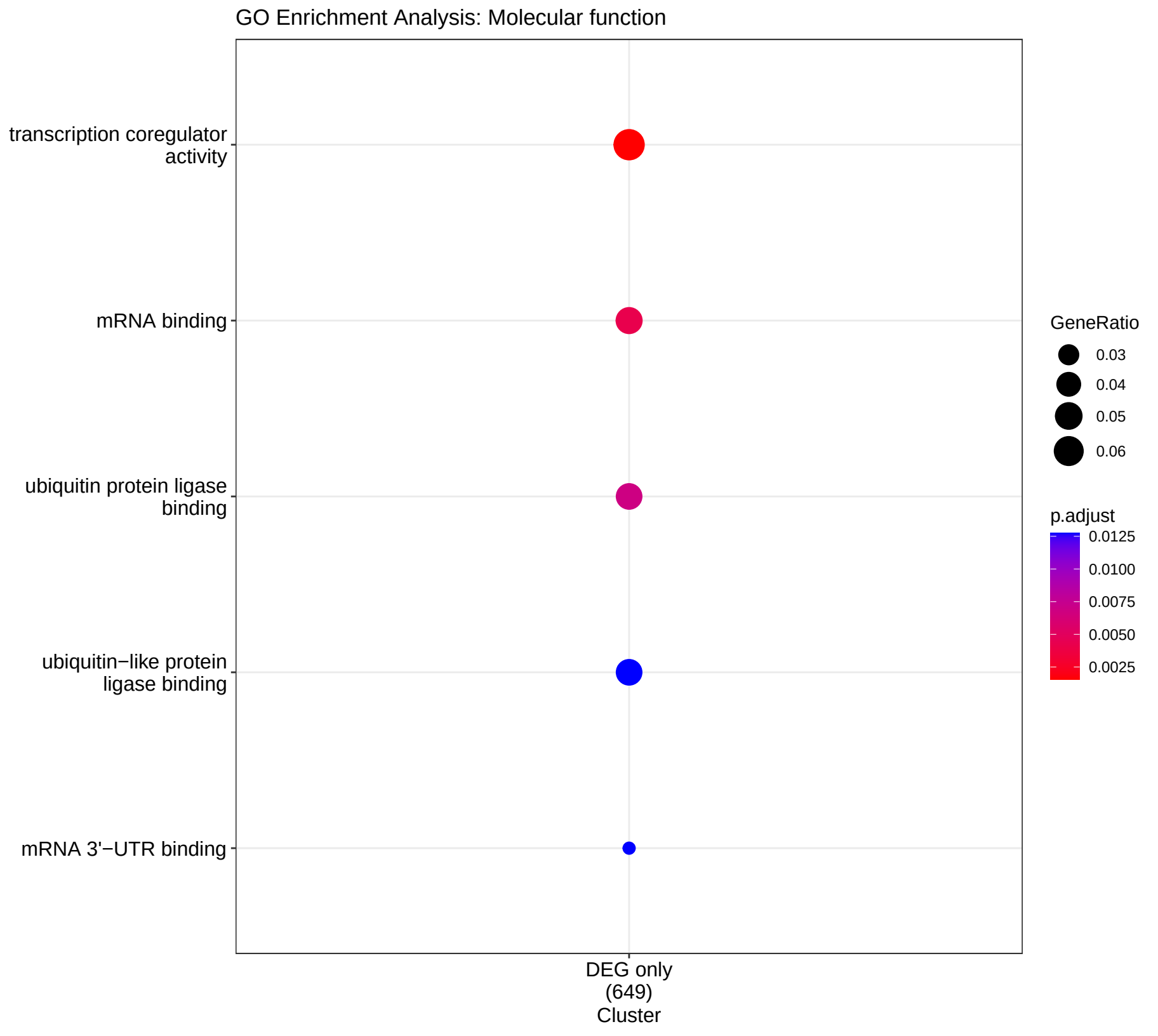
GO Enrichment Analysis: Molecular function
transcription coregulator activity
mRNA binding
ubiquitin protein ligase binding
ubiquitin−like protein ligase binding
mRNA 3'−UTR binding
DEG only
(649)
Cluster
GeneRatio
0.03
0.04
0.05
0.06
p.adjust
0.0125
0.0100
0.0075
0.0050
0.0025