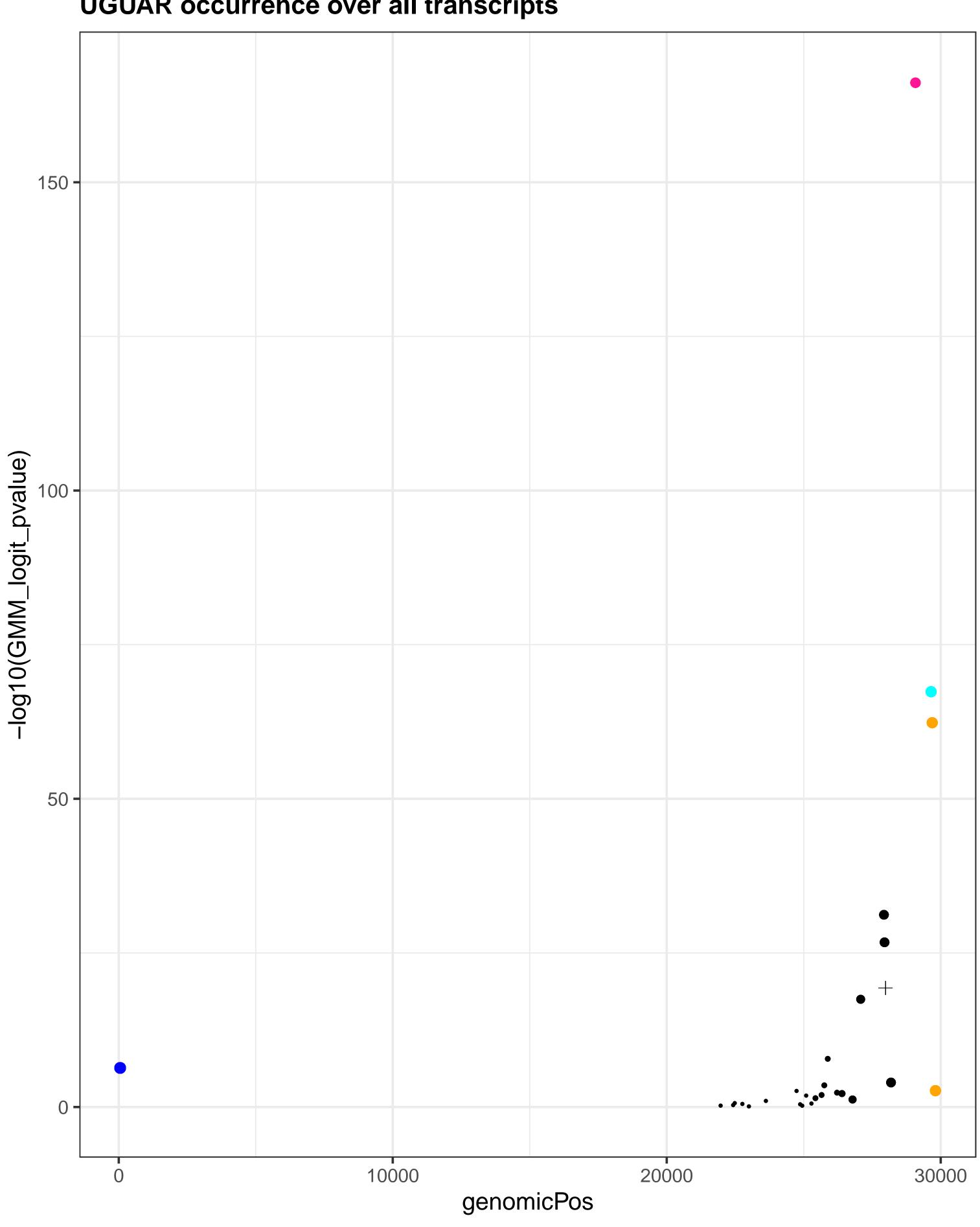
## **UGUAR** occurrence over all transcripts



## fragment\_ID

- No\_Fragment
  Fragment1\_5UTR
  Fragment7\_8
  Fragment9\_10
  Fragment10\_3UTR

## IVT

- IVT junctionNo junction

## N° of transcripts

- 2 3 5 8 10 18 20 14