Camping.b.lo.degree
Camping.b.hi.degree
95% confidence intervals
(a)
por 7.38e-3
abfD 5.91e-3
lacZ 9.25e-3
ACAT 0.011
fhs 2.26e-3
alpha-glucosidase 2.11e-3
xynA 4.36e-3
catalase 1.70e-3
sdhA 1.12e-3
porA 1.22e-3
treS 6.59e-3
bglA 0.017
sucC 7.02e-3
porB 2.22e-3
ppc 0.012
pta 9.81e-4
sucD 5.16e-3
cooS 6.41e-3
catalase-peroxidase 6.62e-3
NAGLU 5.51e-3
E4.2.1.2A 4.88e-3
pyc 0.015
porG 2.06e-3
sdhB 1.29e-3
tktA 0.014
ACO 0.023
bglB 0.035
acnB 3.91e-3
metF 0.015
bgaB 4.32e-4
IDH1 1.72e-3
accC 8.38e-3
E4.2.1.2B 0.023
fumC 0.023
vanD 3.72e-3
K15024 7.88e-4
chitinase 0.033
sdhC 1.56e-3
glpX 0.040
mdh 0.025
mcmA1 0.016
porD 1.80e-3
vanB 2.68e-3
amyA 0.045
korC 1.20e-3
frdA 0.018
PGK 6.42e-3
rpiA 0.015
GAPDH 5.42e-3
rexA 3.11e-3
vanA 4.76e-3
sdhD 9.64e-3
acsB 0.029
celF 9.49e-4
lip 8.16e-3
FBP 2.70e-3
ackA 0.026
glucoamylase 0.010
cdhE 0.041
meh 0.012
korD 1.03e-3
amylosucrase 8.84e-3
ALDO 0.022
rbcL 0.030
vdh 0.018
cdhD 0.029
mct 6.92e-3
ebgA 0.015
putative_chitinase 0.011
fdhB 6.32e-4
mcmA2 0.015
mcl 0.020
PRK 7.55e-4
pycA 5.15e-3
mch 4.08e-3
limB 0.033
ma 0.017
tyrosinase 0.017
frdC 0.034
smtB 4.30e-3
smtA1 5.89e-3
E3.3.2.8 0.028
K15016 6.46e-4
frdD 1.58e-3
K15039 8.47e-3
fdhA 0.034
K15020 0.016
ebgC 3.25e-3
csiR 0.034
K15019 1.59e-3
K15036 0.038
acsE 6.21e-3
K15052 0.018
K14465 0.011
chiA 0.017
K14469 0.035
amyM 0.032
p-value (corrected)
0.0
10.1
Mean proportion (%)
−2 −1 0 1 2 3 4 5
Difference in mean proportions (%)

Camping.specific.b.lo.degree
Camping.specific.b.hi.degree
95% confidence intervals
(b)
por 1.36e-3
abfD 5.80e-3
lacZ 0.017
fhs 4.97e-5
sdhA 1.57e-4
ACAT 0.021
catalase 1.57e-4
bglA 6.70e-3
alpha-glucosidase 3.16e-4
porA 3.70e-3
treS 3.26e-4
sucC 3.86e-5
pta 1.13e-5
ppc 7.64e-3
E4.2.1.2A 3.41e-3
pyc 0.011
sucD 3.56e-5
porB 4.82e-3
sdhB 4.56e-5
catalase-peroxidase 1.48e-4
chitinase 3.51e-5
cooS 0.012
acnB 5.90e-3
mdh 6.73e-3
porG 2.75e-3
E2.3.1.8 8.05e-3
pps 2.96e-3
E4.2.1.2B 0.030
fumC 0.030
ppdK 0.037
accC 9.14e-3
sdhC 3.88e-3
glpX 0.019
E3.2.1.89 3.70e-3
korA 0.022
IDH1 2.22e-3
mcmA1 7.62e-3
cstA 0.043
K15024 0.018
FBP 4.87e-5
vanB 2.57e-4
lacG 0.030
NAGLU 0.046
accA 4.25e-3
accD 2.77e-3
E3.2.1.1A 0.017
vanA 4.32e-6
dexB 0.023
sdhD 2.04e-3
vanD 9.79e-3
E3.2.1.96 0.039
porD 5.17e-3
glucoamylase 2.47e-3
korC 4.05e-3
meh 2.43e-3
rexA 0.048
rbcL 6.61e-3
amylosucrase 3.05e-3
vdh 8.53e-3
cytidine_deaminase 0.011
mct 6.77e-4
Mn-containing_catalase 0.014
putative_chitinase 1.38e-3
korD 7.86e-3
mcmA2 6.95e-3
mcl 0.010
PRK 3.56e-3
fdhB 0.012
frdB 3.62e-3
pycA 3.42e-3
mch 7.22e-7
frdC 7.26e-3
fchA 0.039
limB 0.021
ma 8.43e-3
tyrosinase 6.07e-3
smtB 4.79e-3
smtA1 7.73e-3
E3.2.1.58 0.012
E3.3.2.8 0.018
frdD 2.64e-3
K15020 7.42e-3
K15016 0.038
ebgC 6.77e-3
K15019 1.64e-4
K15036 0.010
acsE 7.98e-3
K15052 8.36e-3
K14465 2.95e-3
chiA 8.45e-3
K14469 0.024
amyM 0.049
p-value (corrected)
0.0
9.9
Mean proportion (%)
−2 −1 0 1 2 3 4 5
Difference in mean proportions (%)