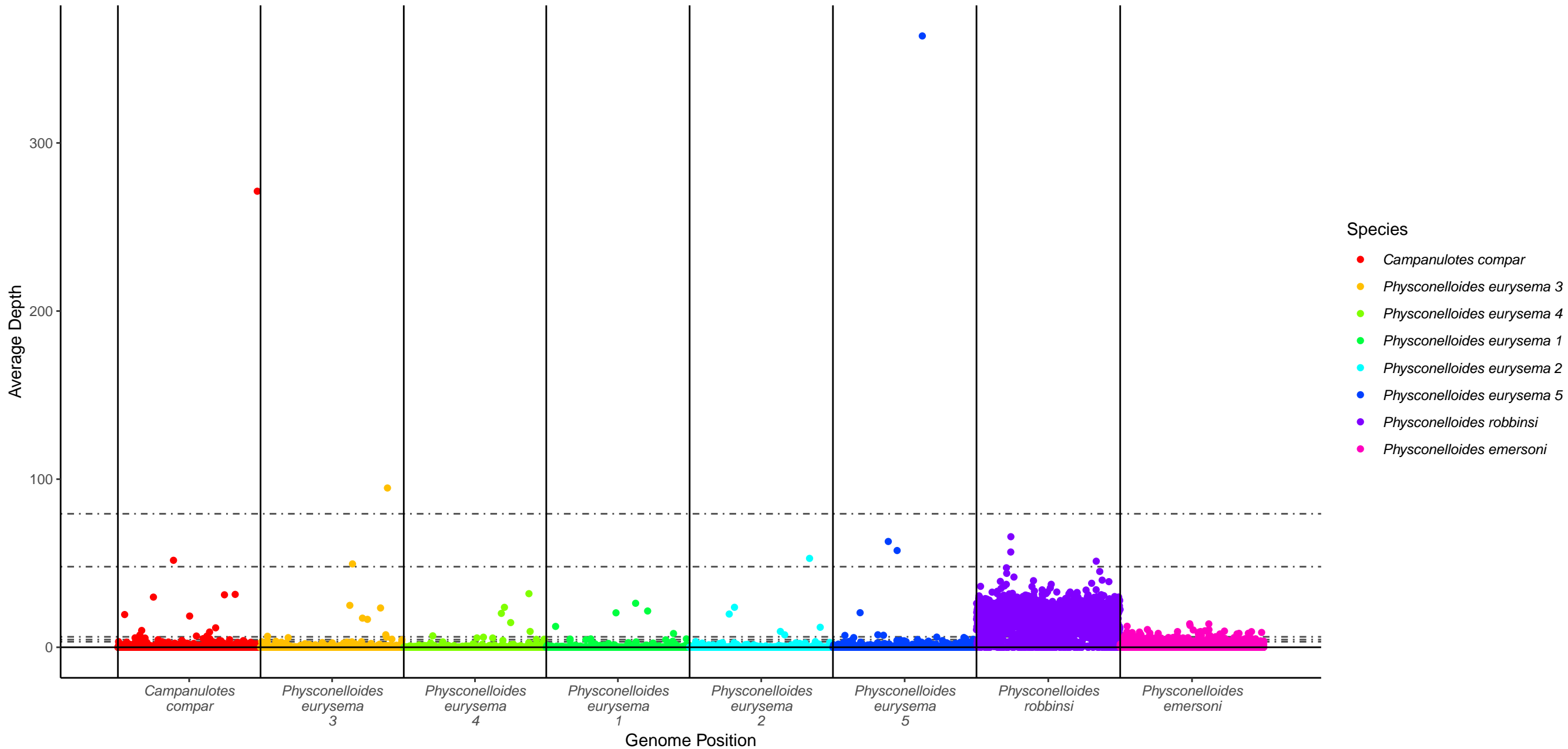
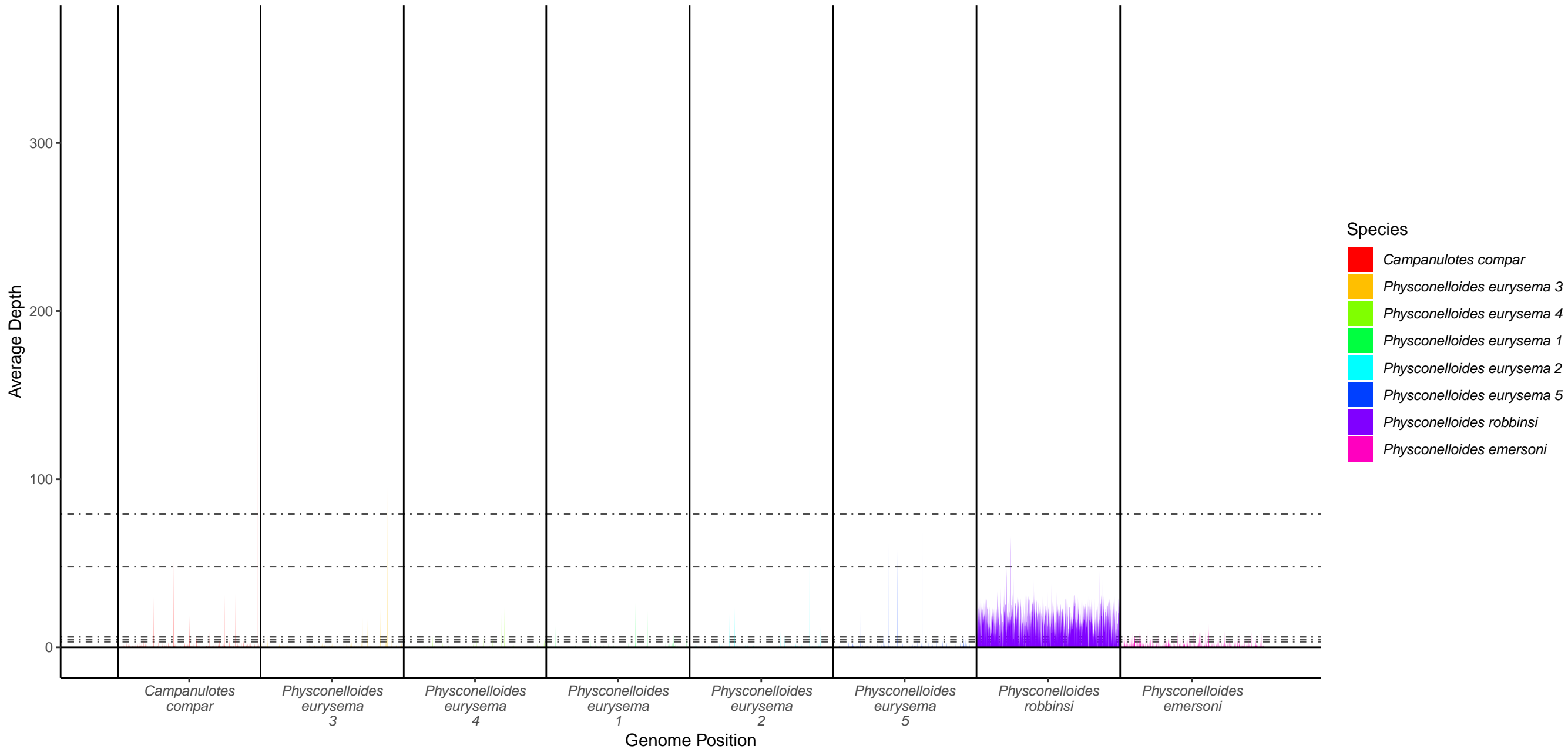


