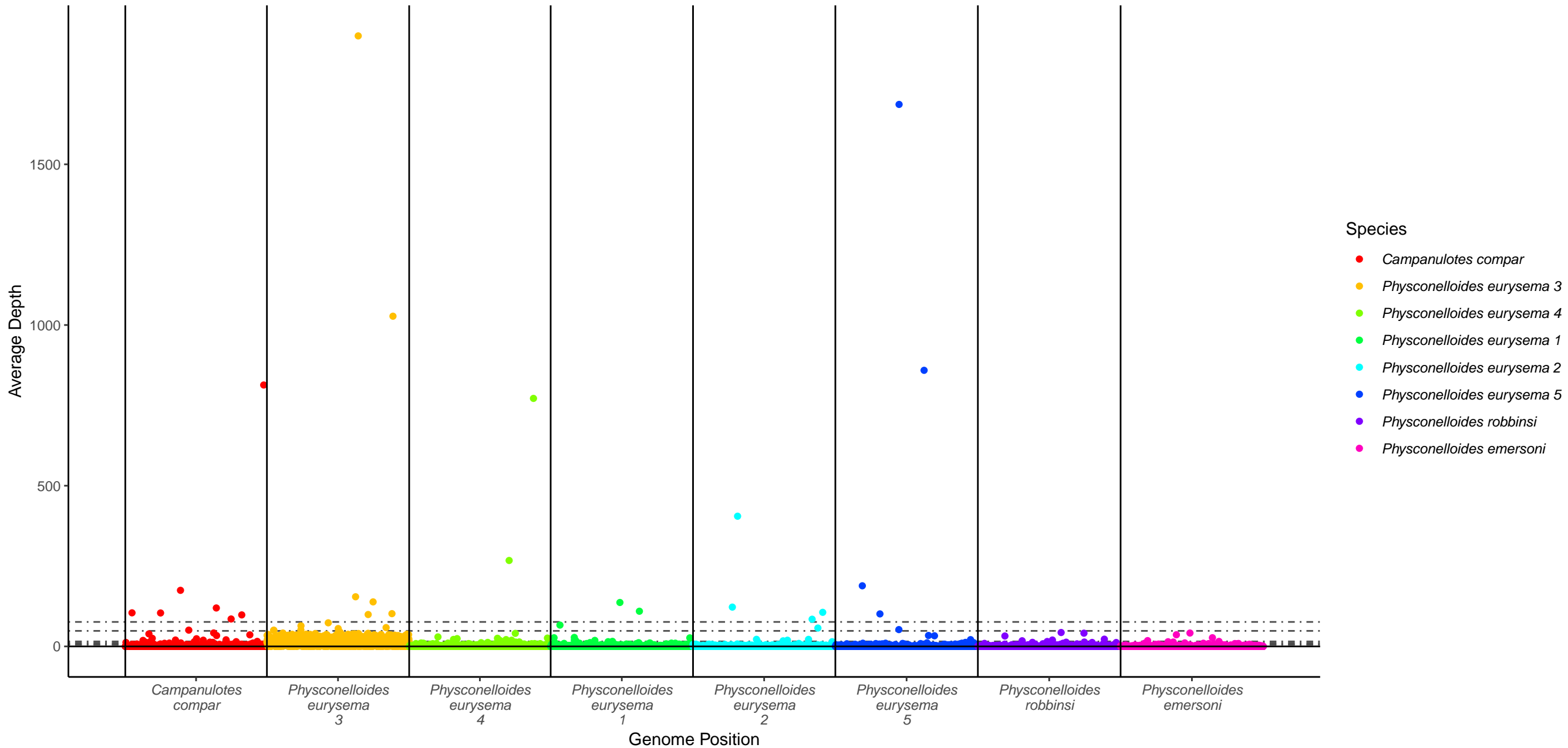
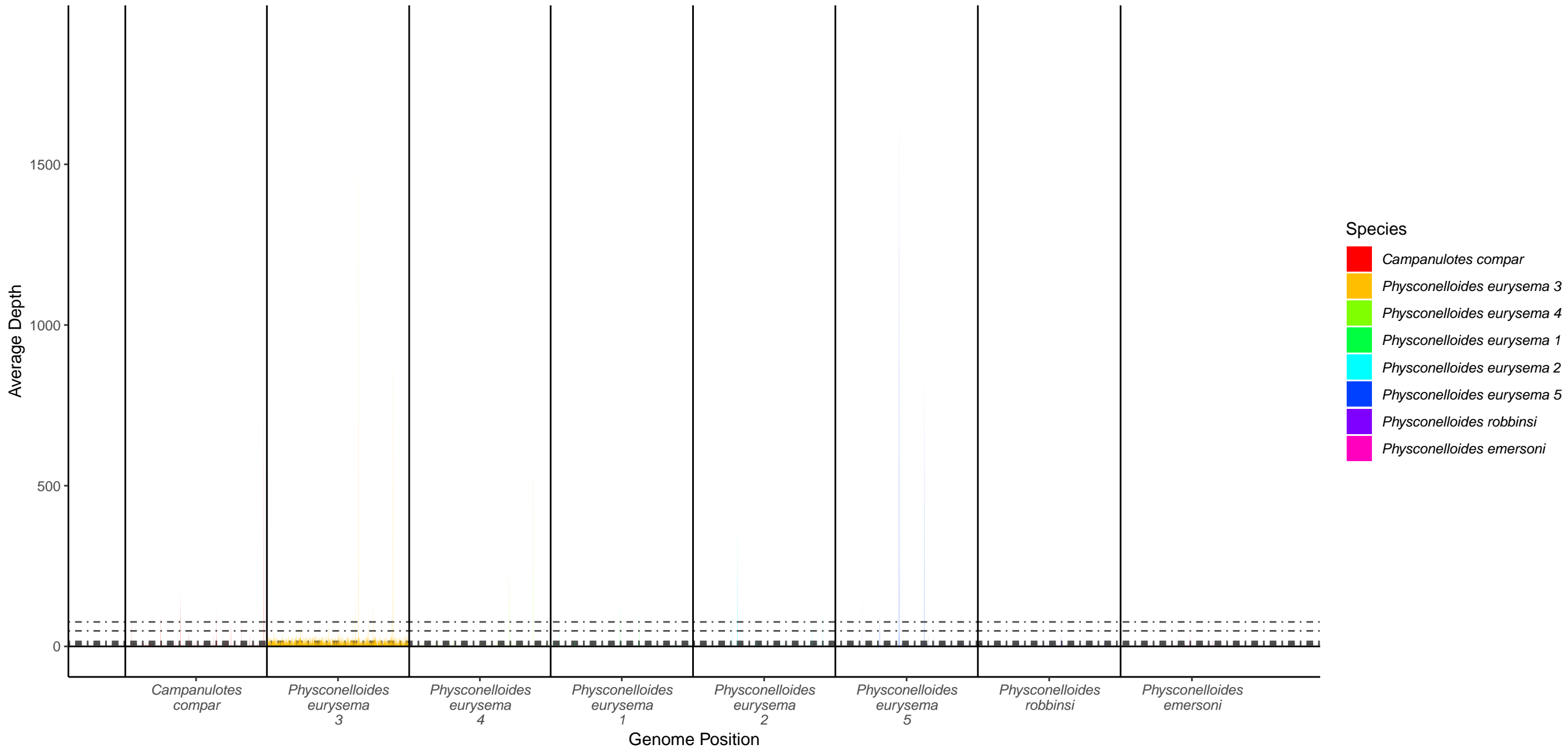


