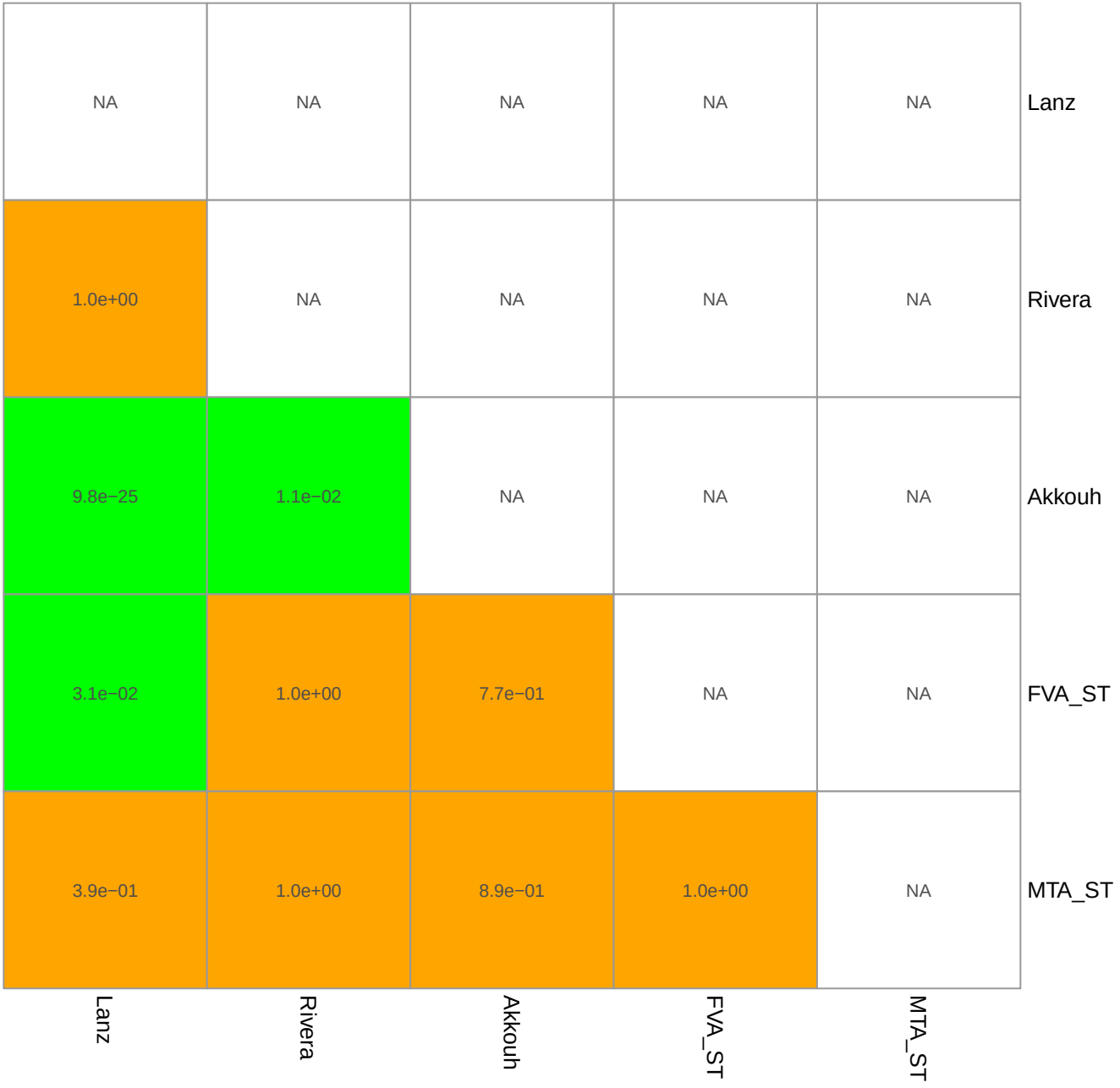