OUT Avg depth of coverage



OUT Avg depth of coverage

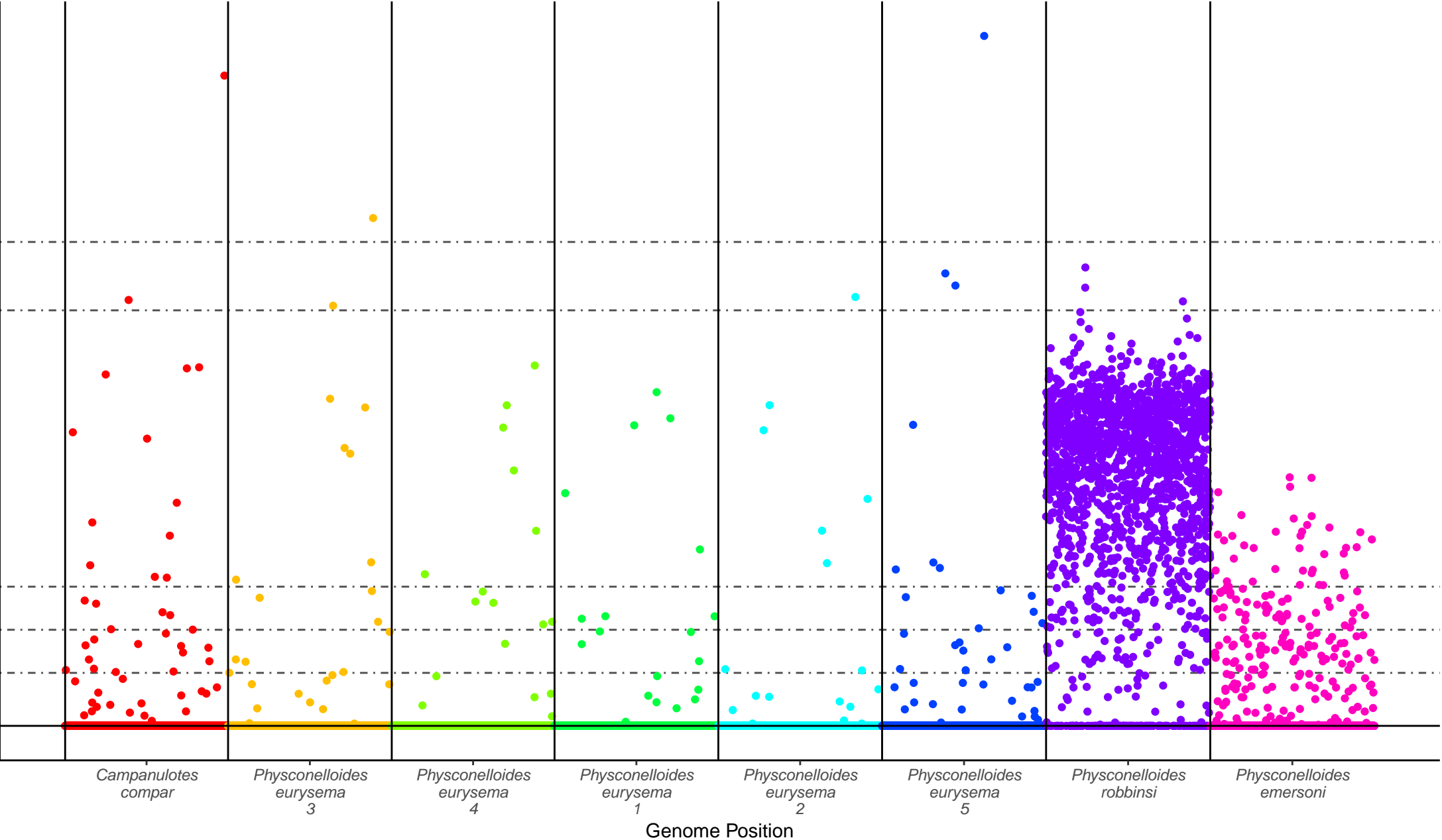


OUT Log2(Average Depth)

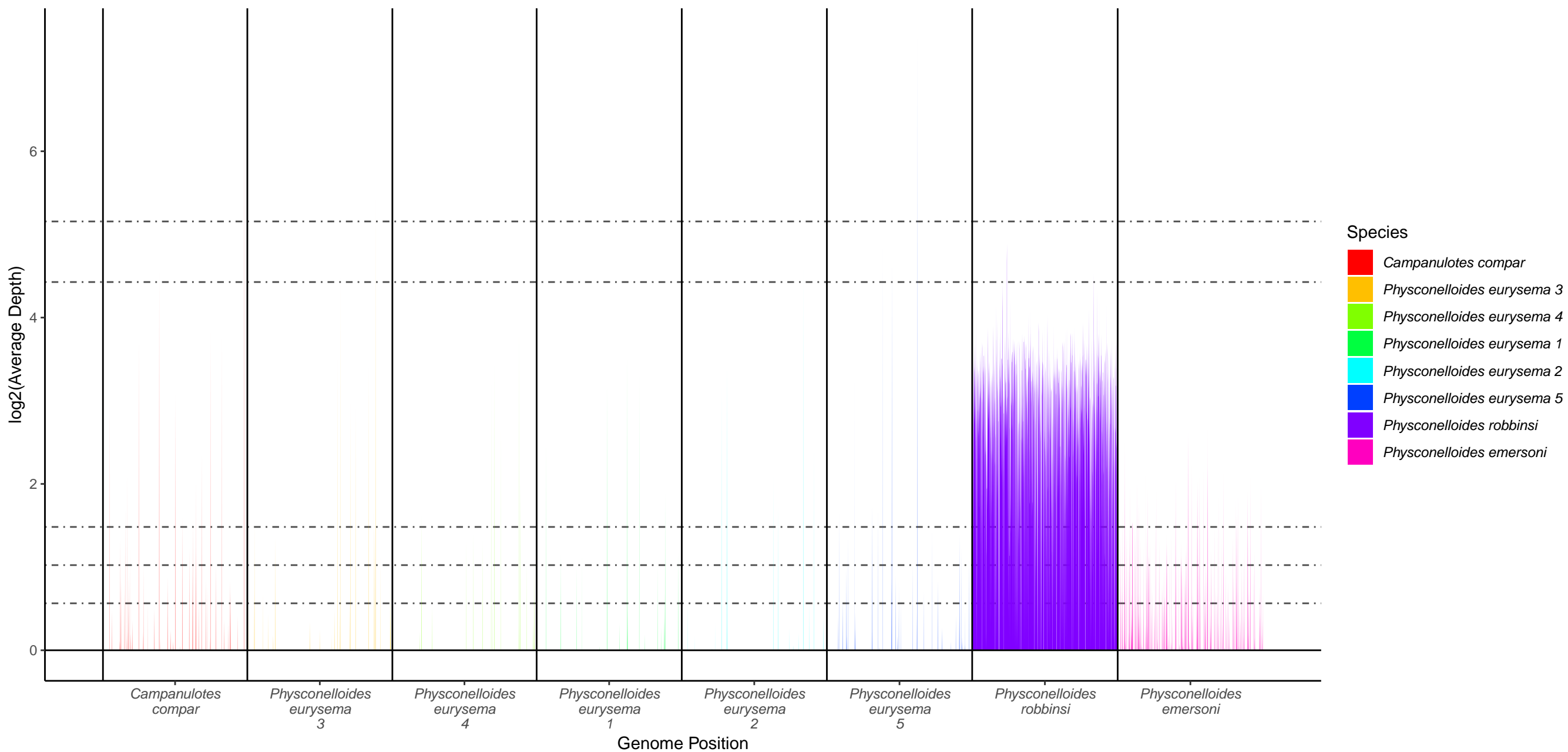
log2(Average Depth)

