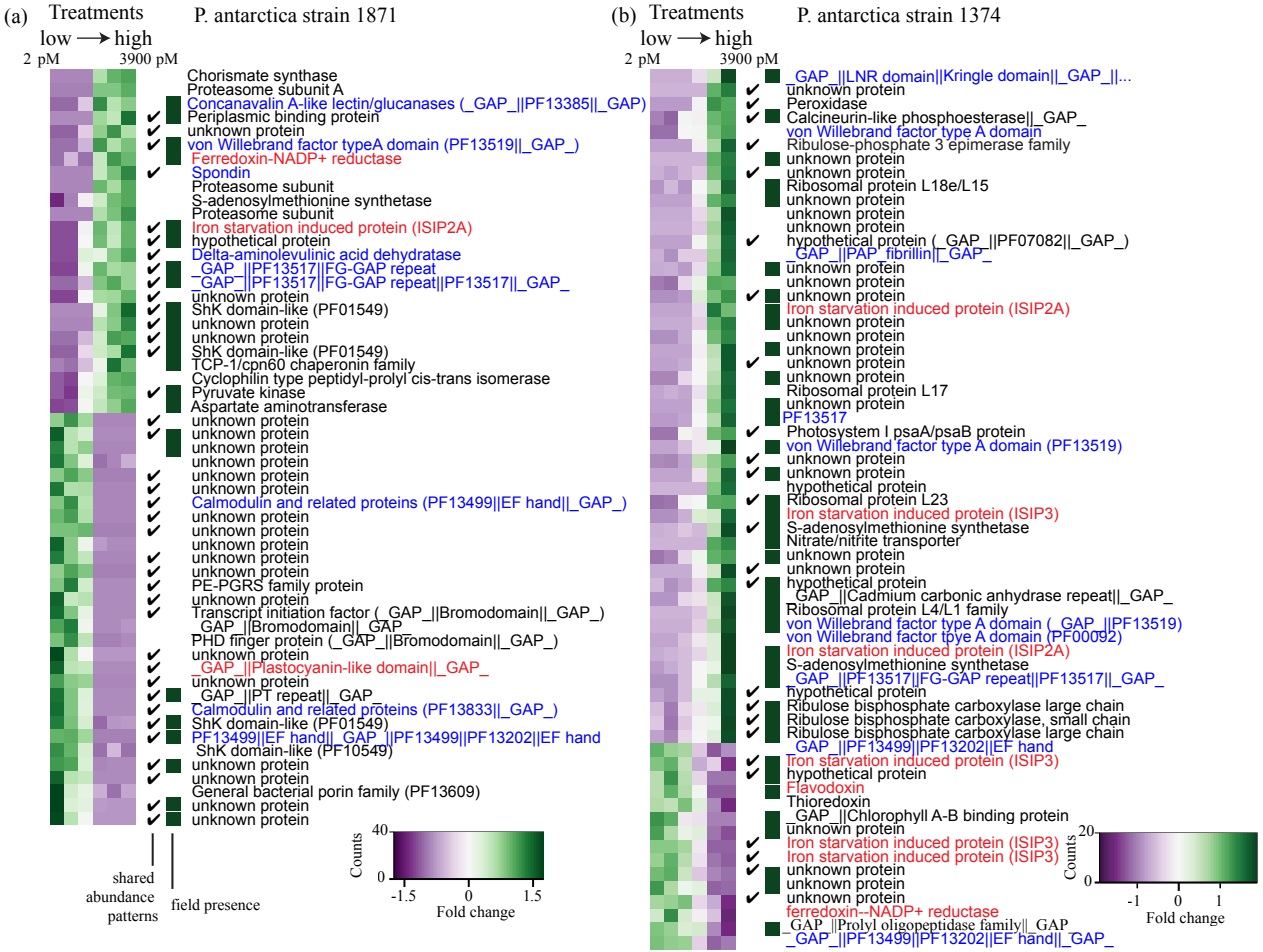
(a)
Treatments
low ⟶ high
2 pM
3900 pM
P. antarctica strain 1871
Chorismate synthase
Proteasome subunit A
Concanavalin A-like lectin/glucanases (_GAP_||PF13385||_GAP)
Periplasmic binding protein
unknown protein
von Willebrand factor typeA domain (PF13519||_GAP_)
Ferredoxin-NADP+ reductase
Spondin
Proteasome subunit
S-adenosylmethionine synthetase
Proteasome subunit
Iron starvation induced protein (ISIP2A)
hypothetical protein
Delta-aminolevulinic acid dehydratase
GAP||PF13517||FG-GAP repeat
GAP||PF13517||FG-GAP repeat||PF13517||_GAP_
unknown protein
ShK domain-like (PF01549)
unknown protein
unknown protein
ShK domain-like (PF01549)
TCP-1/cpn60 chaperonin family
Cyclophilin type peptidyl-prolyl cis-trans isomerase
Pyruvate kinase
Aspartate aminotransferase
unknown protein
unknown protein
unknown protein
unknown protein
unknown protein
unknown protein
Calmodulin and related proteins (PF13499||EF hand||_GAP_)
unknown protein
unknown protein
unknown protein
unknown protein
unknown protein
PE-PGRS family protein
unknown protein
Transcript initiation factor (_GAP_||Bromodomain||_GAP_)
GAP||Bromodomain||_GAP_
PHD finger protein (_GAP_||Bromodomain||_GAP_)
unknown protein
GAP||Plastocyanin-like domain||_GAP_
unknown protein
GAP||PT repeat||_GAP_
Calmodulin and related proteins (PF13833||_GAP_)
ShK domain-like (PF01549)
PF13499||EF hand||_GAP_||PF13499||PF13202||EF hand
ShK domain-like (PF10549)
unknown protein
unknown protein
General bacterial porin family (PF13609)
unknown protein
unknown protein
shared abundance patterns
field presence
Counts
40
0
-1.5
0
1.5
Fold change
(b)
Treatments
low ⟶ high
2 pM
3900 pM
P. antarctica strain 1374
GAP||LNR domain||Kringle domain||_GAP_||...
