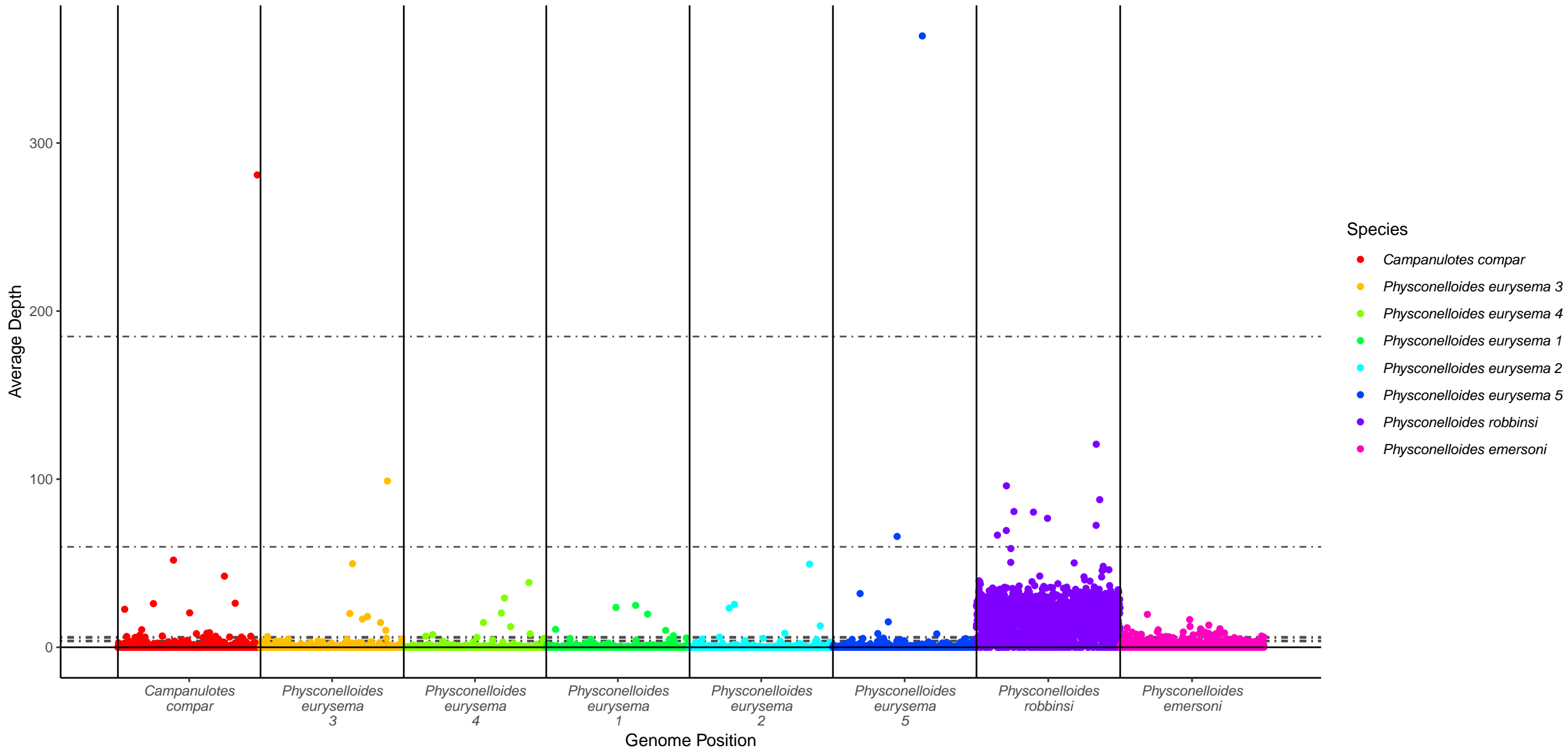
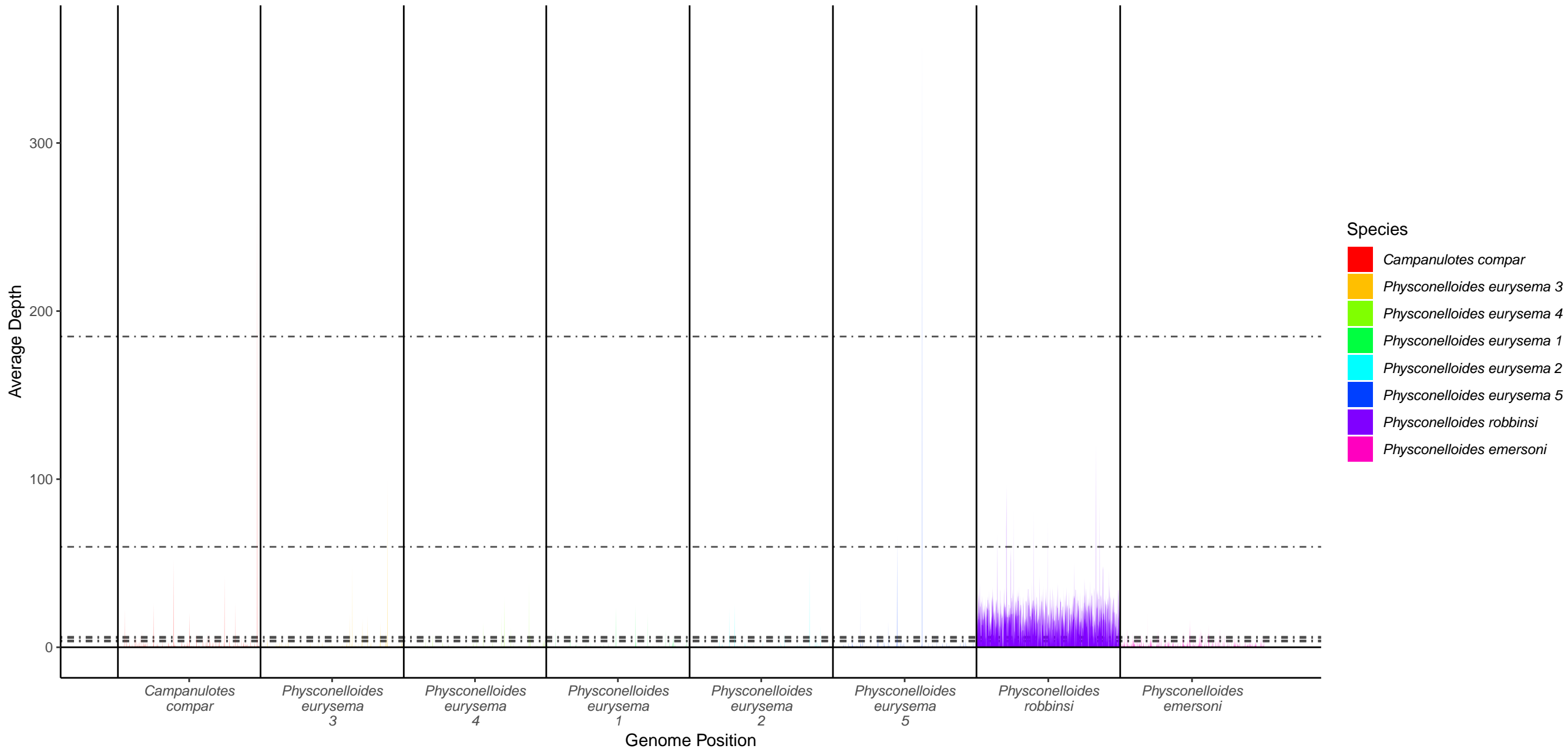