Species

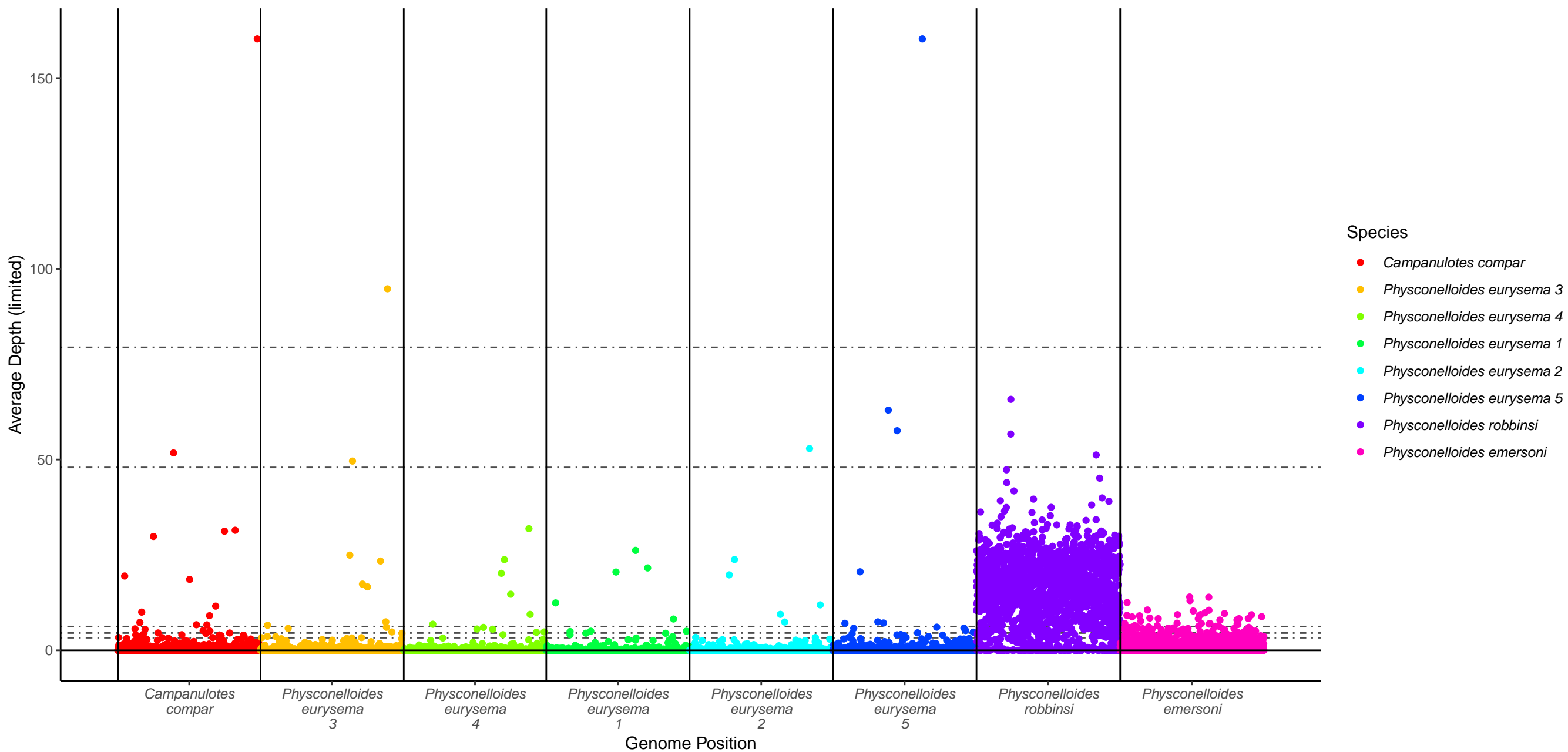
- *Campanulotes compar*
- *Physconelloides eurysema 3*
- *Physconelloides eurysema 4*
- *Physconelloides eurysema 1*
