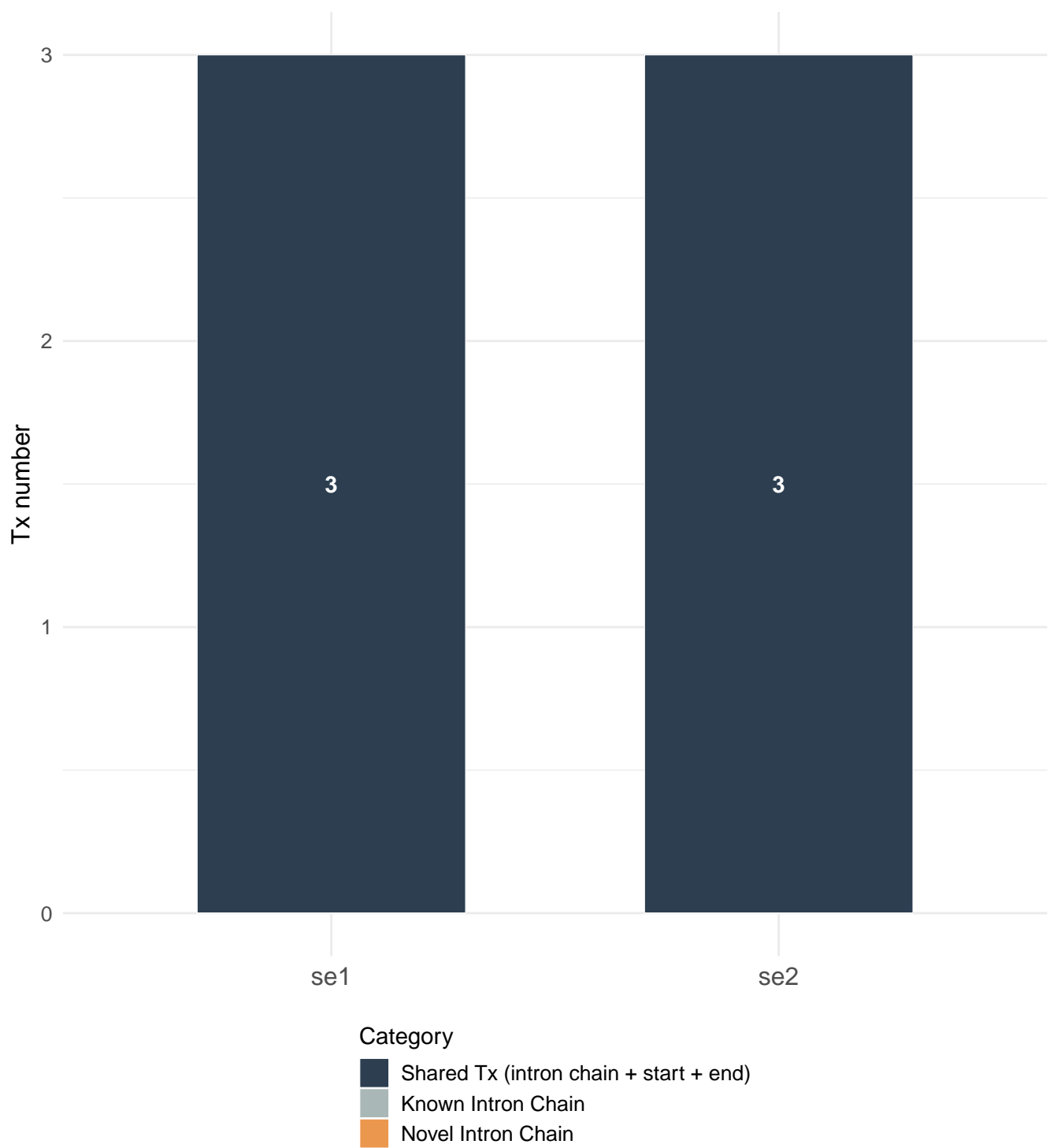
