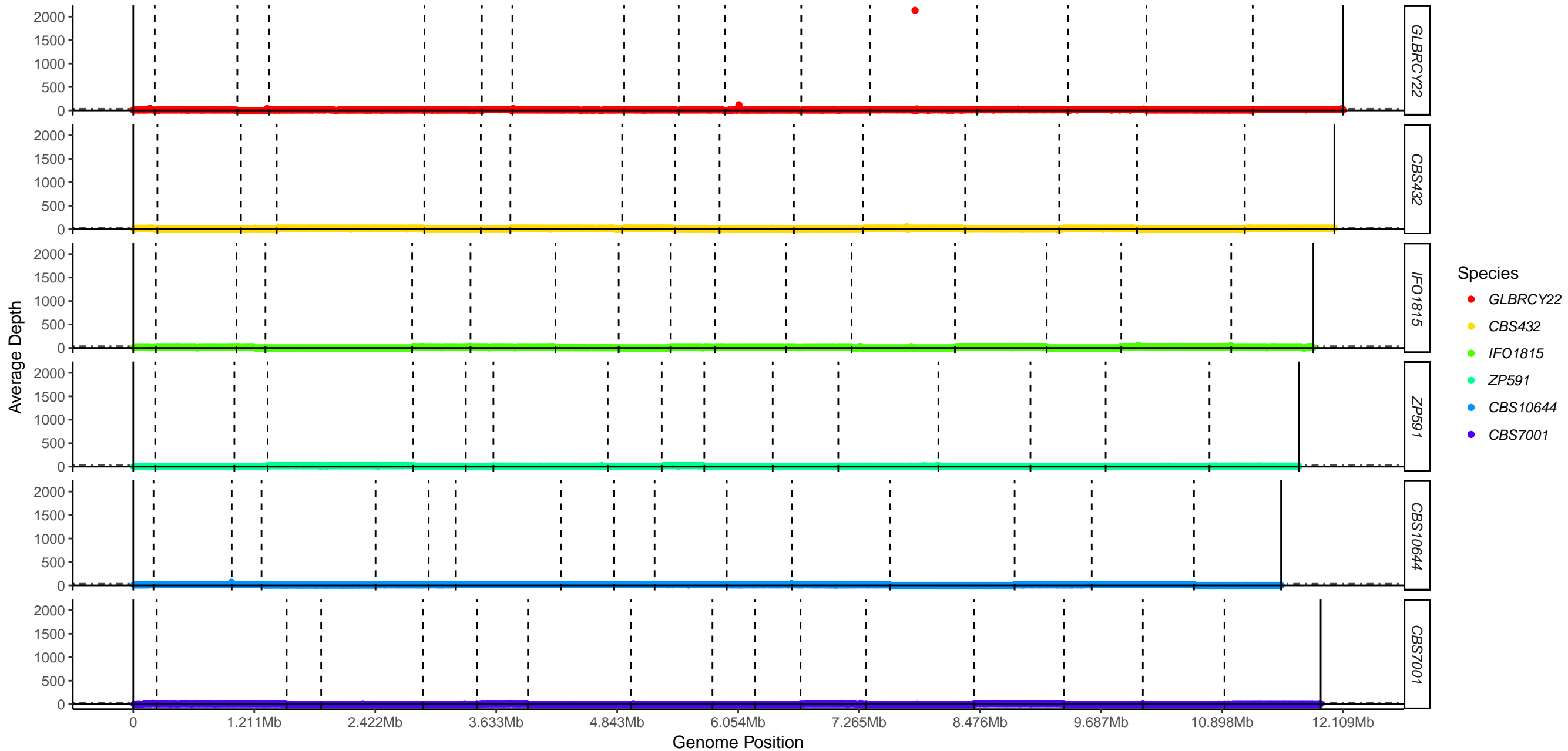
*CBS7001*

*FM1318*

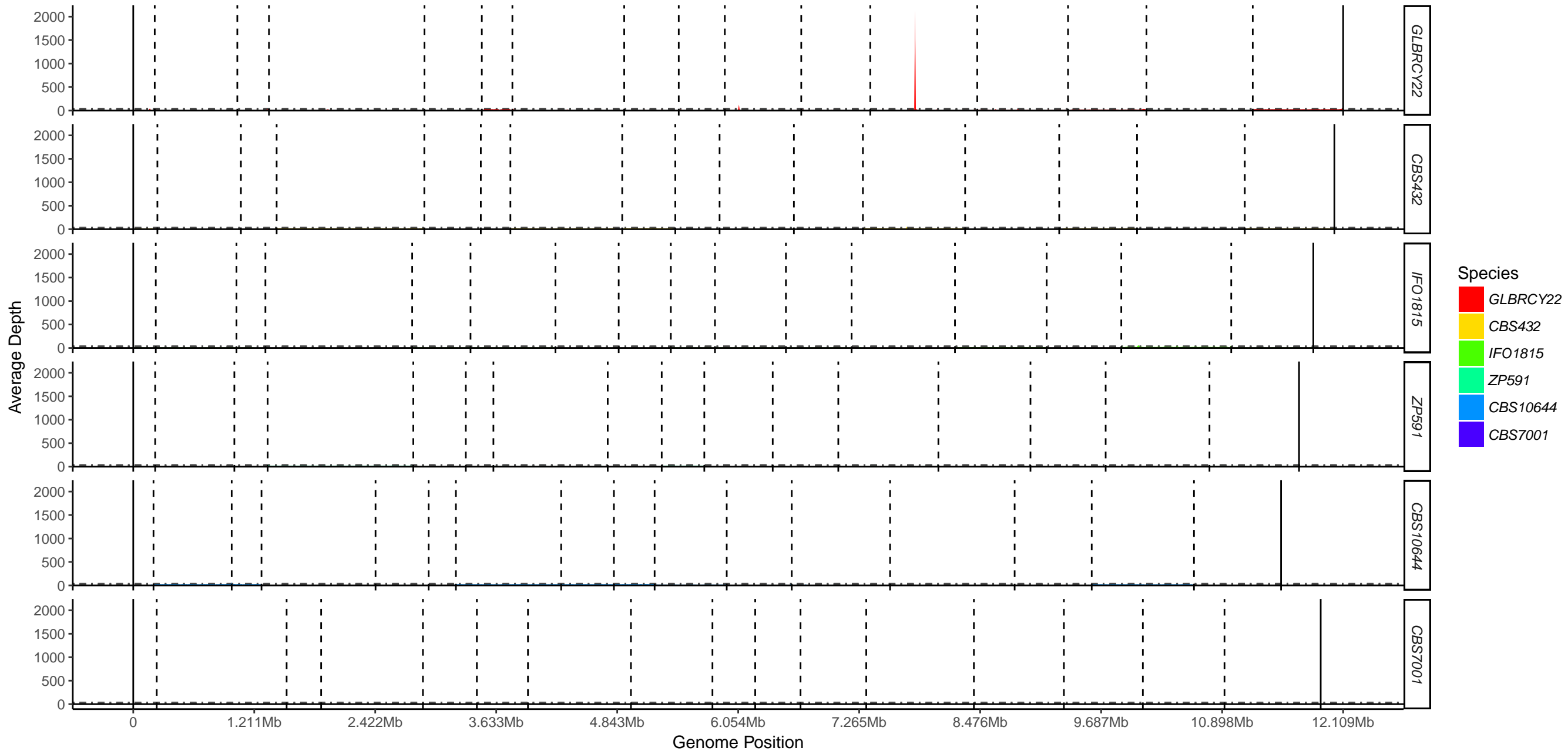