OUT Avg depth of coverage



OUT Avg depth of coverage

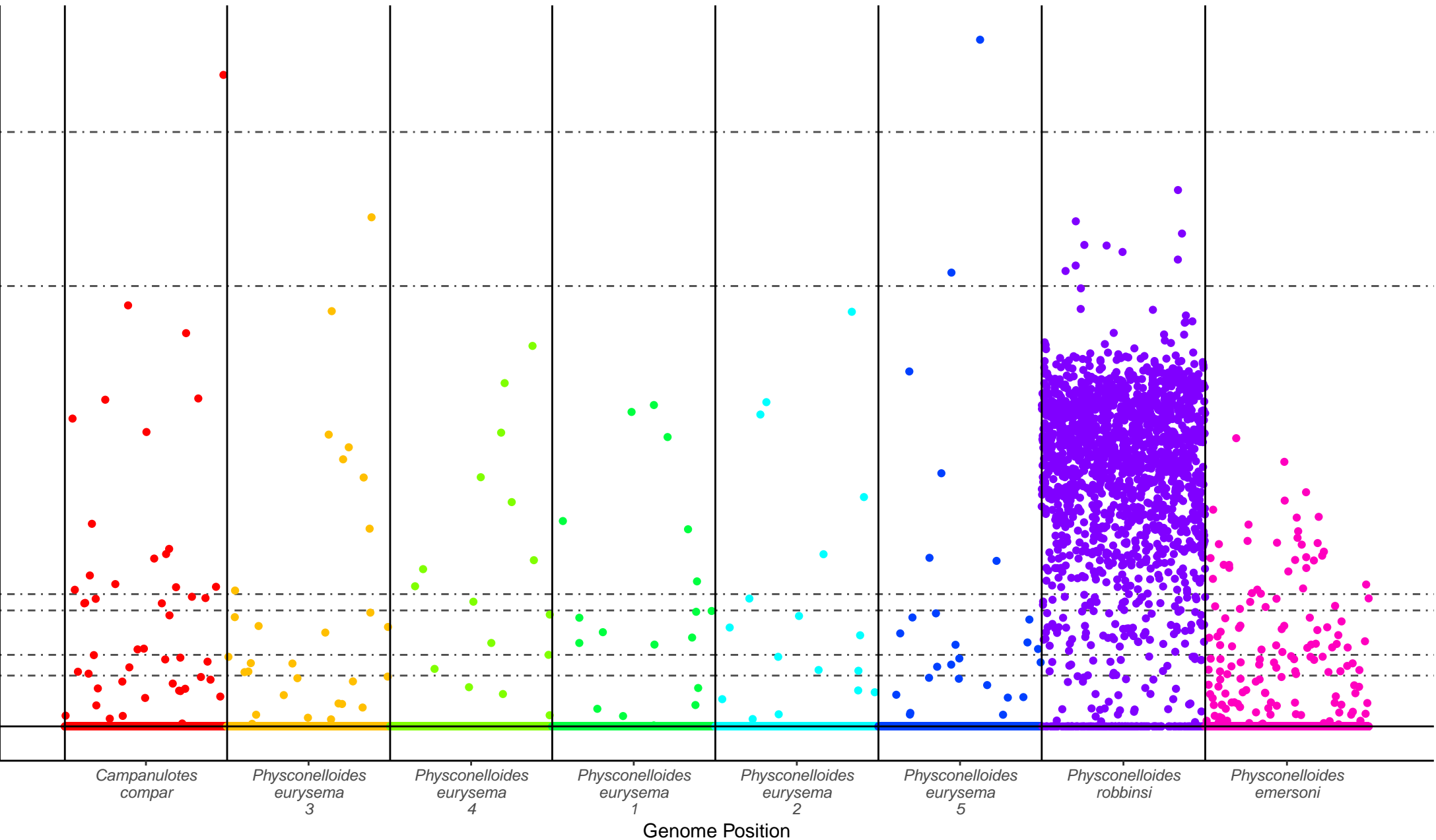


OUT Log2(Average Depth)

