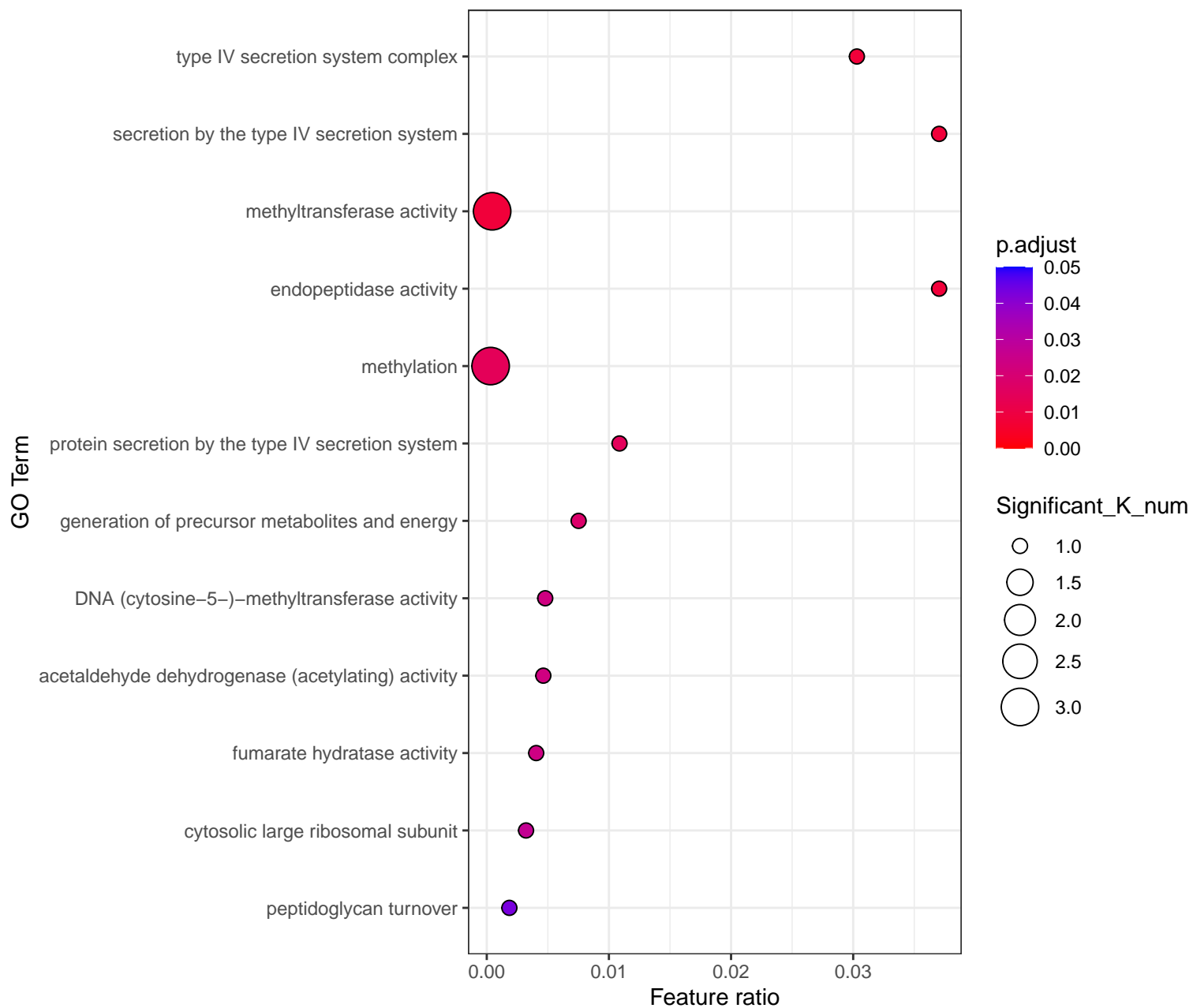