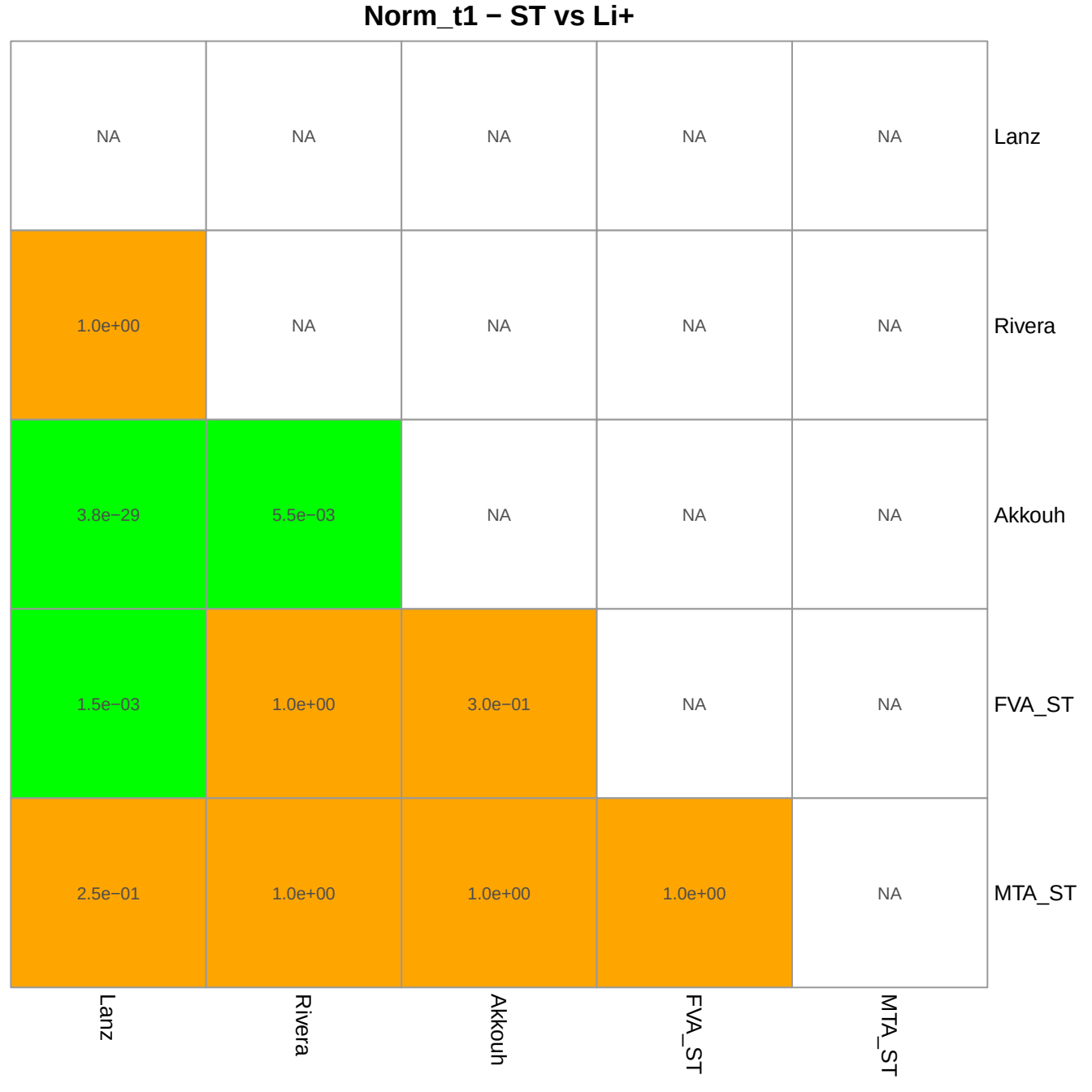
