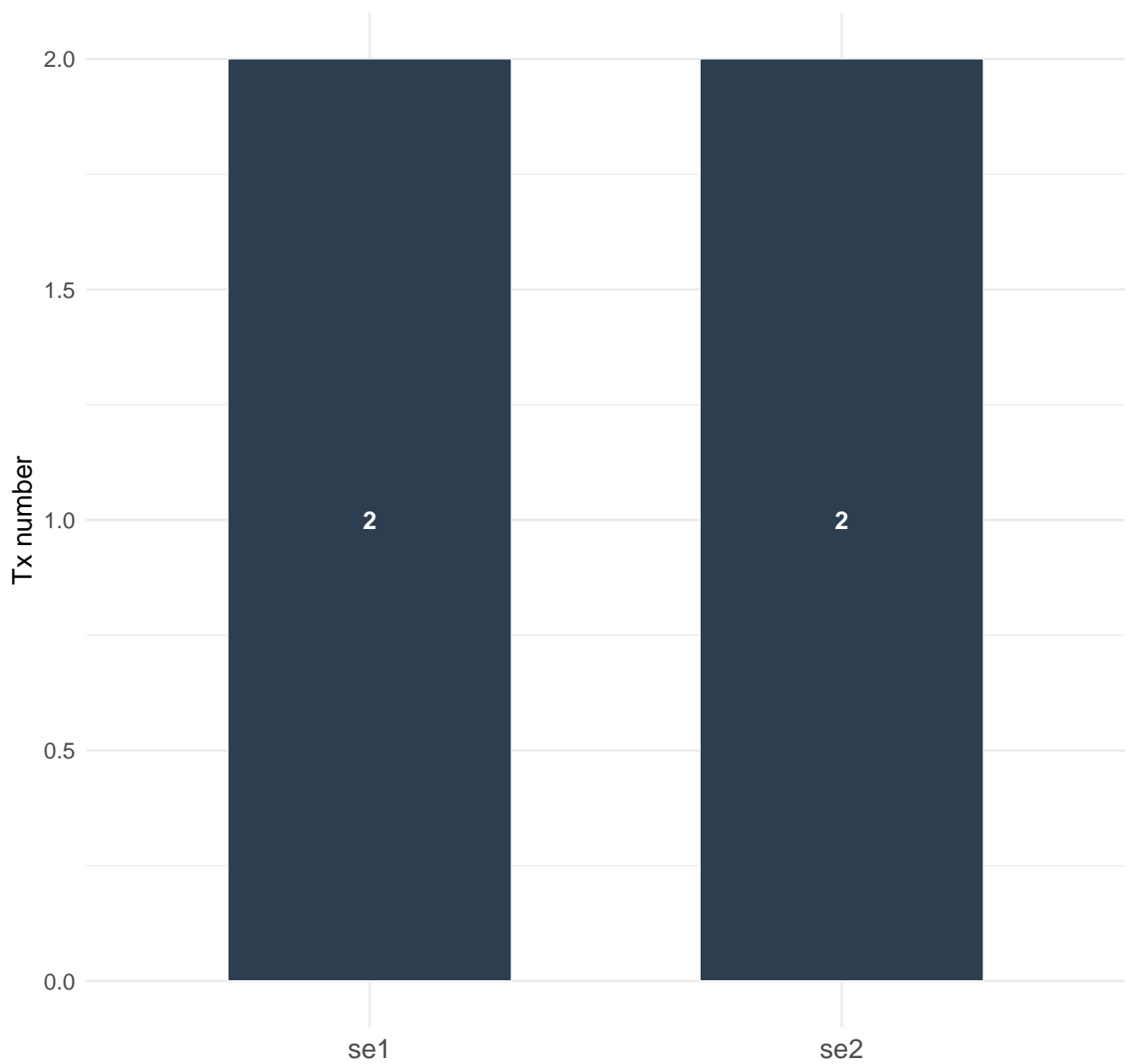


# Transcript Discovery Comparison

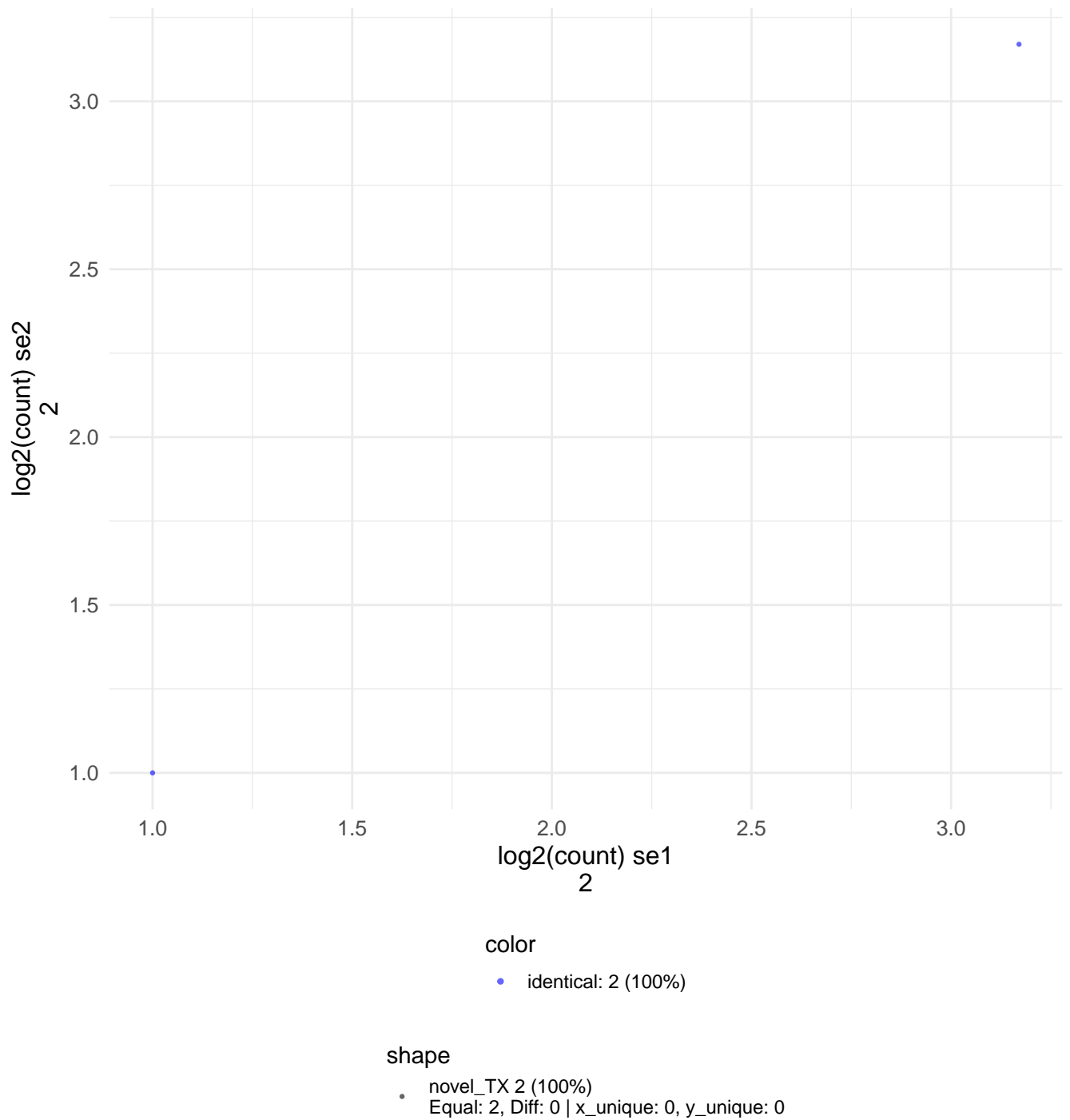


## Category

- Shared Tx (intron chain + start + end)
- Known Intron Chain
- Novel Intron Chain

