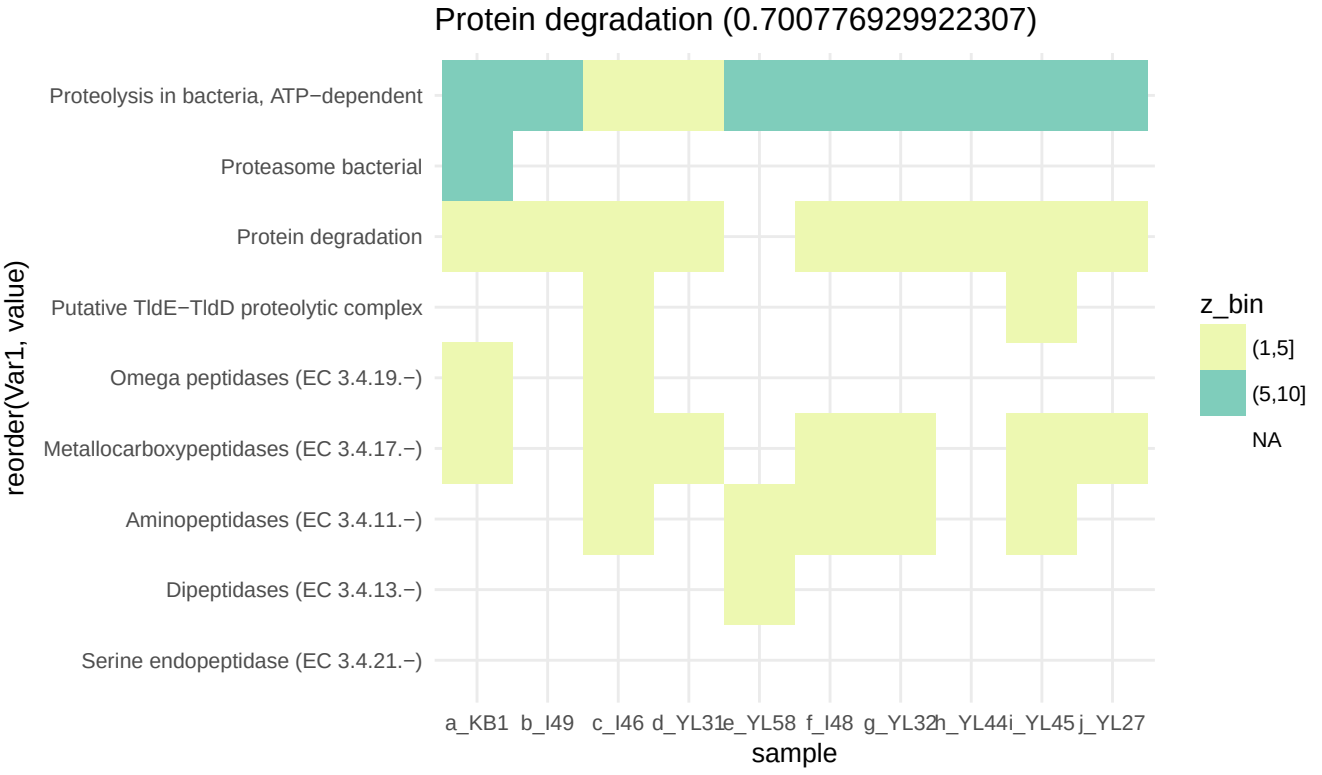
Protein degradation (0.700776929922307)
Proteolysis in bacteria, ATP−dependent
Proteasome bacterial
Protein degradation
Putative TldE−TldD proteolytic complex
Omega peptidases (EC 3.4.19.−)
Metallocarboxypeptidases (EC 3.4.17.−)
Aminopeptidases (EC 3.4.11.−)
Dipeptidases (EC 3.4.13.−)
Serine endopeptidase (EC 3.4.21.−)
reorder(Var1, value)
a_KB1
b_I49
c_I46
d_YL31
e_YL58
f_I48
g_YL32
h_YL44
i_YL45
j_YL27
sample
z_bin
(1,5]
(5,10]
NA