- *Physconelloides eurysema 2*
- *Physconelloides eurysema 5*
- *Physconelloides robbinsi*
- *Physconelloides emersoni*



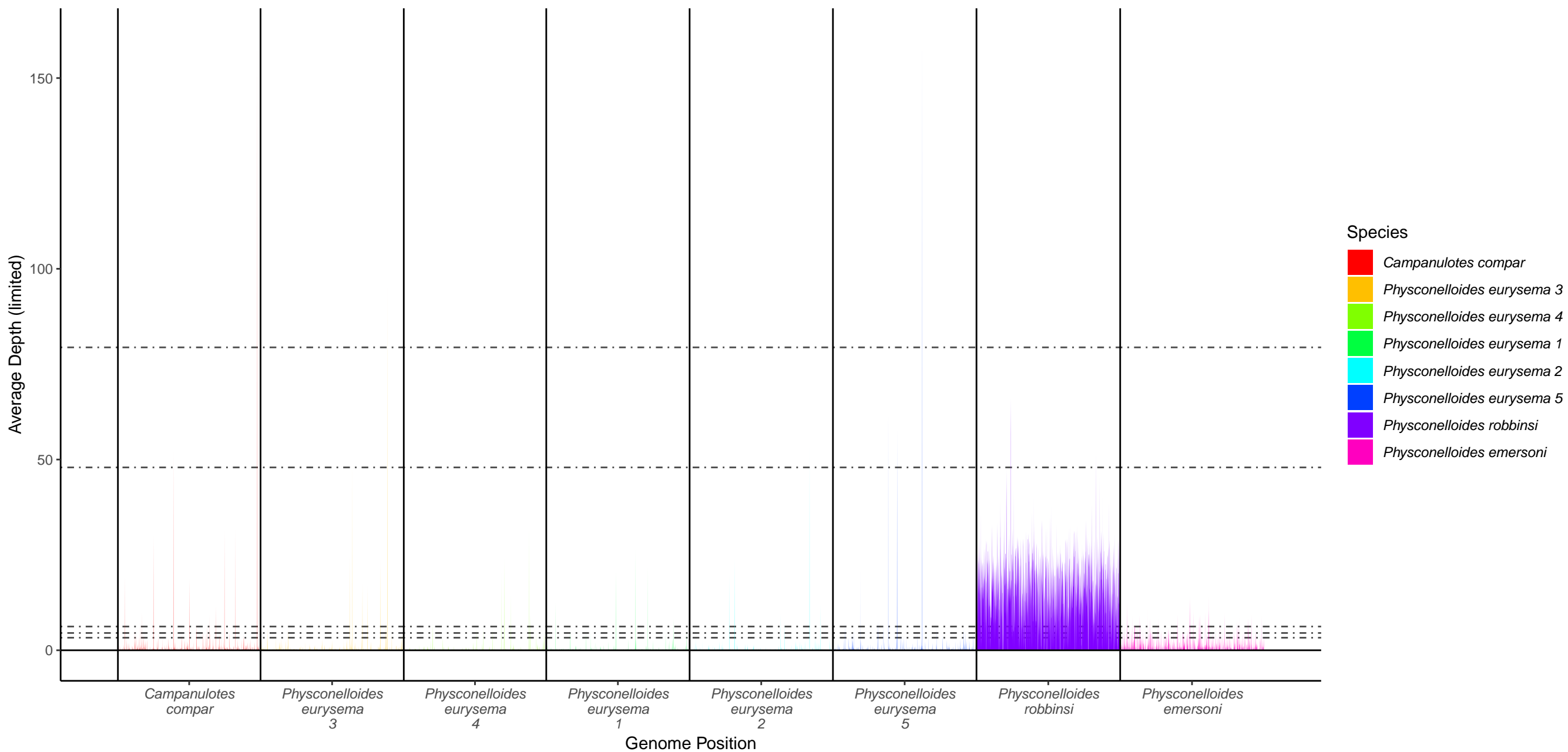
OUT Log2(Average Depth)