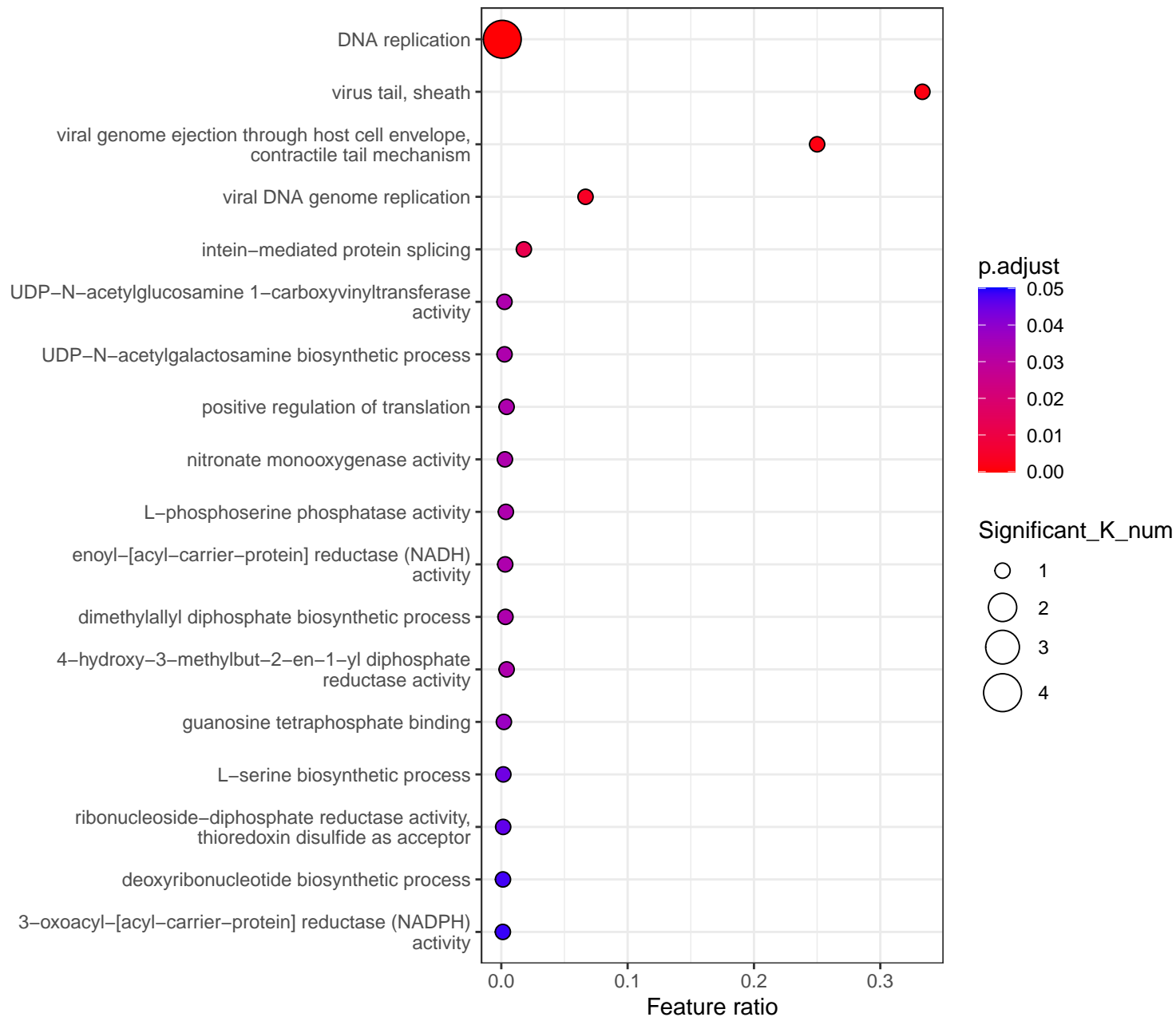