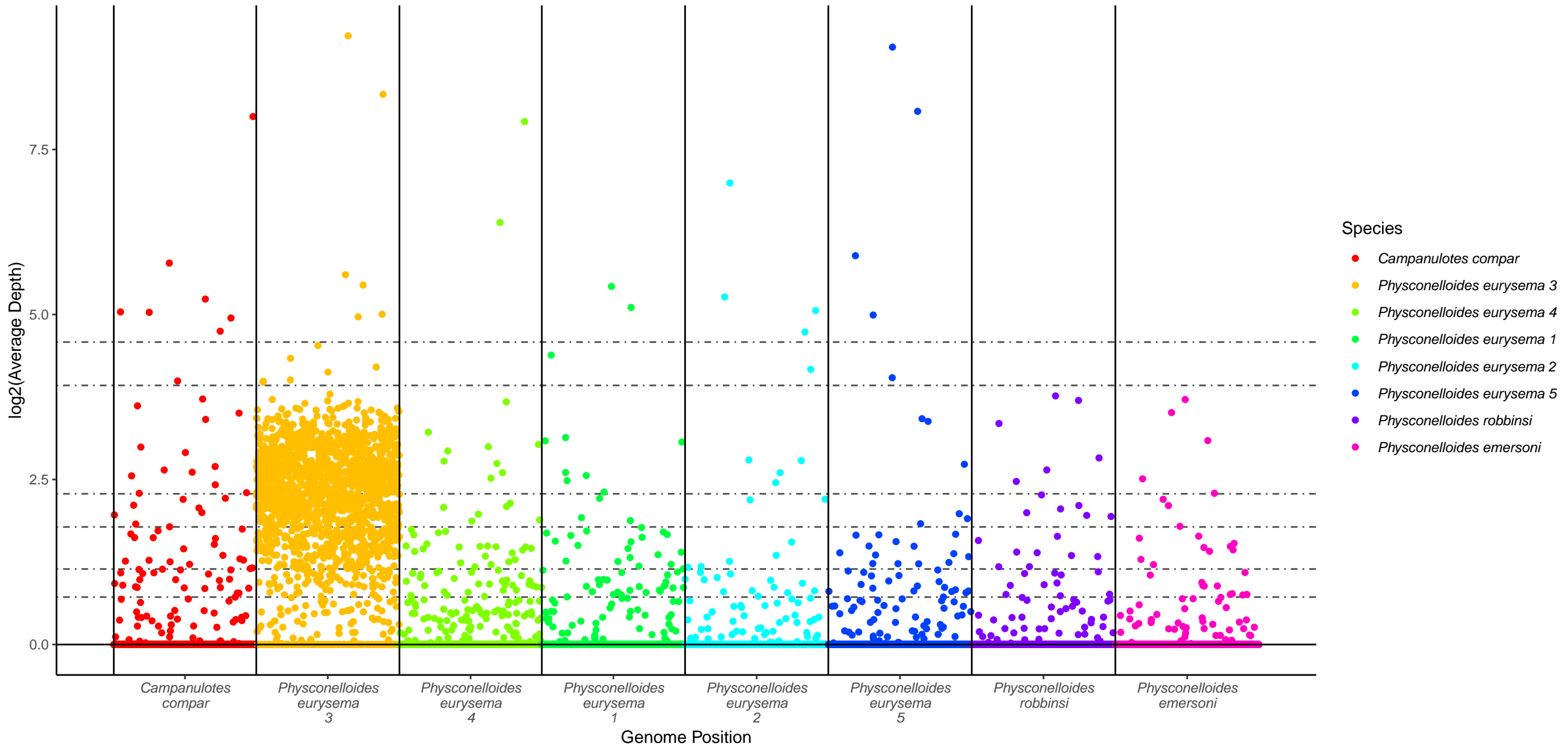
OUT Avg depth of coverage



OUT Avg depth of coverage



OUT Log2(Average Depth)



OUT Log2(Average Depth)

log2(Average Depth)

Species

- *Campanulotes compar*
- *Physconelloides eurysema 3*
- *Physconelloides eurysema 4*
- *Physconelloides eurysema 1*
- *Physconelloides eurysema 2*
- *Physconelloides eurysema 5*
- *Physconelloides robbinsi*
- *Physconelloides emersoni*

