Camping.b.hi.degree
No.camping.b.hi.degree
95% confidence intervals
(a)
lacZ 1.44e-3
por 0.018
bglX 0.016
ACAT 1.37e-3
catalase 3.74e-4
ACO 0.016
ppc 7.35e-3
fhs 0.011
ppdK 4.19e-3
pycB 0.022
pta 7.45e-4
porA 0.037
MUT 0.035
treS 0.012
sucC 0.022
catalase-peroxidase 1.98e-3
E3.2.1.89 0.015
GAPDH 1.10e-3
E4.2.1.2B 9.61e-3
fumC 9.61e-3
acnB 3.46e-3
IDH1 0.029
NAGLU 9.52e-3
sucD 0.024
chitinase 0.023
frdA 4.86e-3
accC 8.44e-3
bglB 0.047
tktA 0.013
E2.3.1.8 0.021
glpX 0.024
porG 0.040
rpiA 0.017
FBP 2.18e-3
accD 0.045
mcmA1 9.16e-3
vanB 2.87e-4
accB 4.16e-3
bgaB 6.91e-3
vanA 4.36e-3
korC 3.46e-3
sdhD 3.33e-3
ALDO 2.29e-3
lip 5.85e-3
ackA 0.030
glucoamylase 0.012
acsB 0.030
K15024 0.016
rbcL 0.024
PGK 0.049
meh 0.012
cytidine_deaminase 0.037
korD 6.34e-3
Mn-containing_catalase 0.022
vdh 0.010
cdhE 0.039
amylosucrase 7.56e-3
PRK 3.90e-3
putative_chitinase 3.04e-3
frdB 5.38e-3
pycA 3.29e-4
mct 0.012
cdhD 0.030
mcl 0.014
frdC 7.82e-3
mch 4.20e-3
MCEE 0.049
mcmA2 0.011
tyrosinase 0.011
E3.2.1.58 8.26e-3
smtB 4.16e-3
smtA1 5.91e-3
frdD 5.91e-3
K15016 2.92e-3
K15039 2.22e-4
fdhA 3.18e-4
csiR 4.31e-4
ebgC 8.15e-3
K15036 0.023
K15020 8.08e-3
K15019 3.83e-4
acsE 0.037
K14465 6.14e-3
K14469 0.043
chiA 0.049
amyM 8.04e-3
0.0
8.6
Mean proportion (%)
−4 −3 −2 −1 0 1 2
Difference in mean proportions (%)
Camping.specific.b.hi.degree
No.camping.specific.b.hi.degree
95% confidence intervals
(b)
lacZ 3.80e-4
por 4.24e-3
bglX 0.018
ACAT 1.07e-3
catalase 7.67e-4
ppc 3.43e-3
fhs 0.011
ACO 0.017
ppdK 3.39e-3
pta 3.79e-3
pycB 0.032
MUT 0.027
treS 7.61e-3
sucC 0.016
porA 0.045
catalase-peroxidase 2.61e-4
acnB 6.28e-3
pps 0.031
chitinase 2.45e-5
E3.2.1.89 0.013
GAPDH 8.03e-4
sucD 0.017
E4.2.1.2B 0.026
fumC 0.026
IDH1 0.025
tktA 0.011
bglB 0.044
NAGLU 5.66e-3
frdA 3.08e-3
E2.3.1.8 7.23e-3
glpX 0.010
FBP 3.53e-4
accC 0.034
porG 0.041
mcmA1 2.21e-3
vanB 5.96e-5
rpiA 0.012
accB 4.75e-3
vanA 4.74e-6
sdhD 4.03e-4
korC 1.38e-3
lip 2.45e-3
glucoamylase 2.03e-3
rbcL 0.012
ALDO 0.022
meh 3.69e-3
acsB 0.022
K15024 0.018
folD 0.032
korD 3.42e-4
Mn-containing_catalase 4.10e-3
cytidine_deaminase 0.038
vdh 2.47e-3
amylosucrase 1.87e-3
cdhE 0.023
PRK 1.02e-5
putative_chitinase 1.90e-4
frdB 7.28e-3
pycA 9.33e-4
mct 0.013
mcl 4.18e-3
frdC 0.011
mcmA2 3.44e-3
cdhD 0.021
mch 2.28e-7
tyrosinase 2.80e-3
ma 0.036
E3.2.1.58 1.31e-3
smtB 4.71e-3
smtA1 7.75e-3
frdD 7.47e-3
K15016 5.29e-3
K15039 1.25e-3
csiR 2.10e-4
ebgC 0.011
K15020 1.68e-3
K15036 3.06e-3
fdhA 0.010
K15019 9.81e-5
acsE 0.028
K14465 4.37e-4
K14469 0.020
chiA 0.024
amyM 0.011
0.0
8.7
Mean proportion (%)
−4 −3 −2 −1 0 1 2 3
Difference in mean proportions (%)
p-value (corrected)