unknown protein
Peroxidase
Calcineurin-like phosphoesterase||_GAP_
von Willebrand factor type A domain
Ribulose-phosphate 3 epimerase family
unknown protein
unknown protein
Ribosomal protein L18e/L15
unknown protein
unknown protein
unknown protein
hypothetical protein (_GAP_||PF07082||_GAP_)
GAP||PAP_fibrilin||_GAP_
unknown protein
unknown protein
unknown protein
Iron starvation induced protein (ISIP2A)
unknown protein
unknown protein
unknown protein
unknown protein
unknown protein
Ribosomal protein L17
unknown protein
PF13517
Photosystem I psaA/psaB protein
von Willebrand factor type A domain (PF13519)
unknown protein
unknown protein
hypothetical protein
Ribosomal protein L23
Iron starvation induced protein (ISIP3)
S-adenosylmethionine synthetase
Nitrate/nitrite transporter
unknown protein
unknown protein
hypothetical protein
GAP||Cadmium carbonic anhydrase repeat||_GAP_
Ribosomal protein L4/L1 family
von Willebrand factor type A domain (_GAP_||PF13519)
von Willebrand factor type A domain (PF00092)
Iron starvation induced protein (ISIP2A)
S-adenosylmethionine synthetase
GAP||PF13517||FG-GAP repeat||PF13517||_GAP_
hypothetical protein
Ribulose bisphosphate carboxylase large chain
Ribulose bisphosphate carboxylase, small chain
Ribulose bisphosphate carboxylase large chain
GAP||PF13499||PF13202||EF hand
Iron starvation induced protein (ISIP3)
hypothetical protein
Flavodoxin
Thioredoxin
GAP||Chlorophyll A-B binding protein
unknown protein
Iron starvation induced protein (ISIP3)
Iron starvation induced protein (ISIP3)
unknown protein
unknown protein
unknown protein
ferredoxin--NADP+ reductase
GAP||Prolyl oligopeptidase family||_GAP_
GAP||PF13499||PF13202||EF hand||_GAP_
Counts
20
0
-1
0
1
Fold change