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*

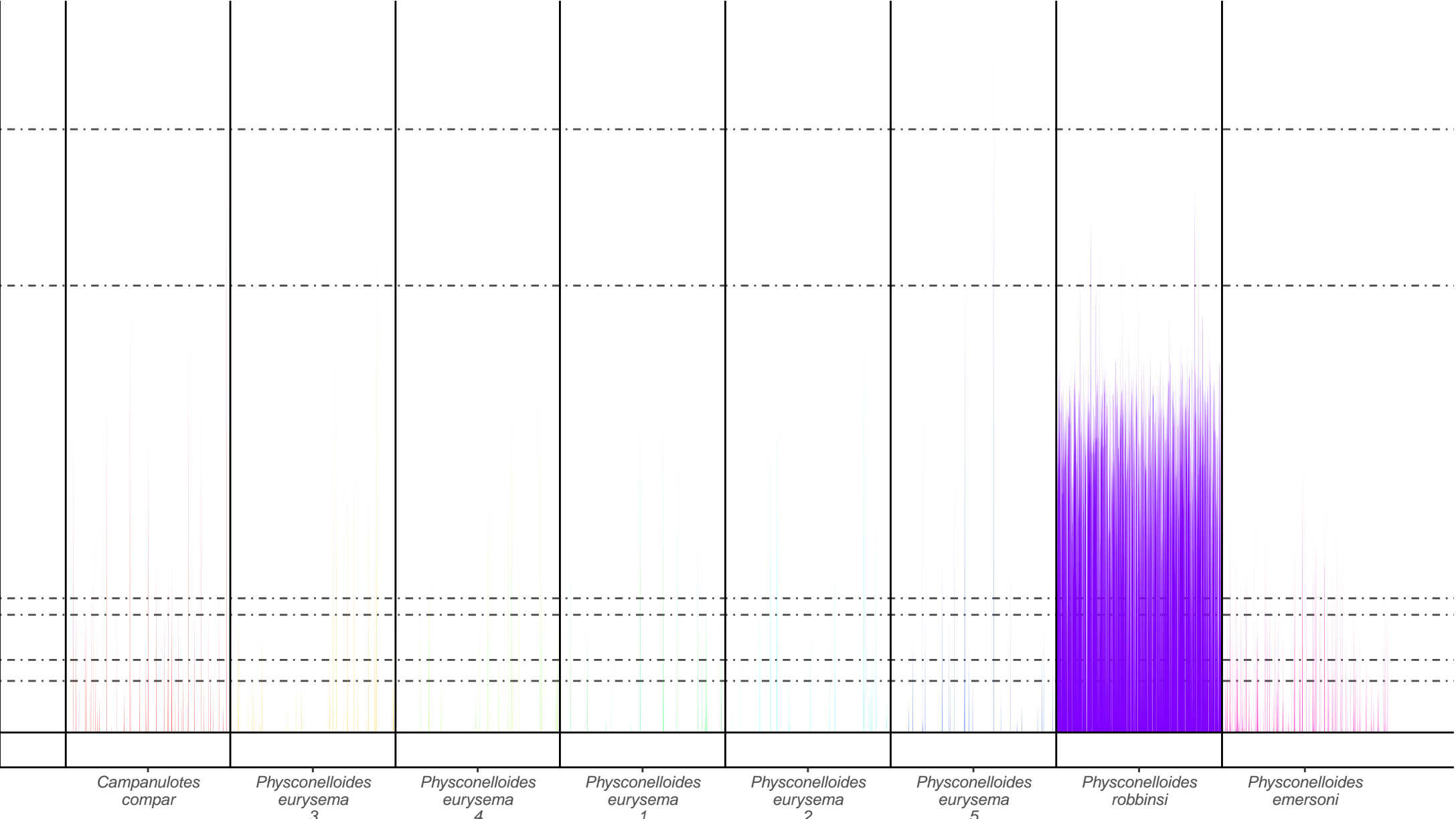


OUT Log2(Average Depth)

log2(Average Depth)

