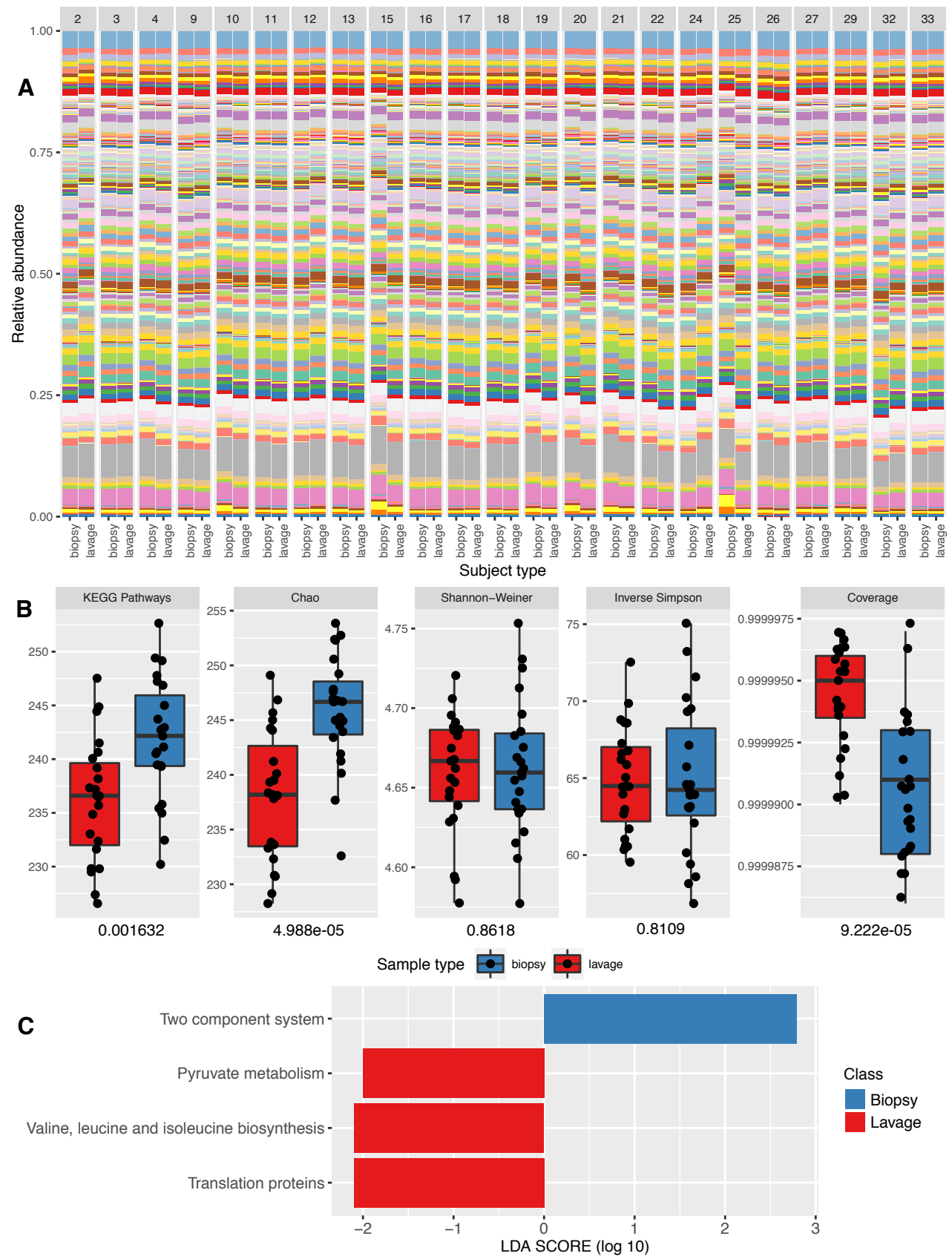

A
Relative abundance
1.00
0.75
0.50
0.25
0.00
2
3
4
9
10
11
12
13
15
16
17
18
19
20
21
22
24
25
26
27
29
32
33
biopsy
lavage
Subject type
B
KEGG Pathways
Chao
Shannon−Weiner
Inverse Simpson
Coverage
0.001632
4.988e-05
0.8618
0.8109
9.222e-05
Sample type
biopsy
lavage
C
Two component system
Pyruvate metabolism
Valine, leucine and isoleucine biosynthesis
Translation proteins
LDA SCORE (log 10)
Class
Biopsy
Lavage