# s\_\_Escherichia coli



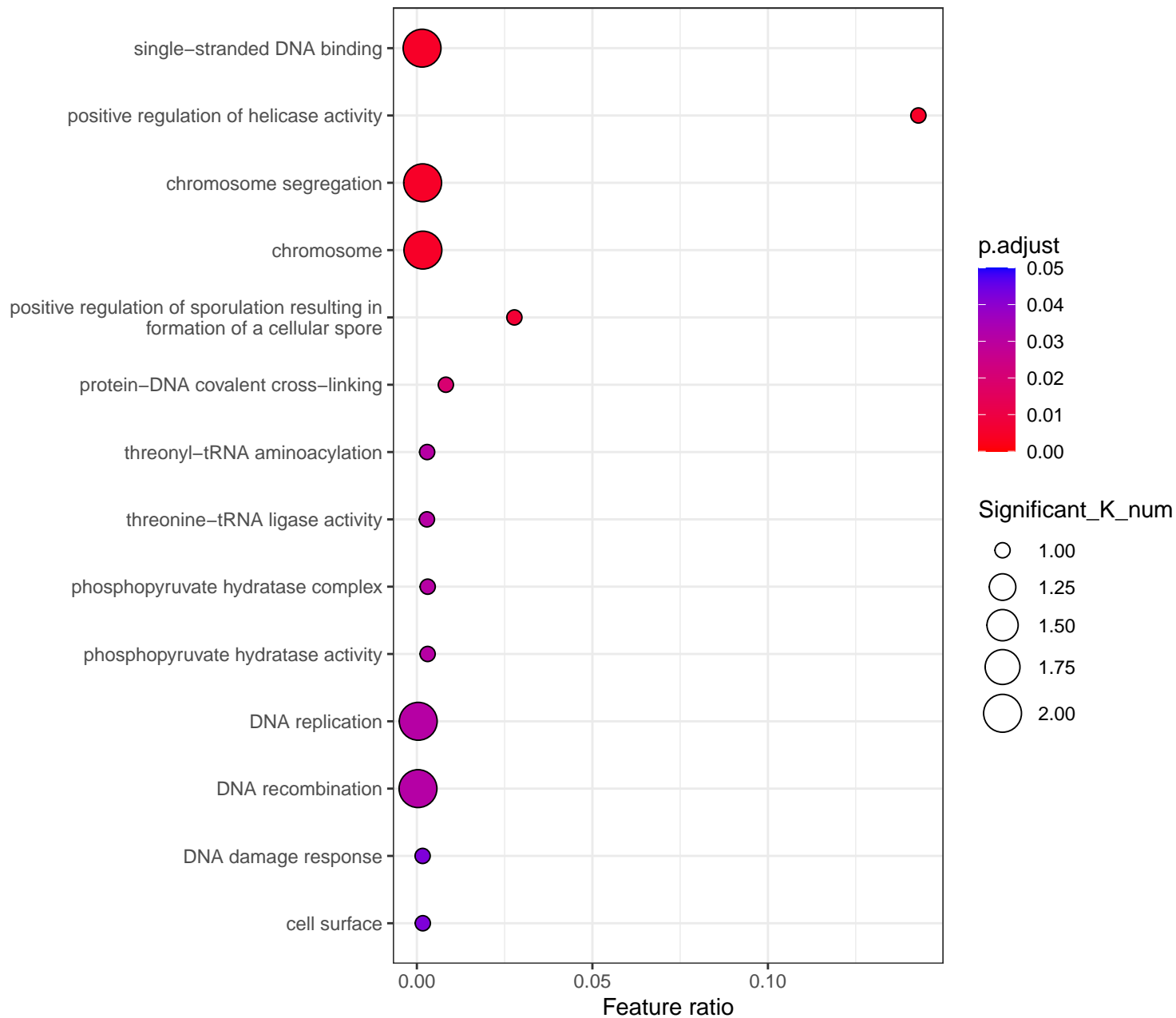
# s\_\_Campylobacter coli

GO Term



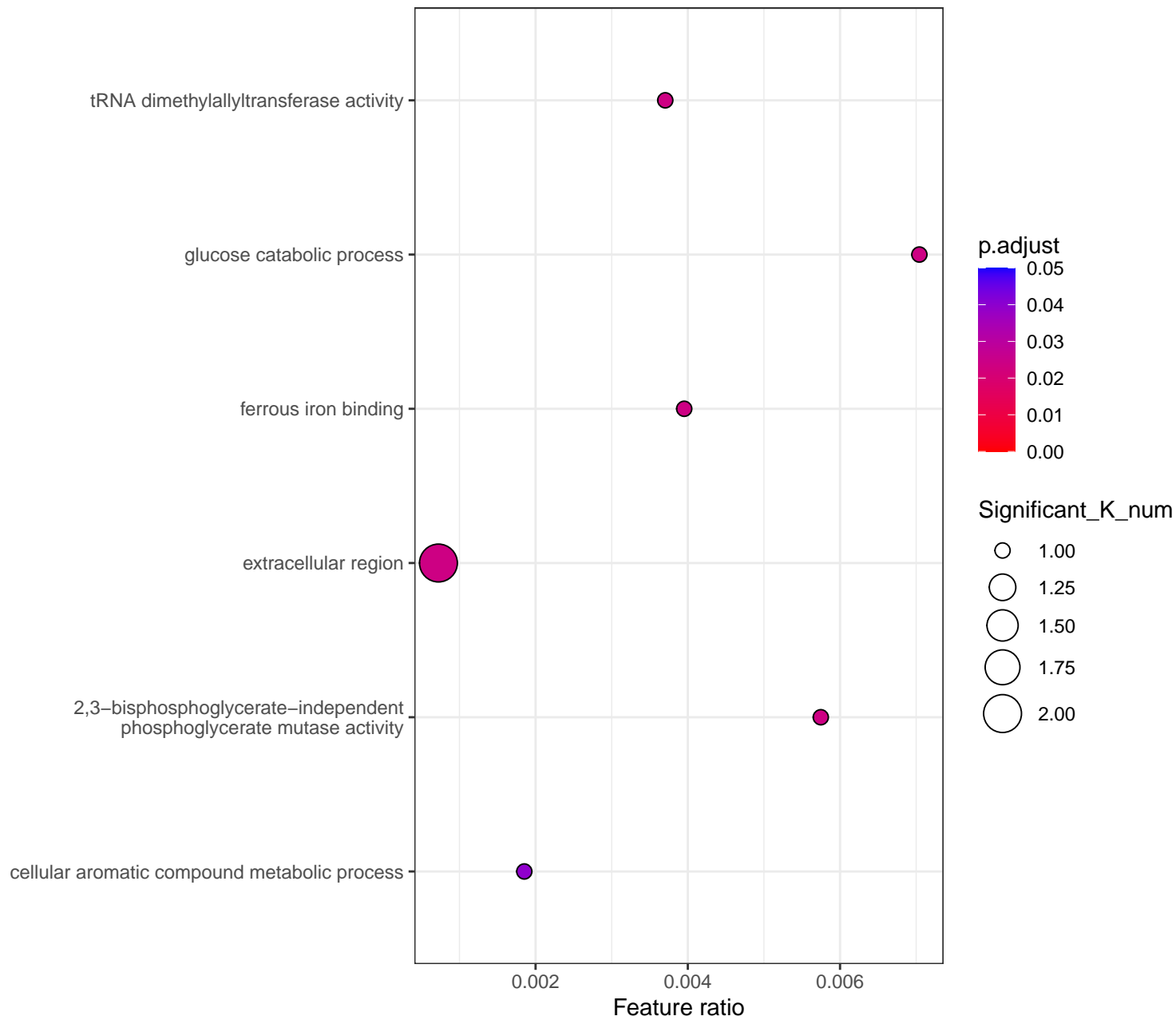
# s\_\_Salmonella enterica

GO Term



# s\_\_Klebsiella pneumoniae

GO Term