Genome Position



## yHRWh56 Avg depth of coverage

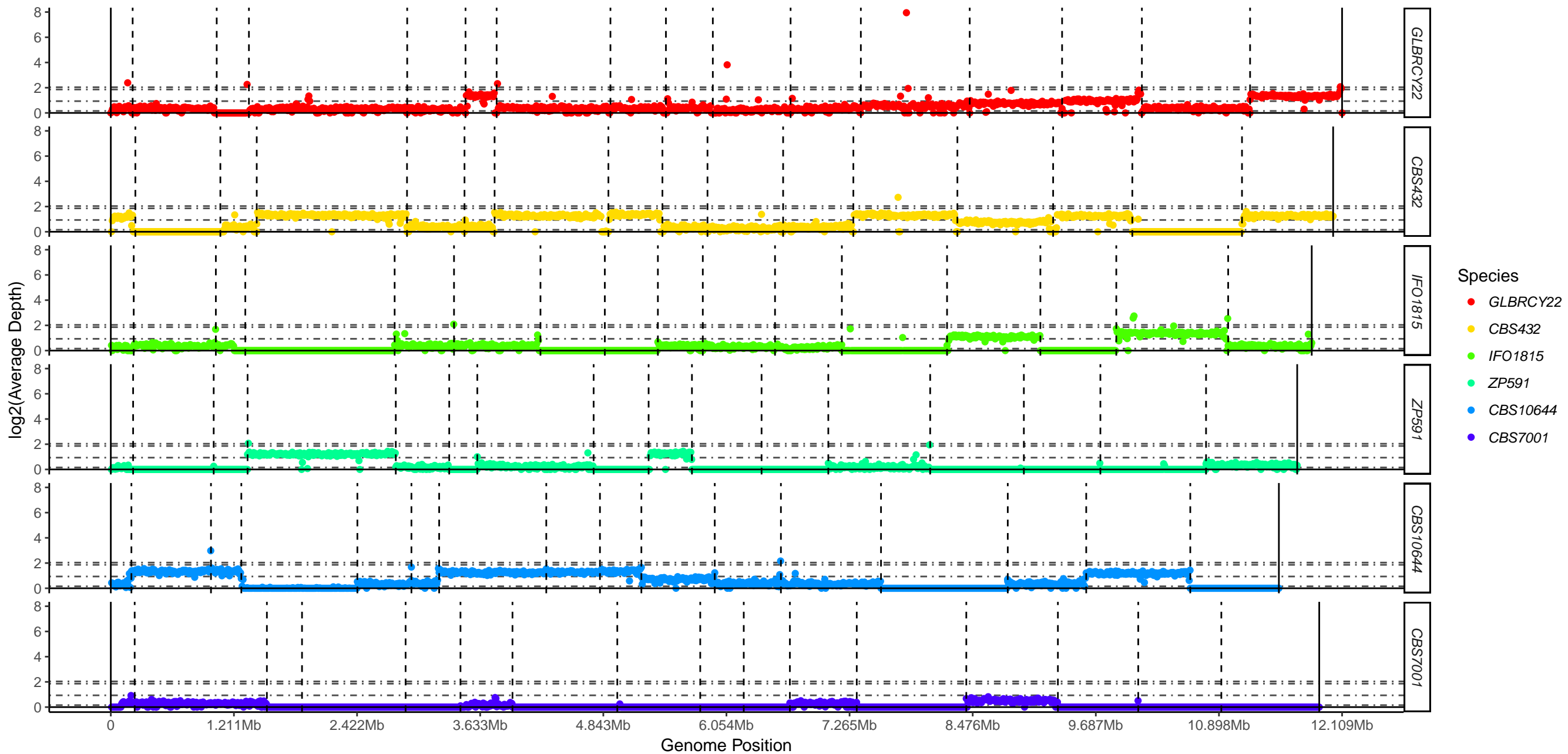