7.5

5.0

2.5

0.0

*Campanulotes
compar*

*Physconelloides
eurysema
3*

*Physconelloides
eurysema
4*

*Physconelloides
eurysema
1*

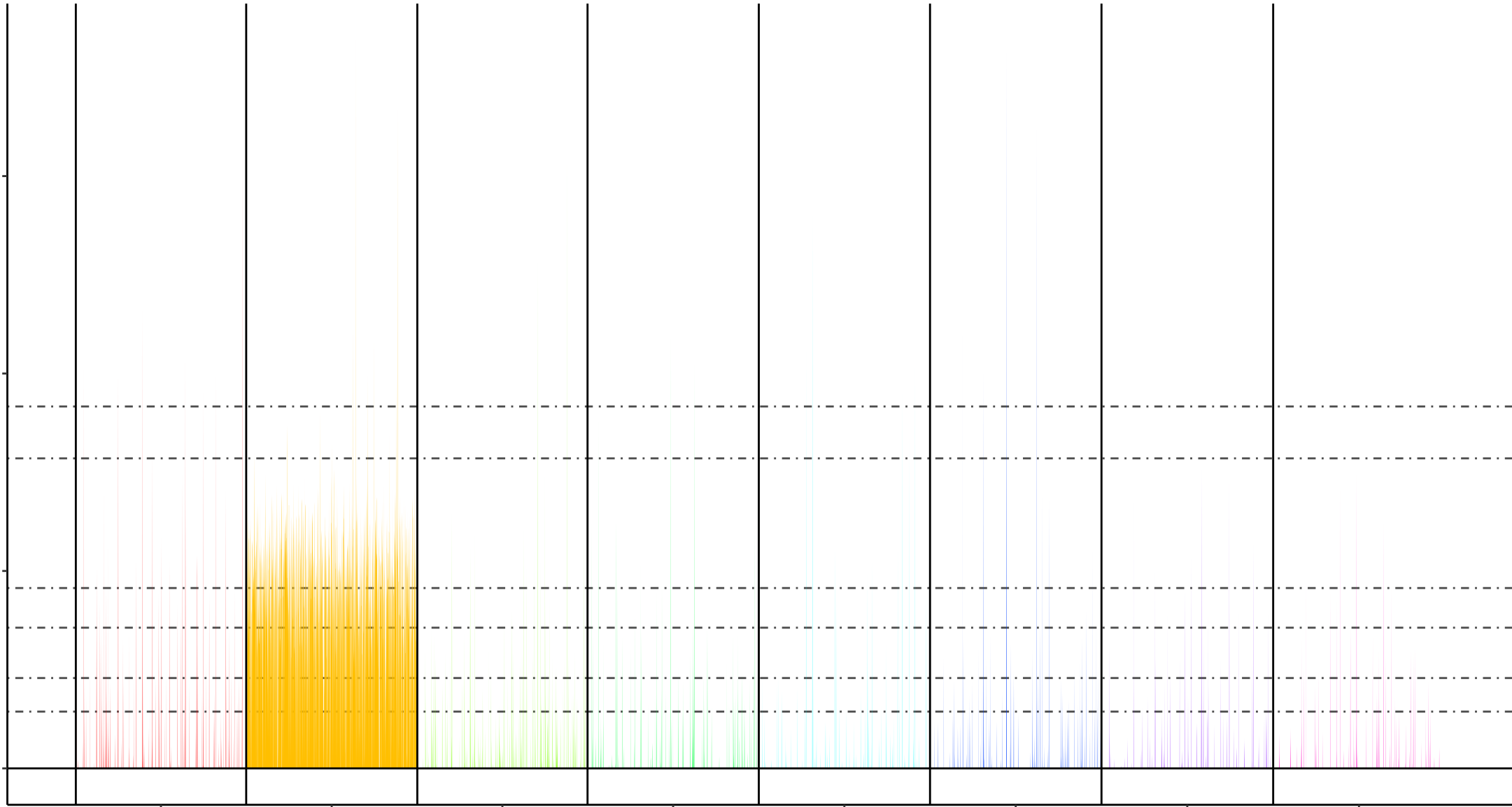
*Physconelloides
eurysema
2*

*Physconelloides
eurysema
5*

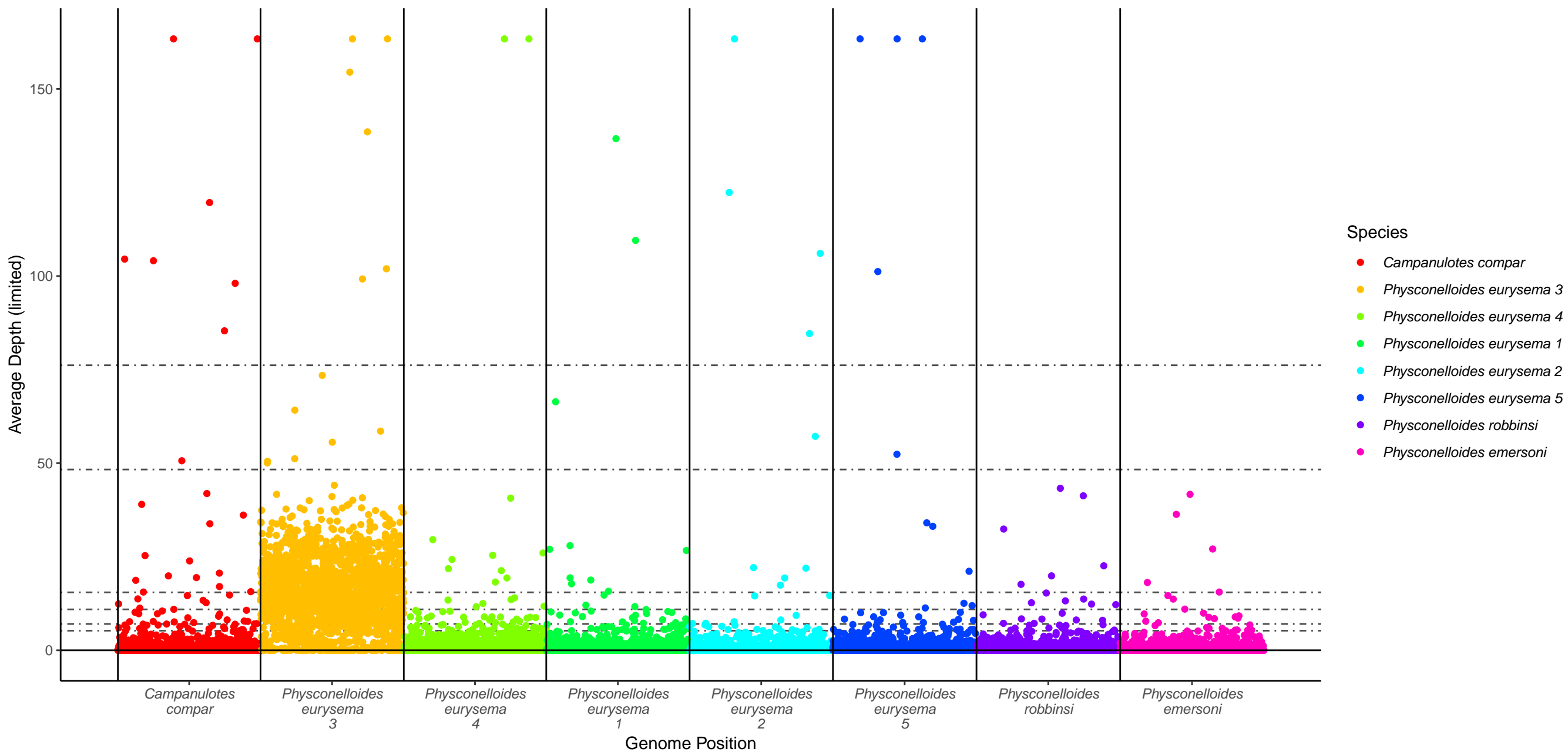