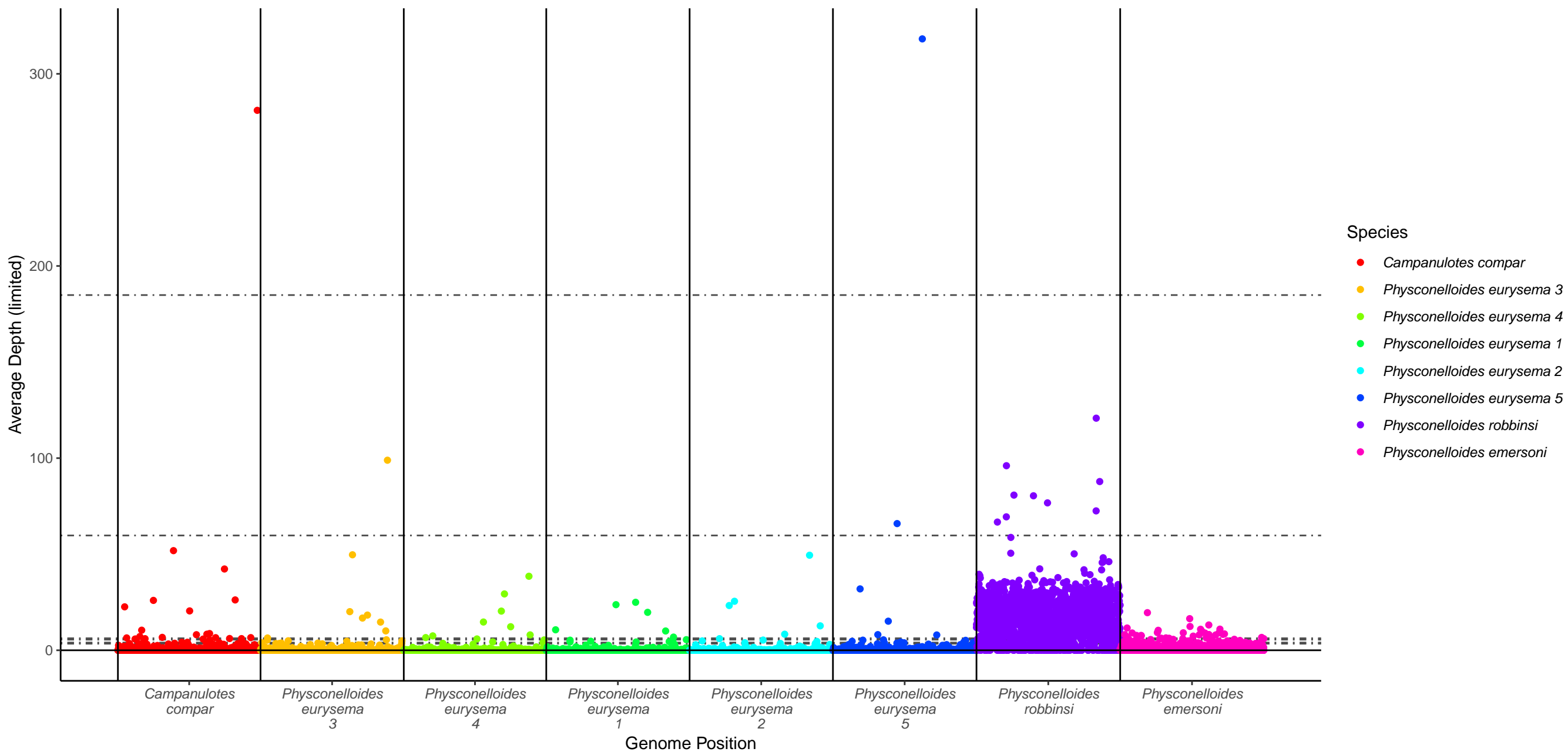
Species

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