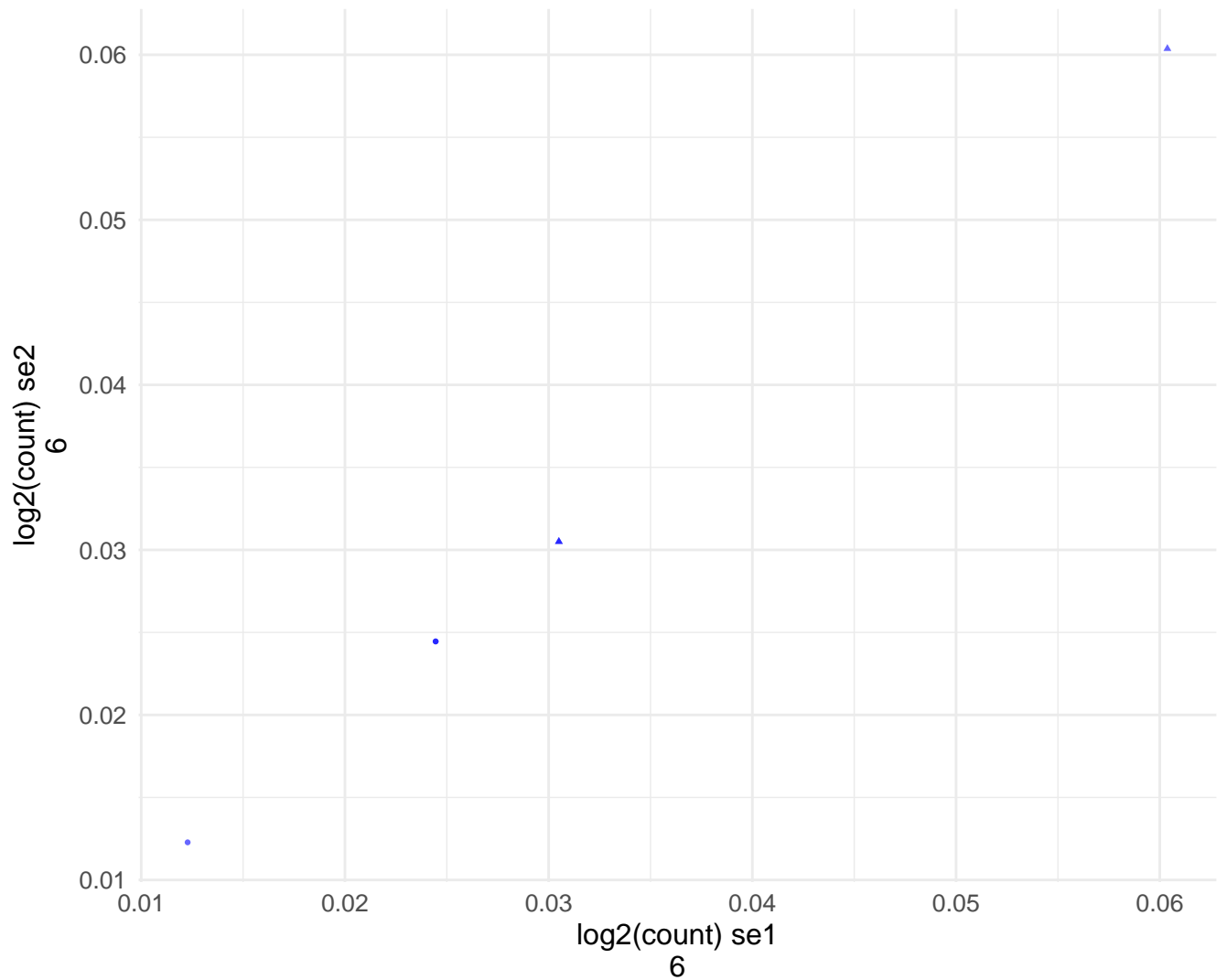