*Physconelloides
robbinsi*

*Physconelloides
emersoni*

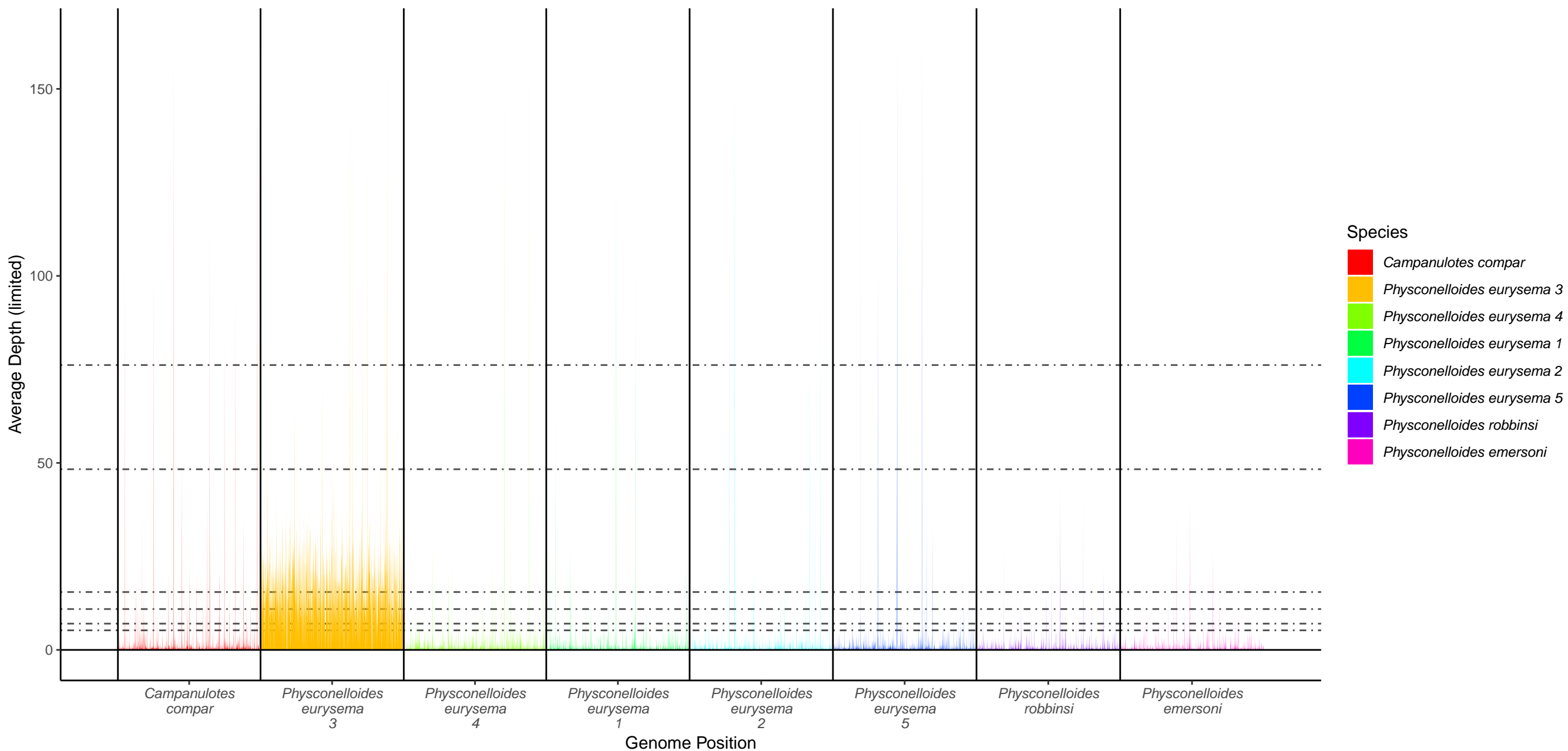
Genome Position