# yHRWh56 Avg depth of coverage

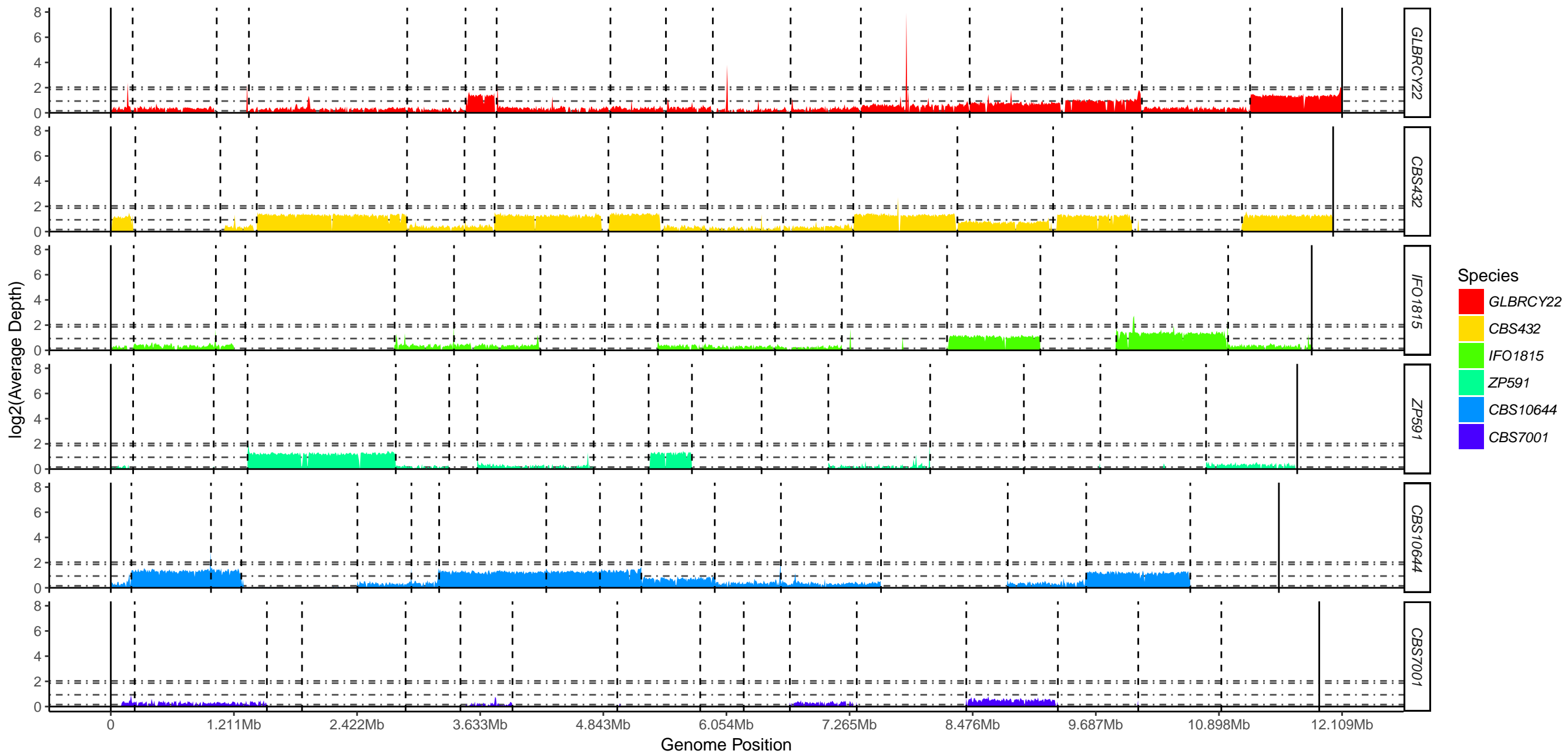