# s\_\_Neisseria meningitidis

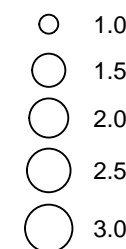
GO Term



p.adjust

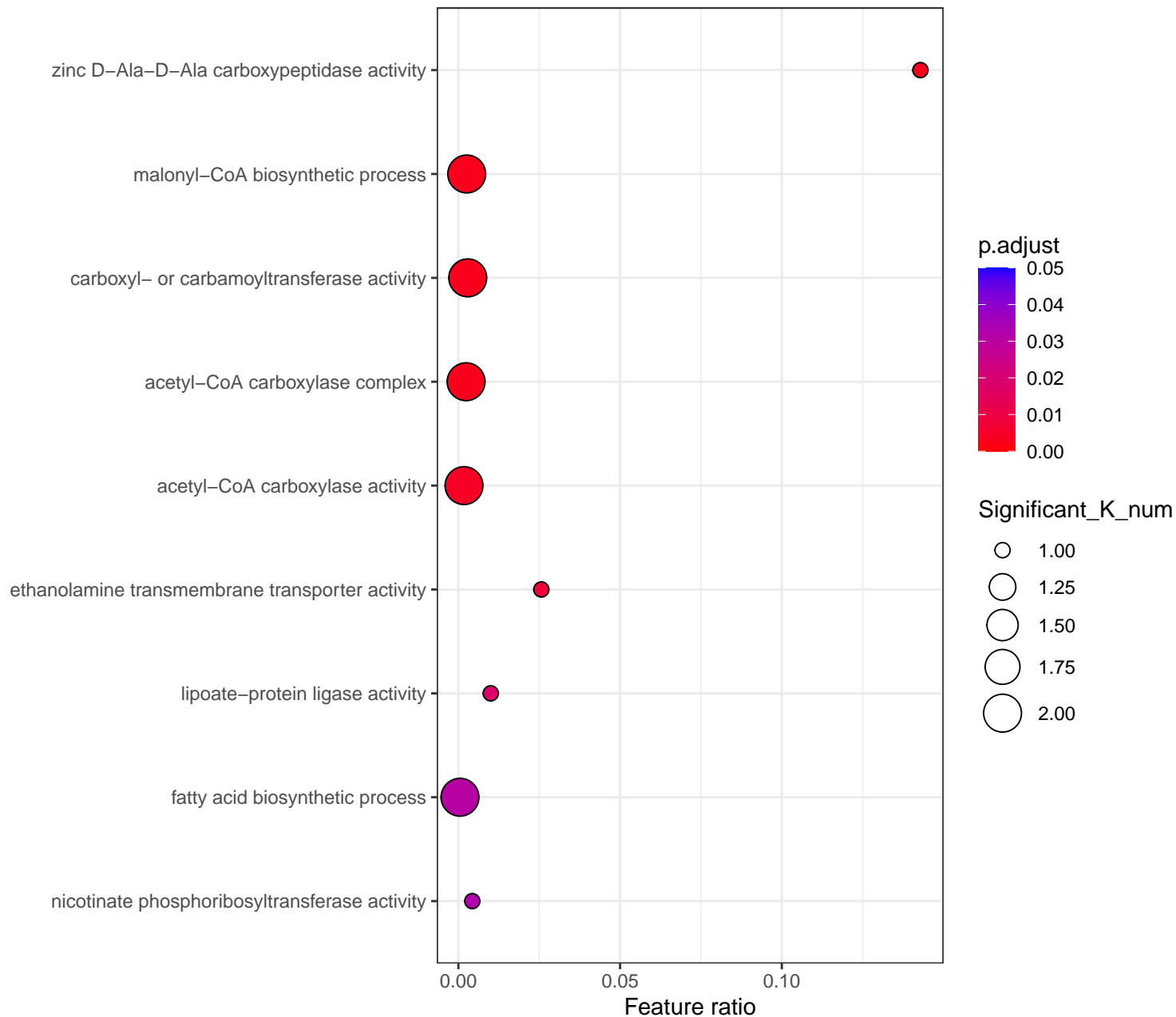


Significant\_K\_num



# s\_\_Clostridioides difficile

GO Term



# s\_\_Parabacteroides distasonis

GO Term

