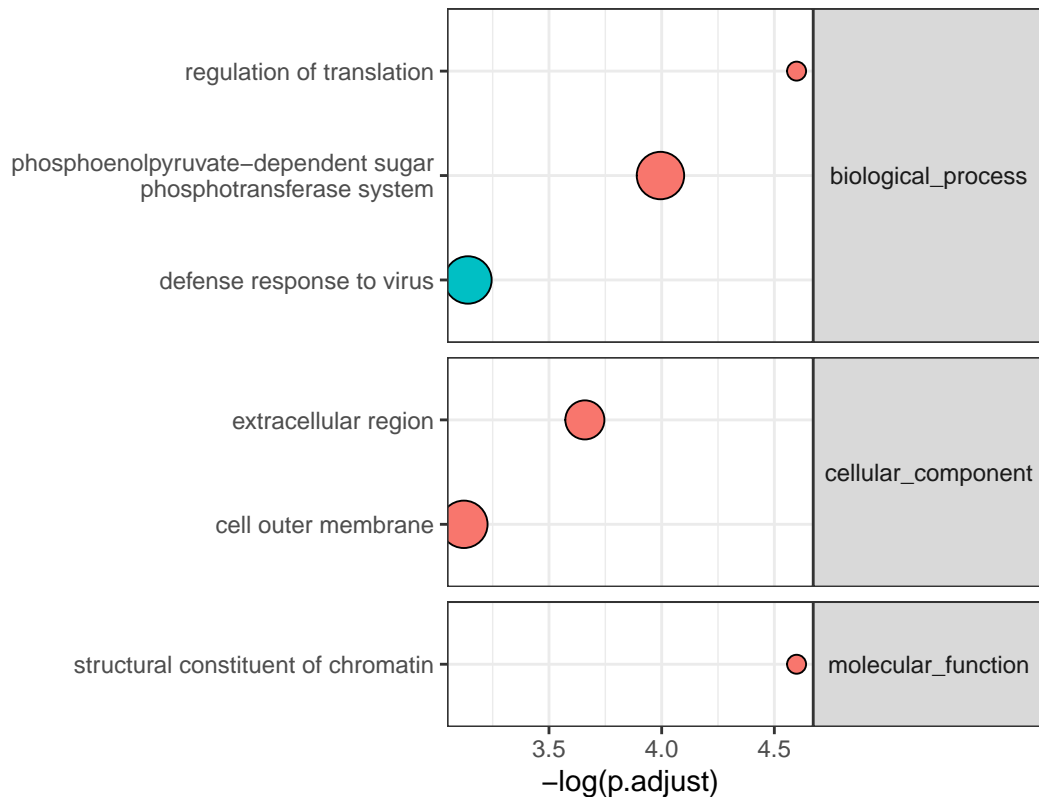


GO Enrichment

GO Term



Group

- gene
- inter

Significant_K_num

- 2.0
- 2.5
- 3.0
- 3.5
- 4.0