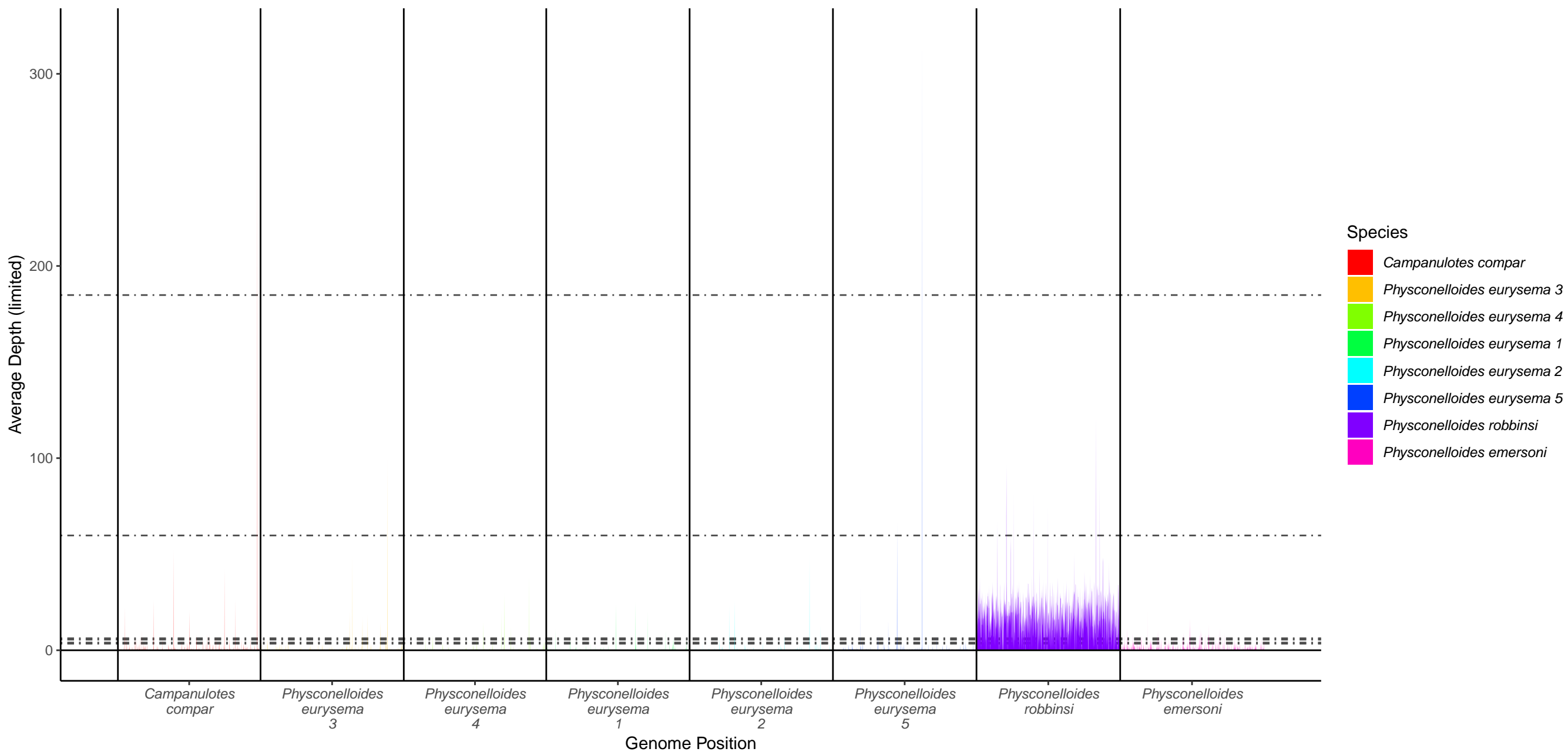


Genome Position

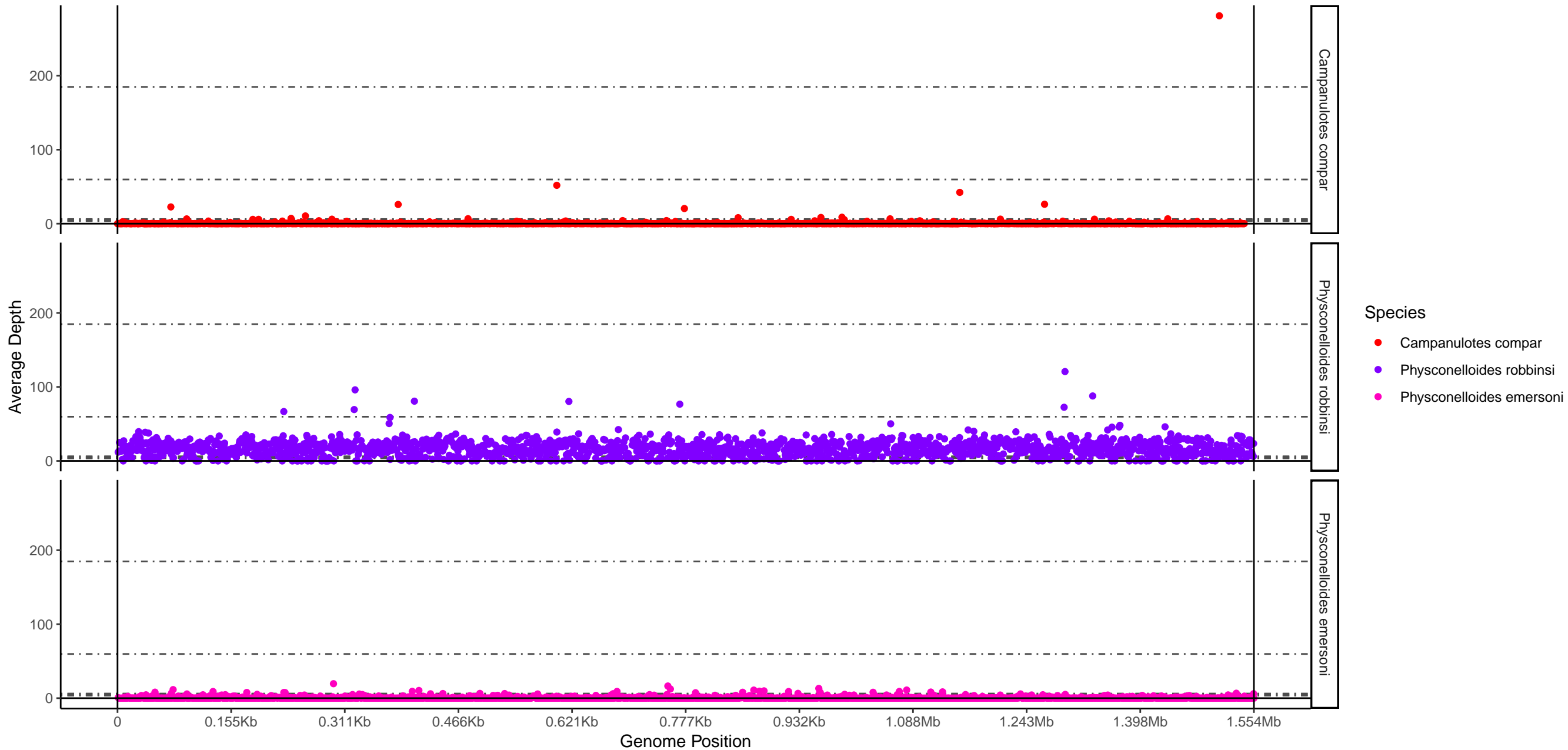
OUT Avg depth of coverage