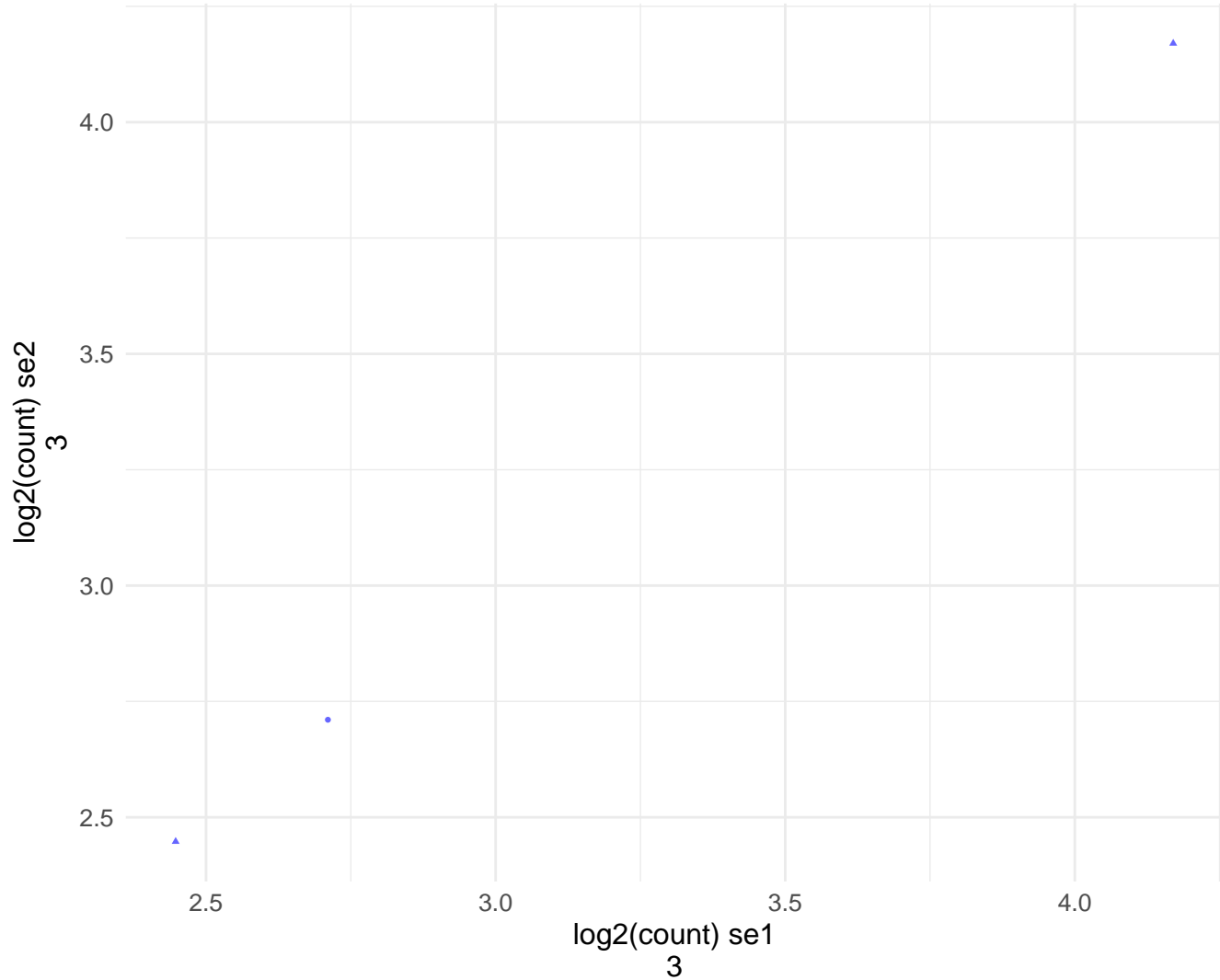
# Transcript Quantification Comparison

Sample: GIS\_ARG\_I002C\_Rep1\_Run1\_grch38\_sorted\_demo



# Transcript Quantification Comparison

Sample: GIS\_ARG\_I002C\_Rep1\_Run2\_grch38\_sorted\_demo



## shape

- anno\_TX 1 (33.3%)  
Equal: 3, Diff: 0 | x\_unique: 0, y\_unique: 0
- ▲ novel\_TX 2 (66.7%)  
Equal: 3, Diff: 0 | x\_unique: 0, y\_unique: 0

## color

- identical: 3 (100%)