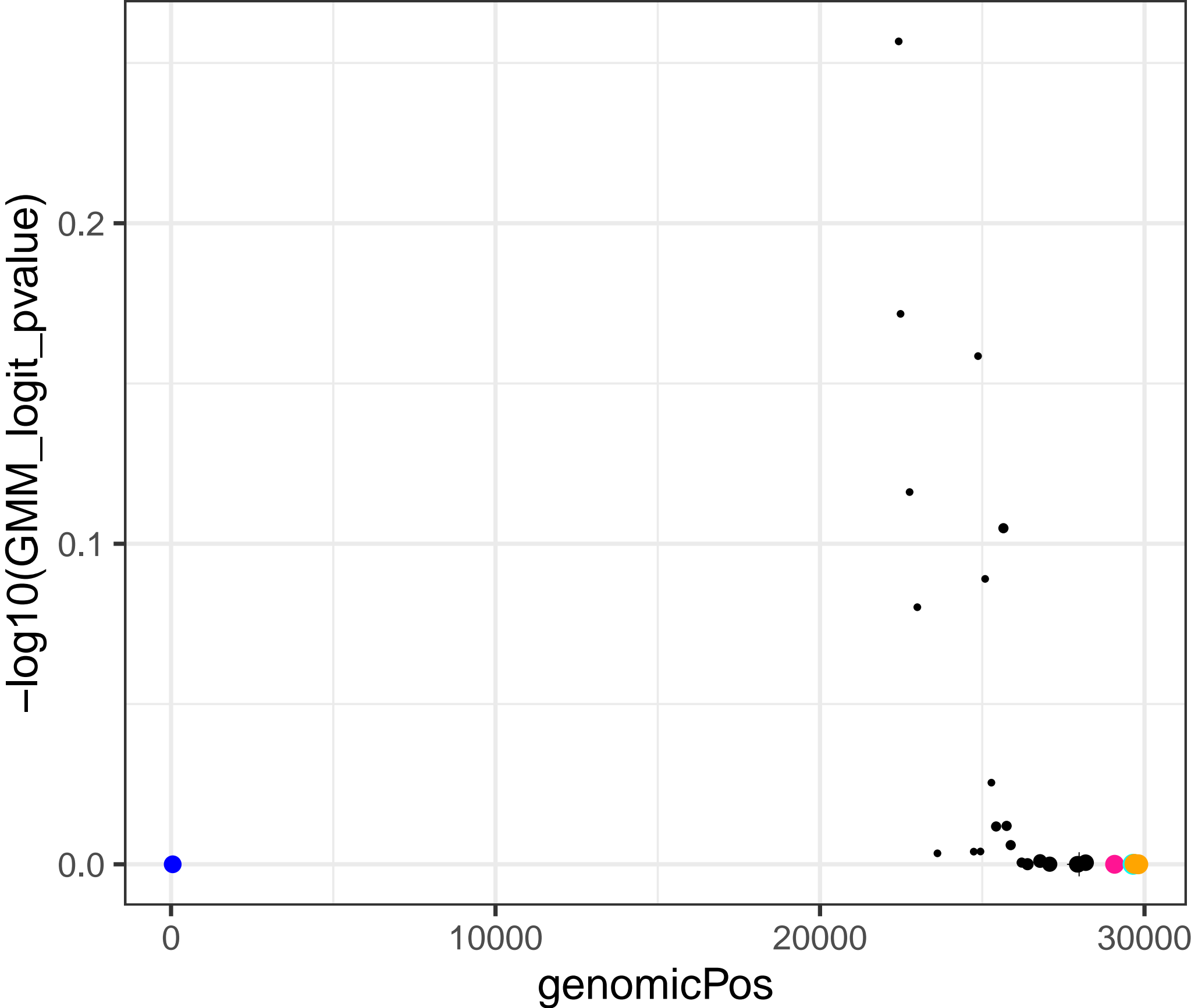


UGUAR occurrence over all transcripts



- fragment_ID
- No_Fragment
 - Fragment1_5UTR
 - Fragment7_8
 - Fragment9_10
 - Fragment10_3UTR

- IVT
- IVT junction
 - No junction

- N° of transcripts
- 1
 - 2
 - 3
 - 5
 - 8
 - 10
 - 14
 - 17
 - 19
 - 20