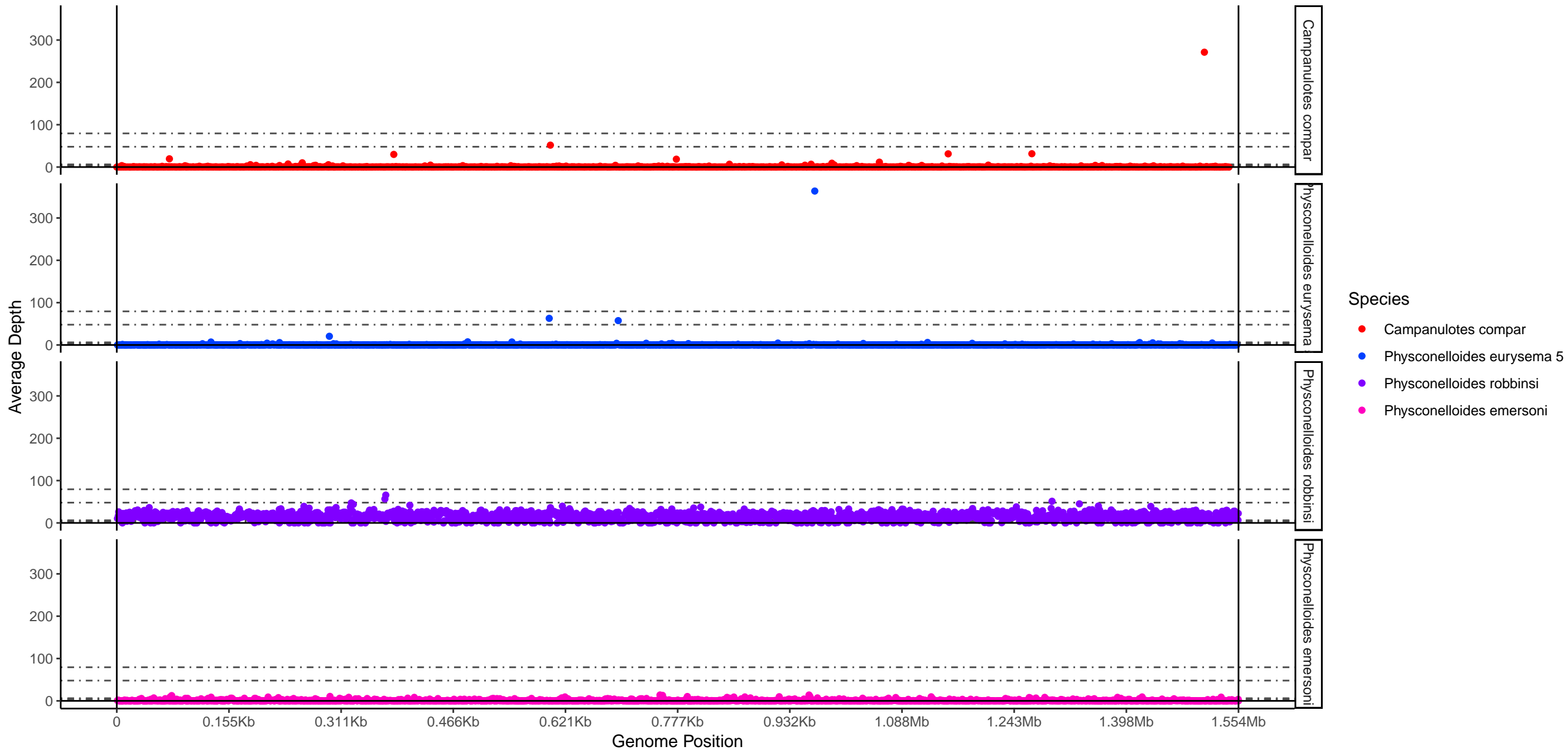
OUT Avg depth of coverage



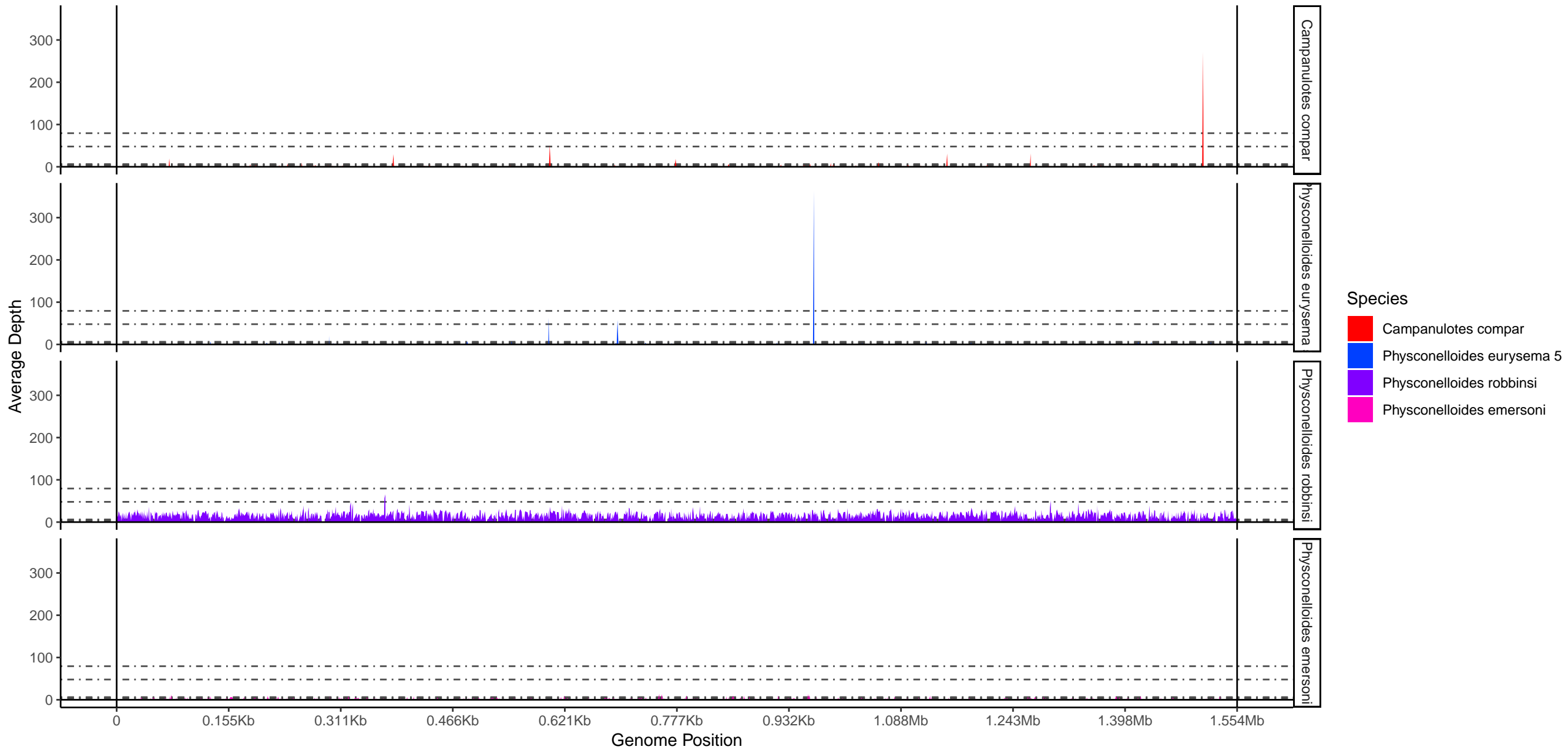
OUT Avg depth of coverage



OUT Avg depth of coverage



OUT Avg depth of coverage



OUT Log2(Avg depth of coverage)

log2(Average Depth)

0 0.155Kb 0.311Kb 0.466Kb 0.621Kb 0.777Kb 0.932Kb 1.088Mb 1.243Mb 1.398Mb 1.554Mb