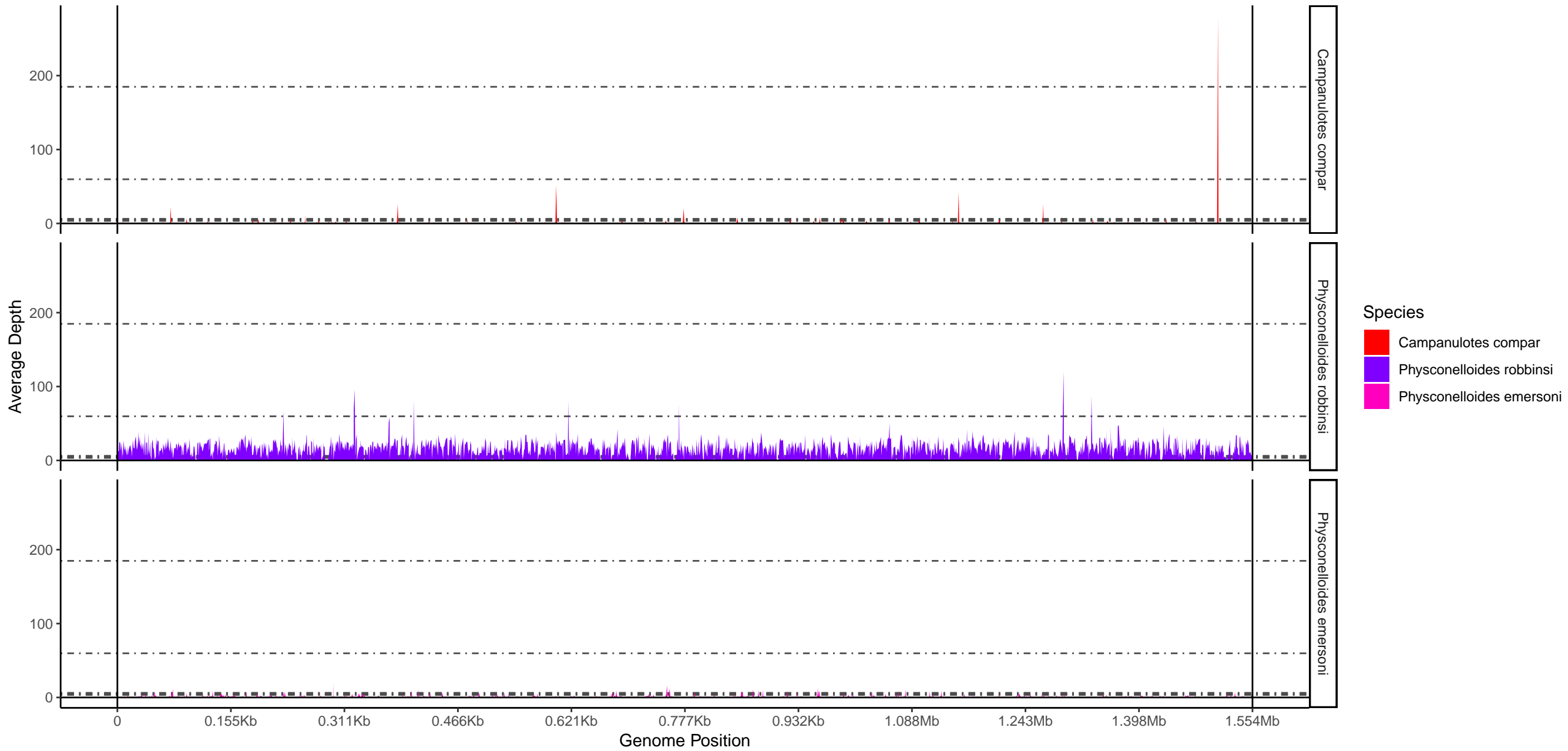
OUT Avg depth of coverage



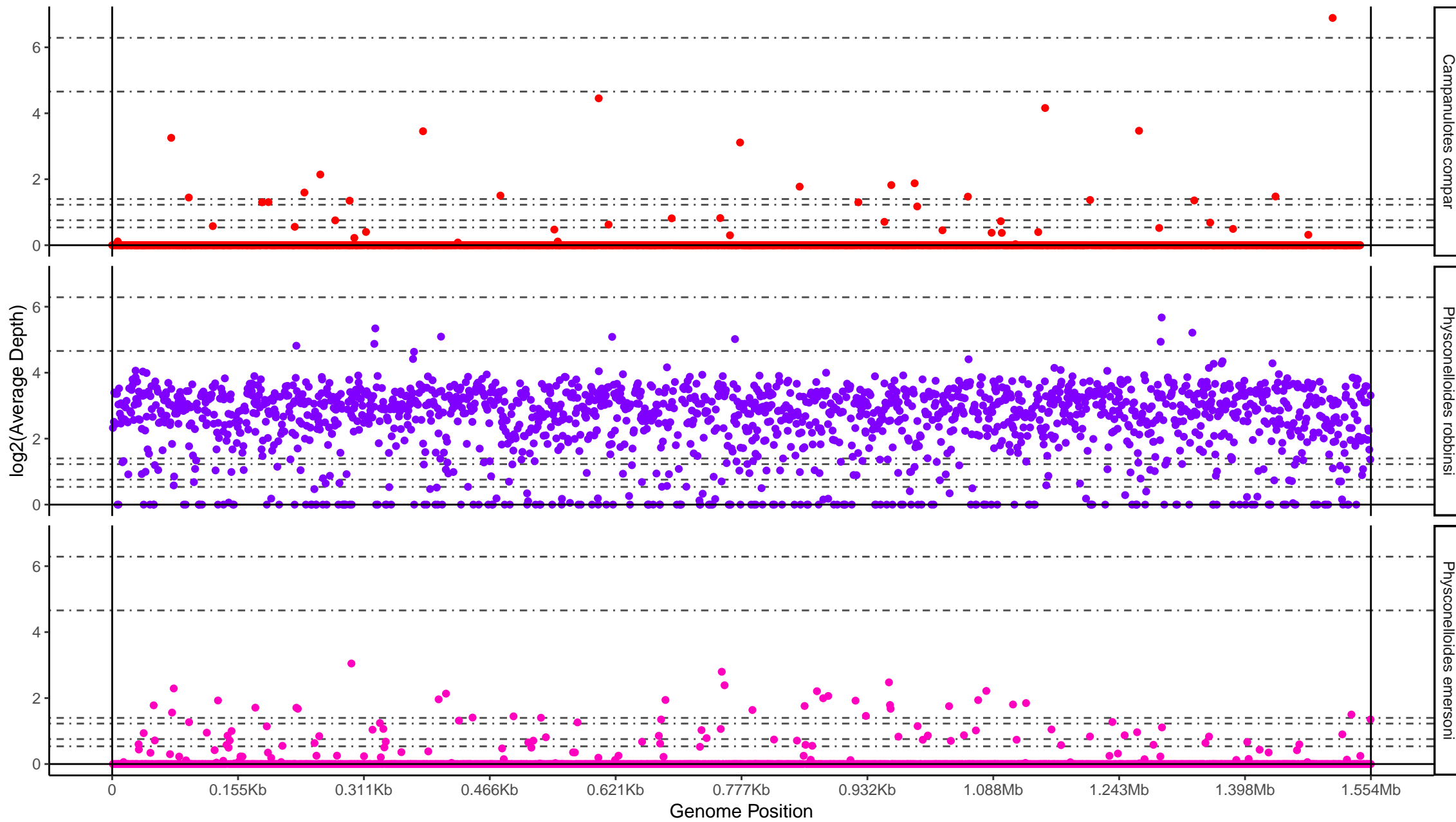