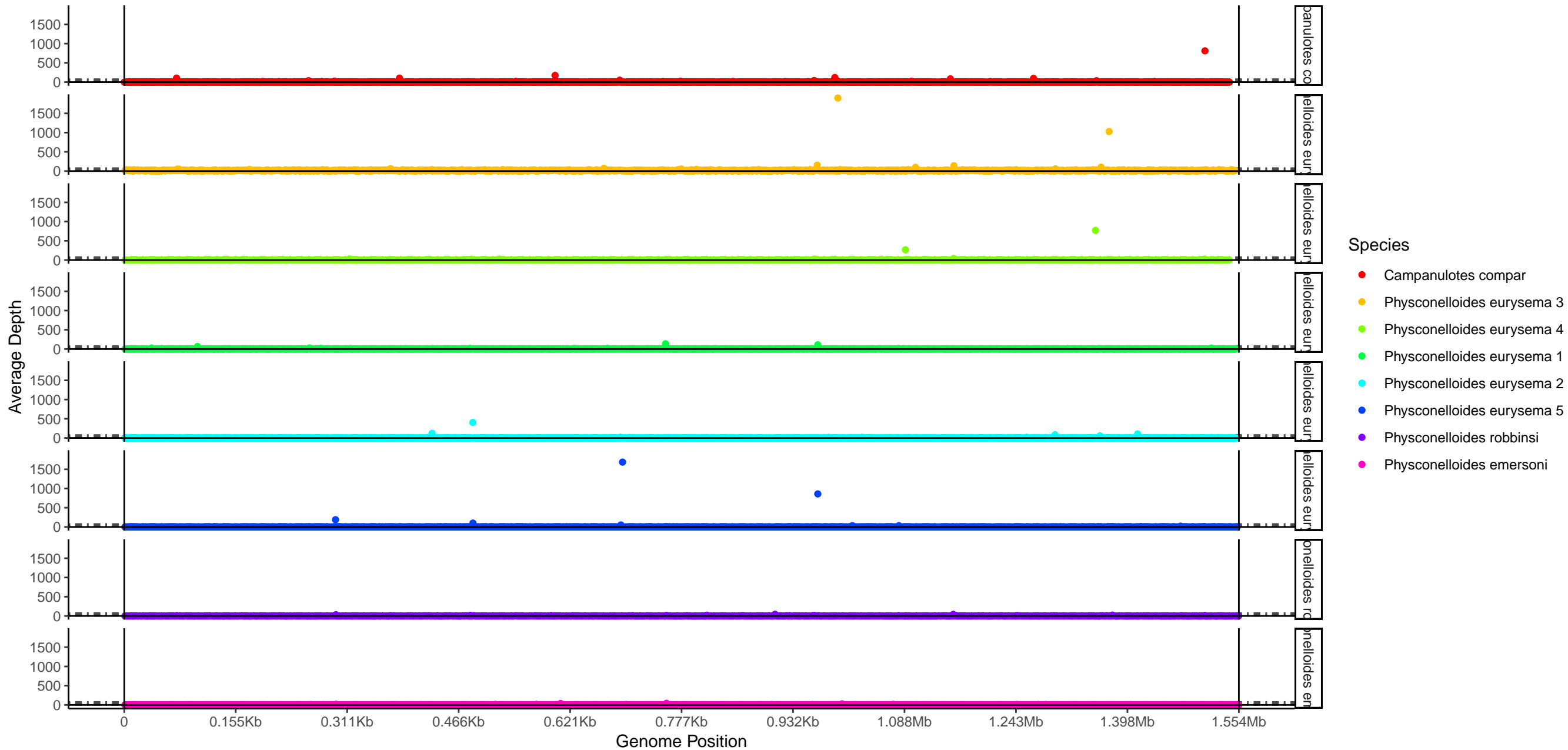
OUT Avg depth of coverage



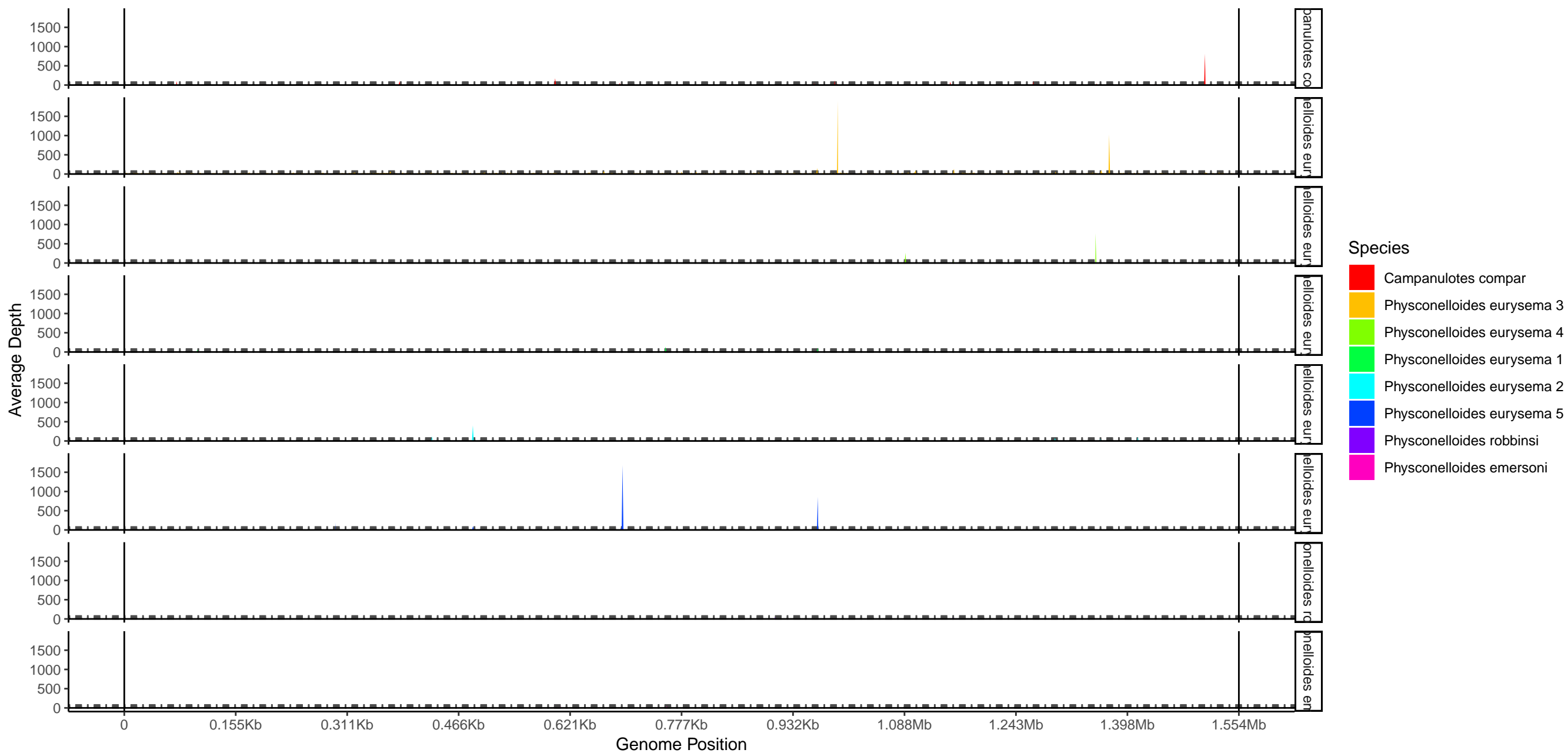
OUT Avg depth of coverage