# Transcript Discovery Comparison



# Transcript Quantification Comparison

Sample: GIS\_cellMix\_HepG2-A549-H9-HEYA8\_5primeSingleCellcDNA\_Rep1\_Run



## shape

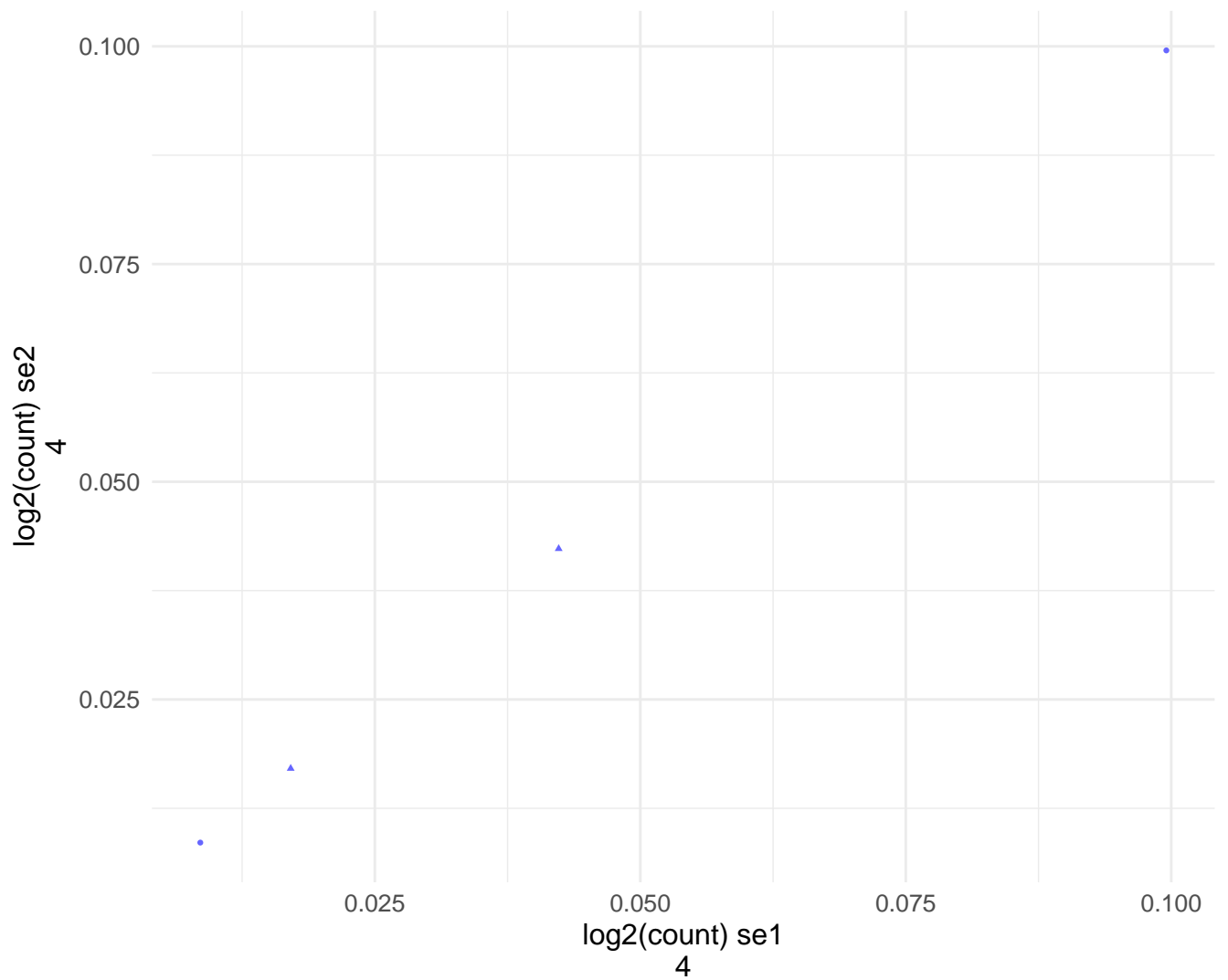
- anno\_TX 3 (50%)  
Equal: 6, Diff: 0 | x\_unique: 0, y\_unique: 0
- ▲ novel\_TX 3 (50%)  
Equal: 6, Diff: 0 | x\_unique: 0, y\_unique: 0

## color

- identical: 6 (100%)

# Transcript Quantification Comparison

Sample: GIS\_cellMix\_HepG2-A549-H9-HEYA8\_5primeSingCellcDNAPacBio\_Rep



color

• identical: 4 (100%)

shape

- anno\_TX 2 (50%)  
Equal: 4, Diff: 0 | x\_unique: 0, y\_unique: 0
- ▲ novel\_TX 2 (50%)  
Equal: 4, Diff: 0 | x\_unique: 0, y\_unique: 0