OUT Avg depth of coverage



OUT Avg depth of coverage



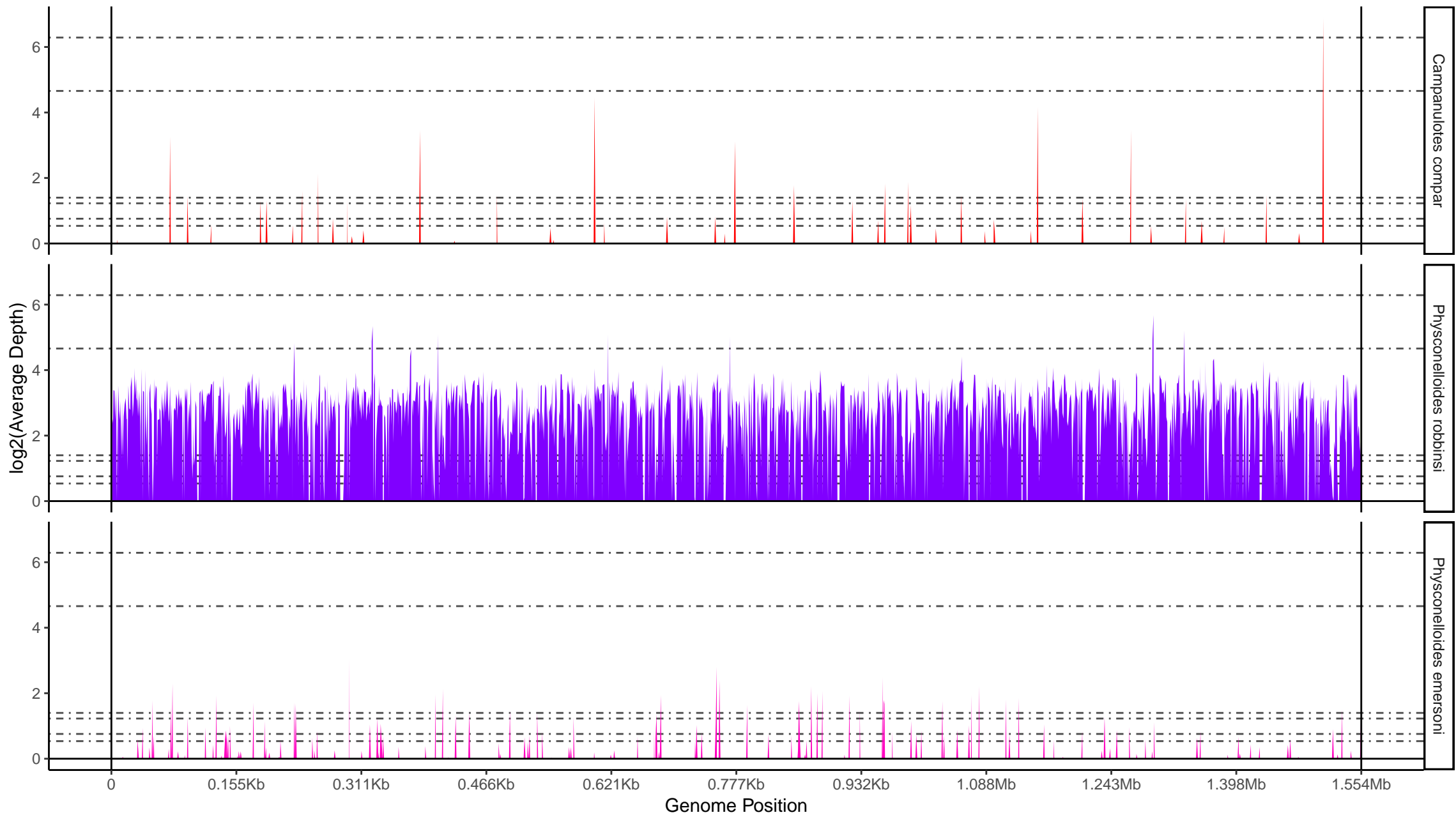
OUT Log2(Avg depth of coverage)