Genome Position

Campanulotes compar

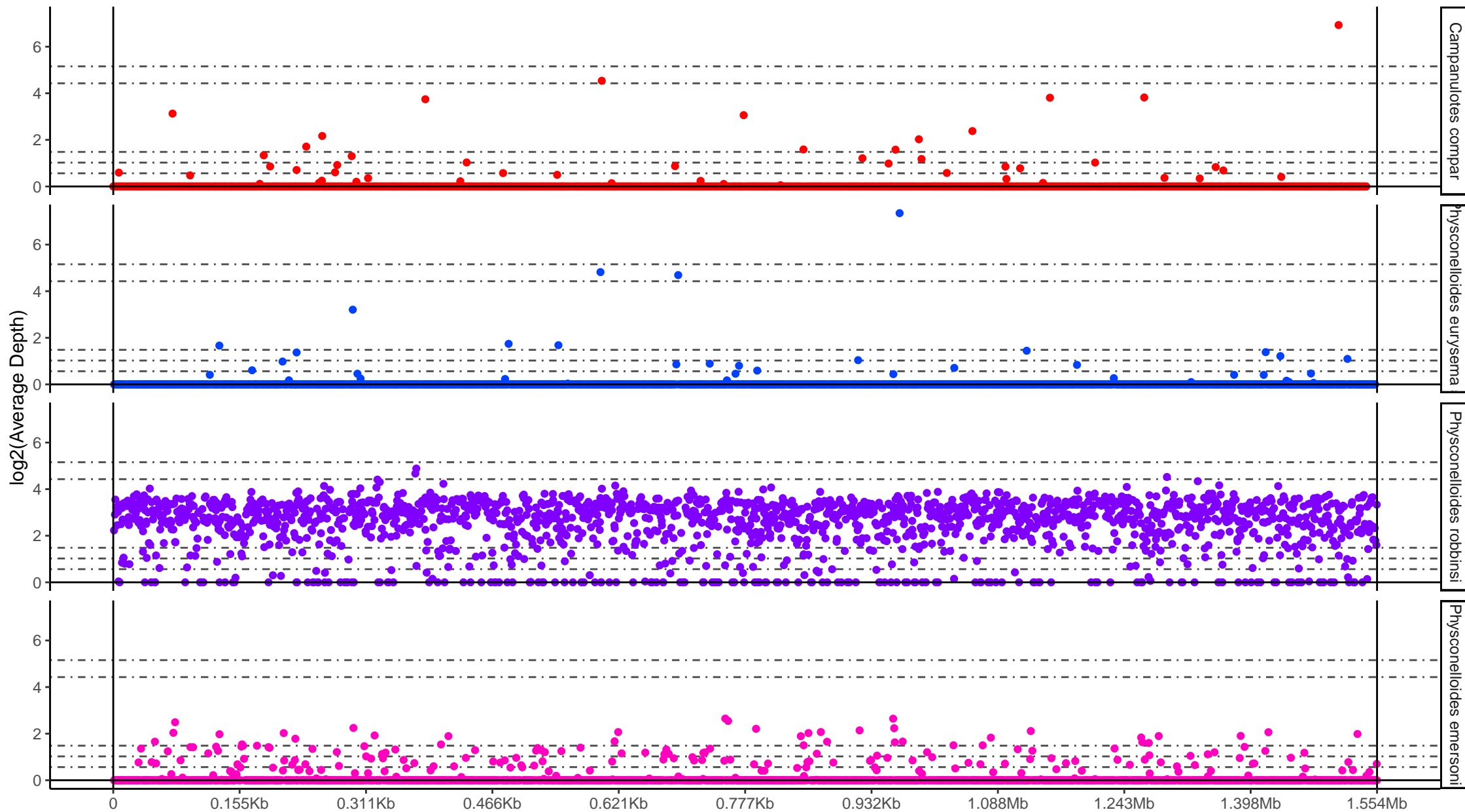
Physconelloides eurysema

Physconelloides robbinsi

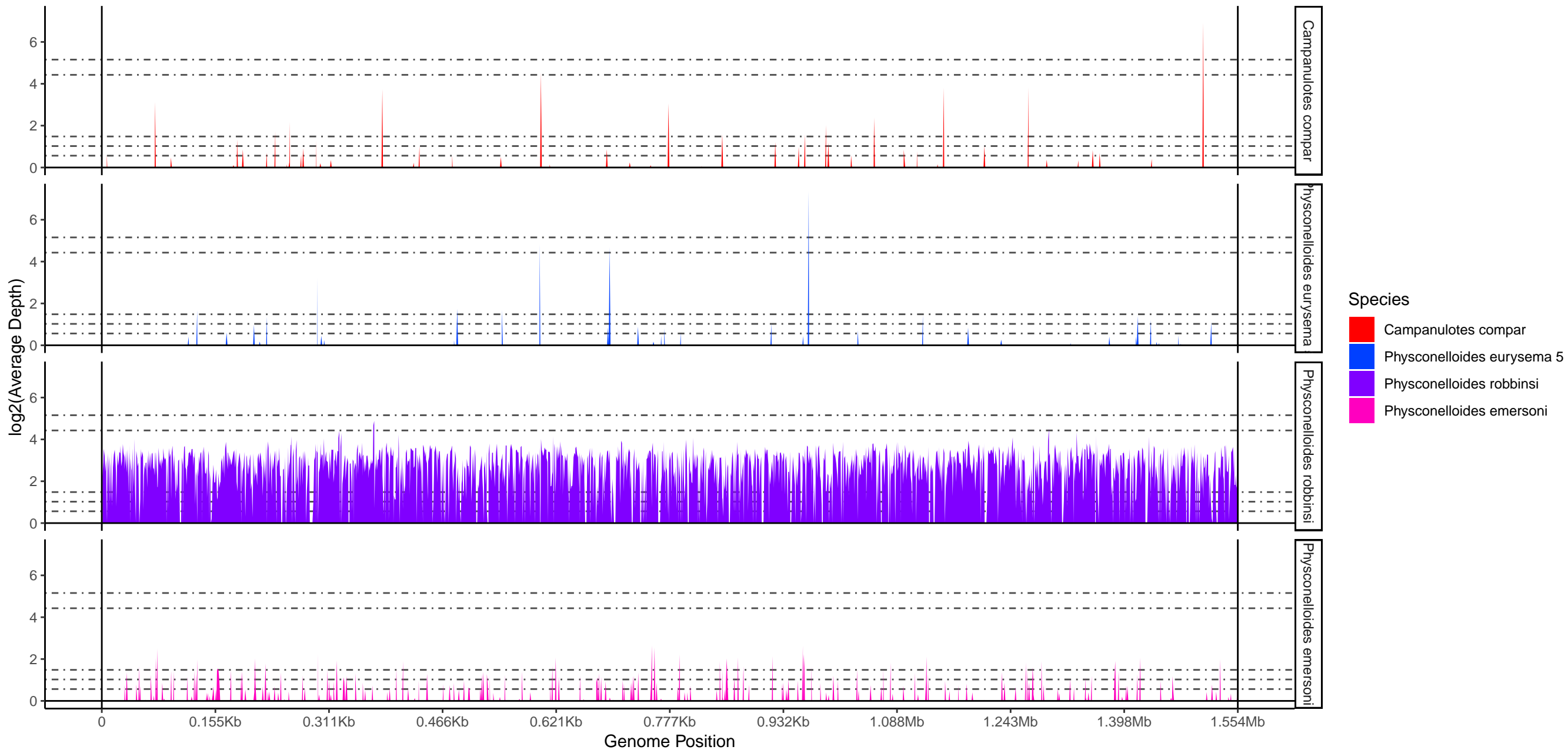
Physconelloides emersoni

Species