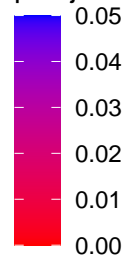
translation initiation factor activity

sodium ion transport

Significant\_K\_num

○ 2

p.adjust



0.00172

0.00174

0.00176

0.00178

Feature ratio

# s\_\_Acinetobacter baumannii

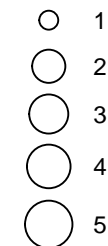
GO Term



p.adjust



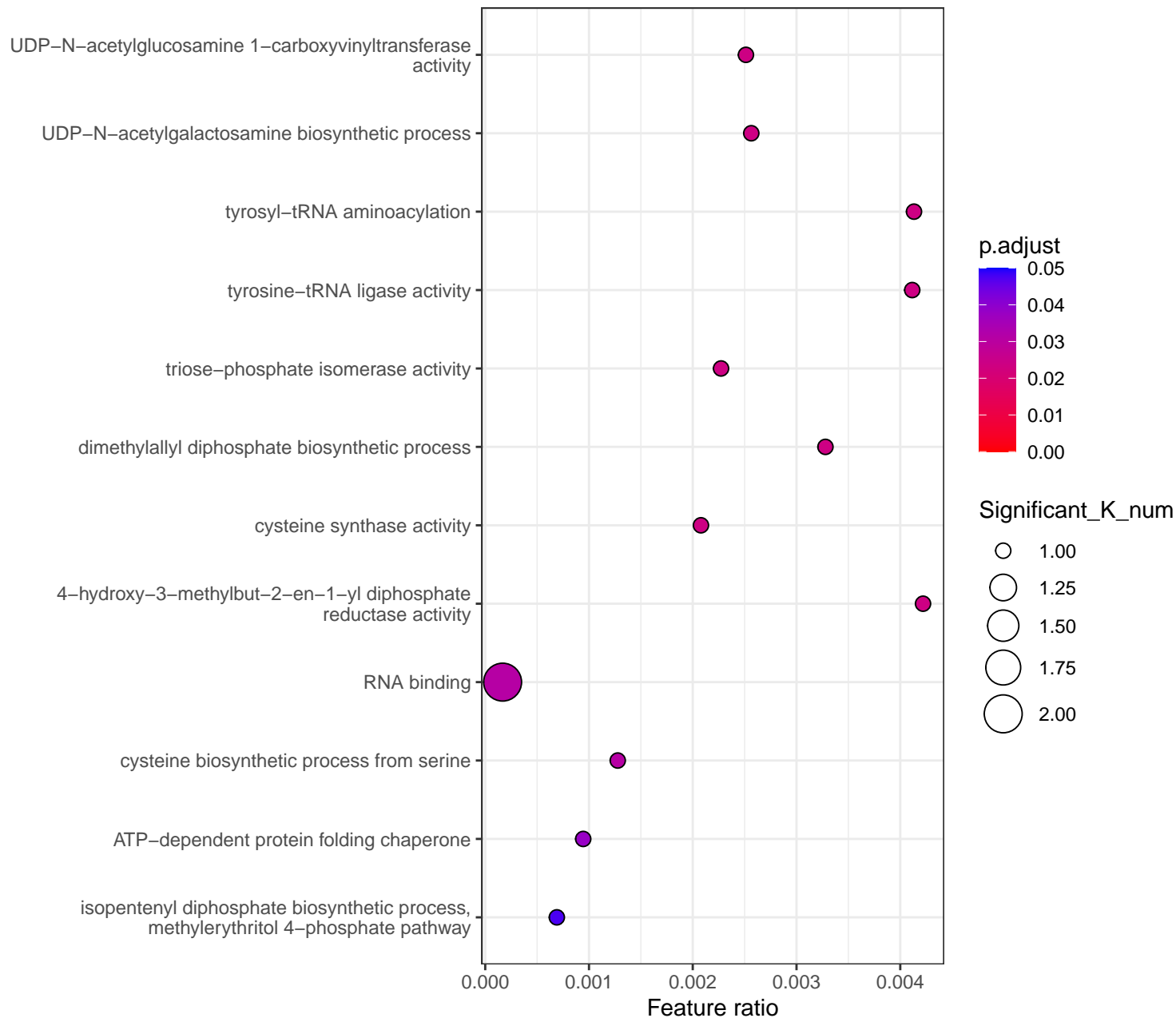
Significant\_K\_num





# s\_\_Megasphaera elsdenii

GO Term



# s\_\_Aggregatibacter actinomycetemcomitans

GO Term