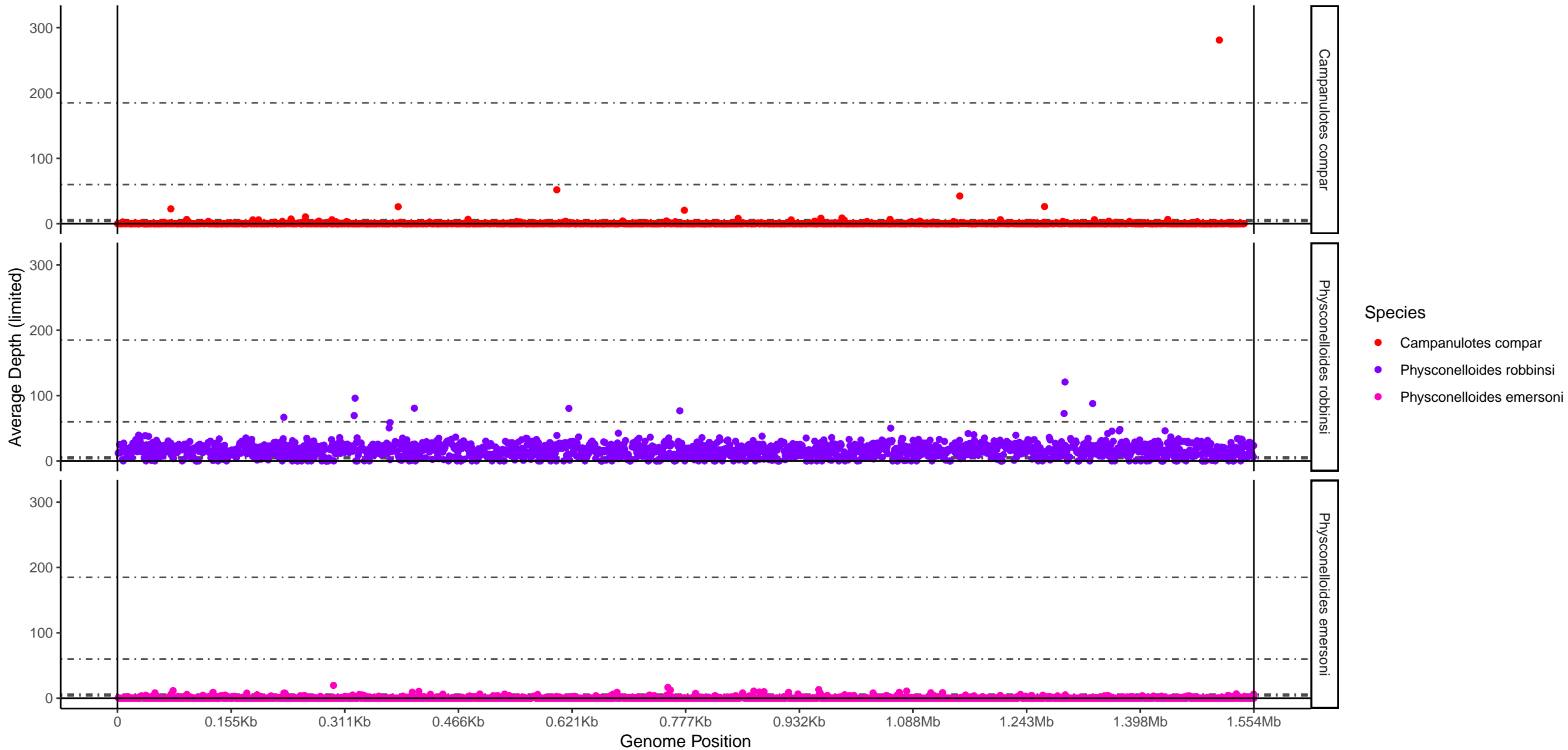
Species

- Campanulotes compar
- Physconelloides robbinsi
- Physconelloides emersoni

OUT Log2(Avg depth of coverage)



OUT Avg depth of coverage



OUT Avg depth of coverage