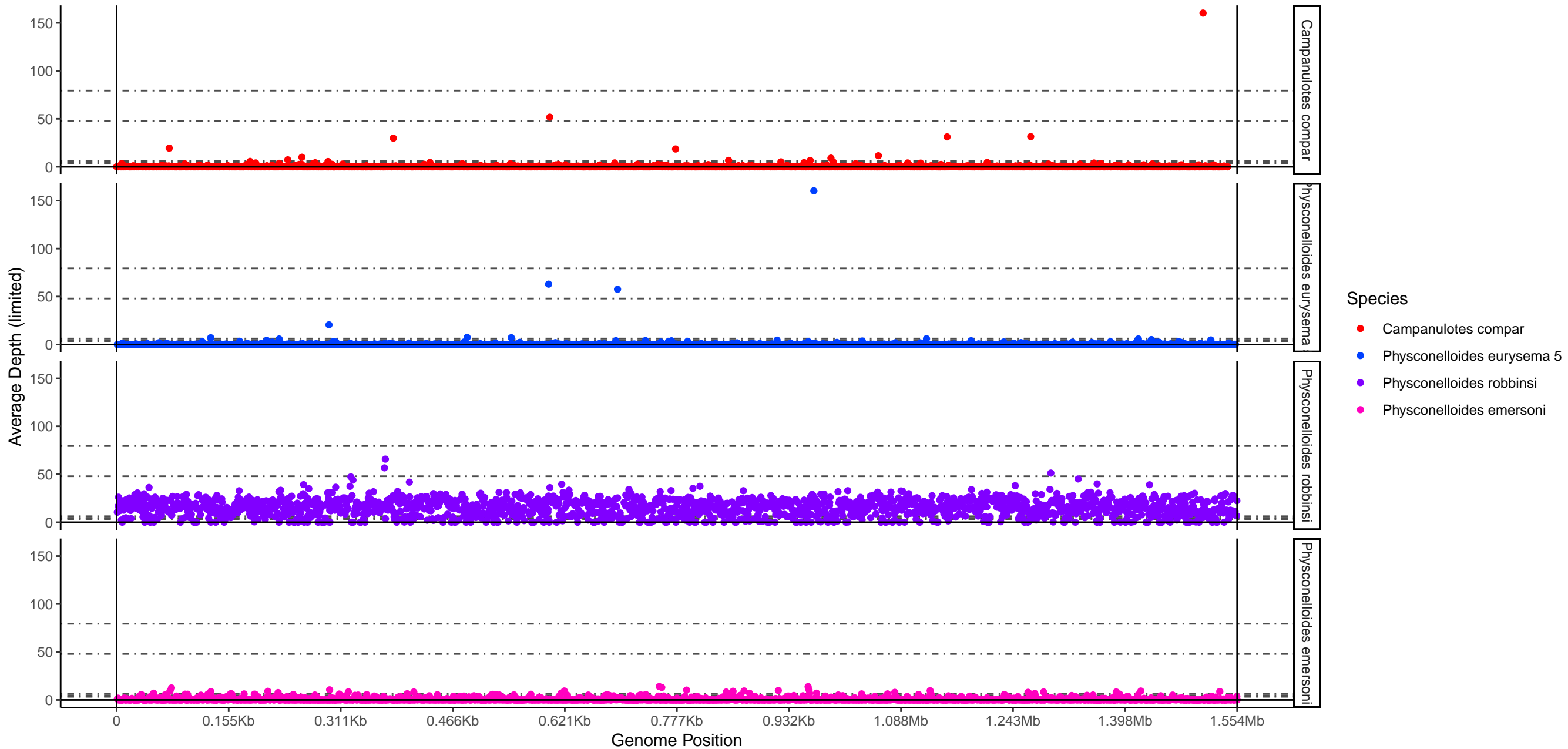
- Campanulotes compar
- Physconelloides eurysema 5
- Physconelloides robbinsi
- Physconelloides emersoni



OUT Log2(Avg depth of coverage)



OUT Avg depth of coverage



OUT Avg depth of coverage