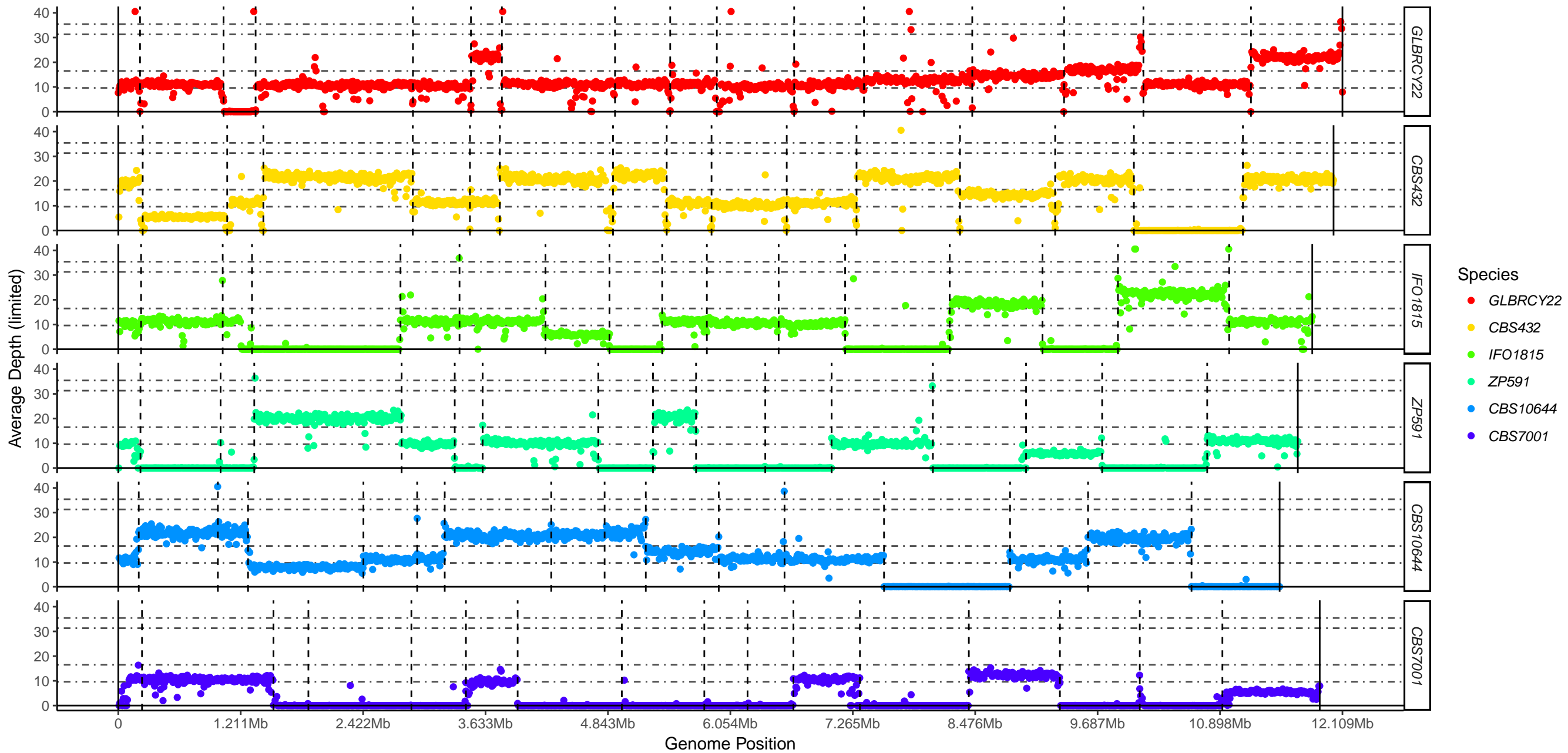
yHRWh56 Log2(Avg depth of coverage)



yHRWh56 Log2(Avg depth of coverage)



yHRWh56 Avg depth of coverage



yHRWh56 Avg depth of coverage