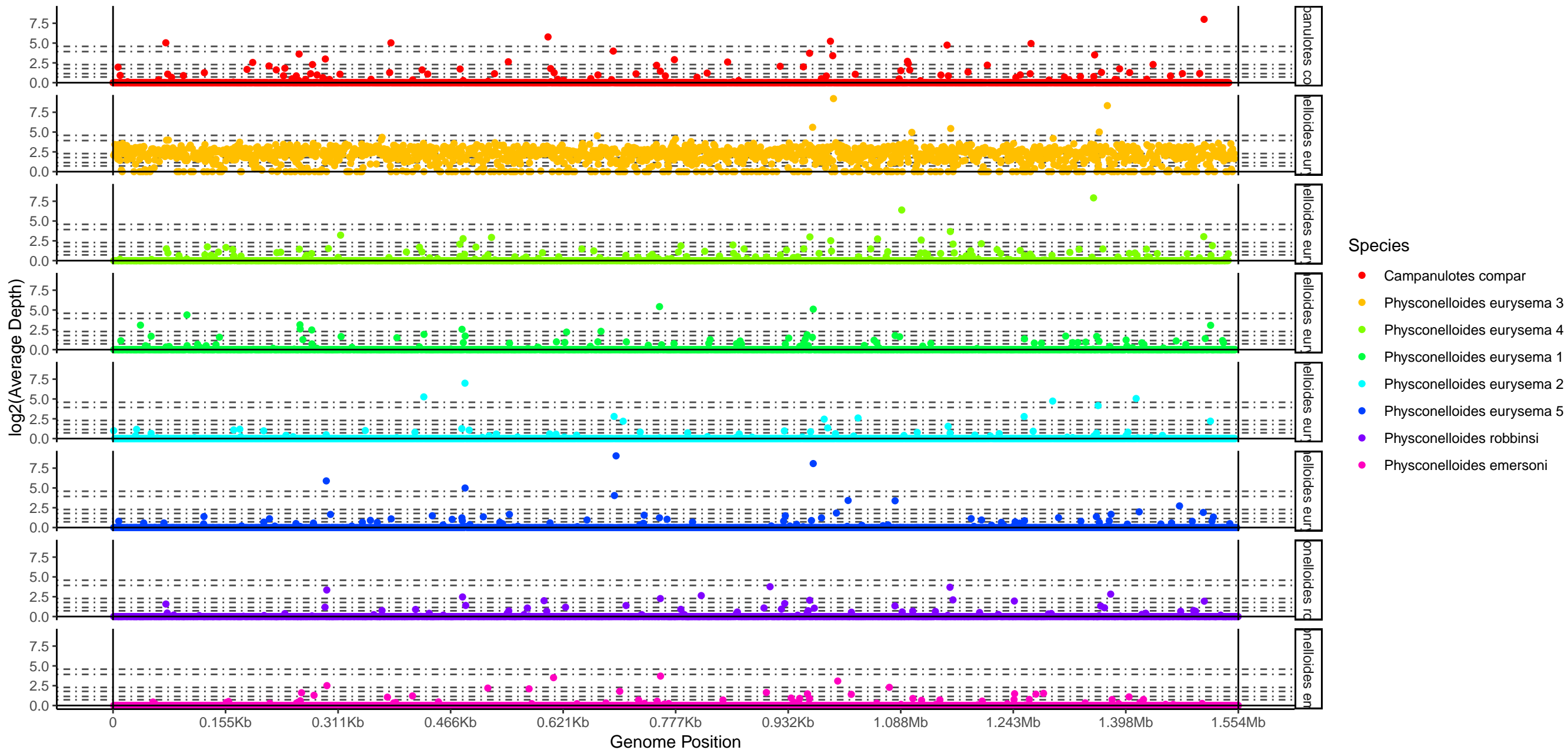
OUT Avg depth of coverage



OUT Avg depth of coverage



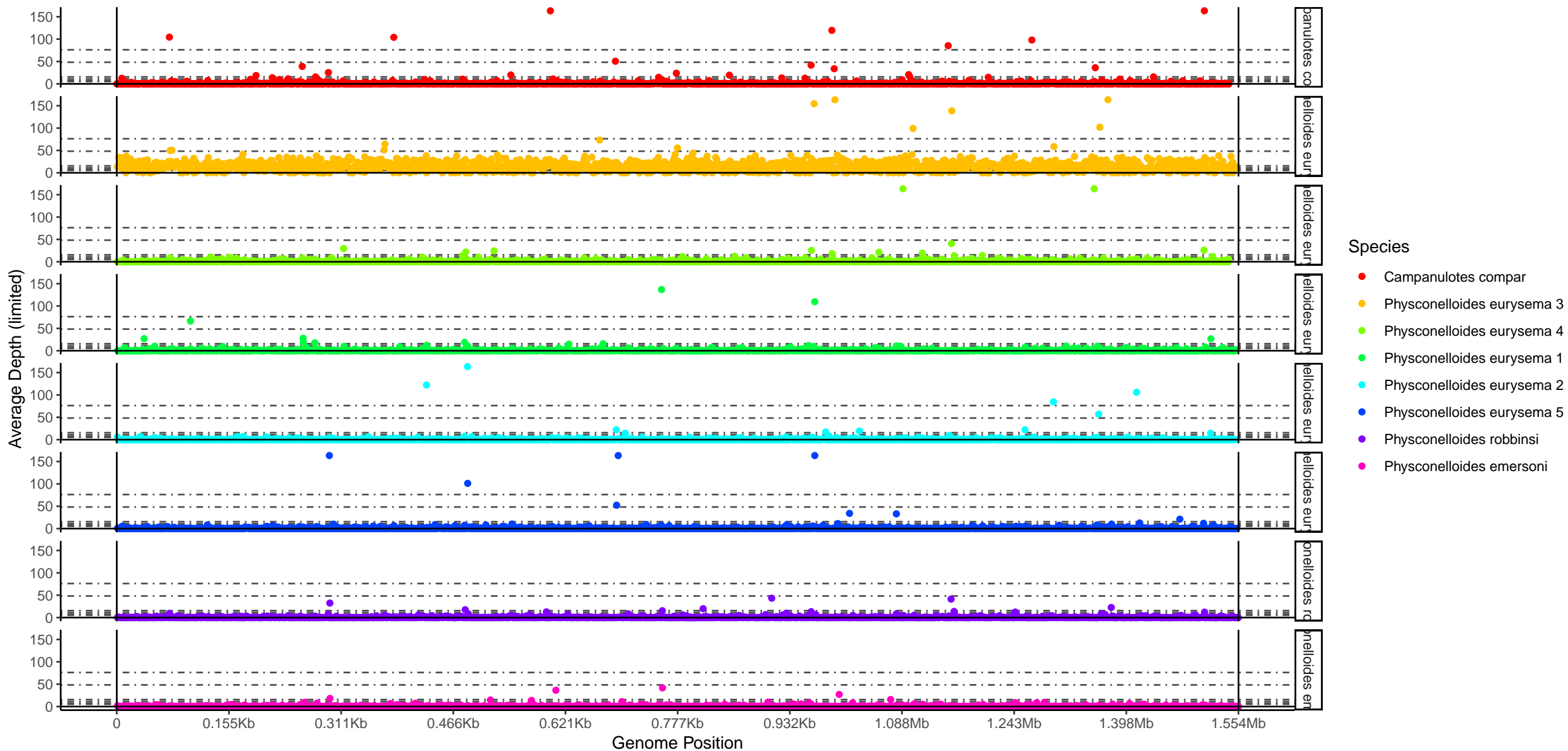
OUT Log2(Avg depth of coverage)



OUT Log2(Avg depth of coverage)



OUT Avg depth of coverage



OUT Avg depth of coverage