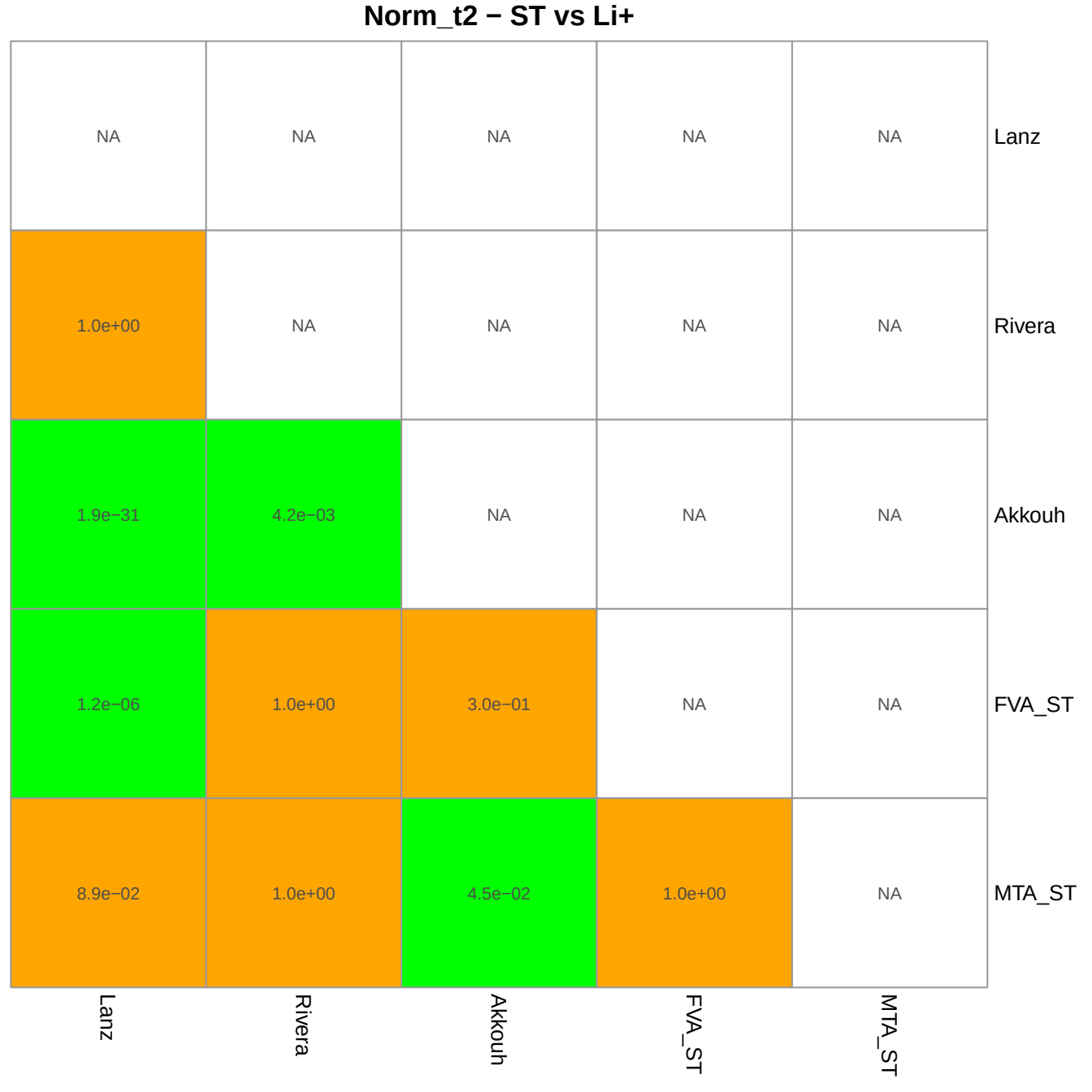
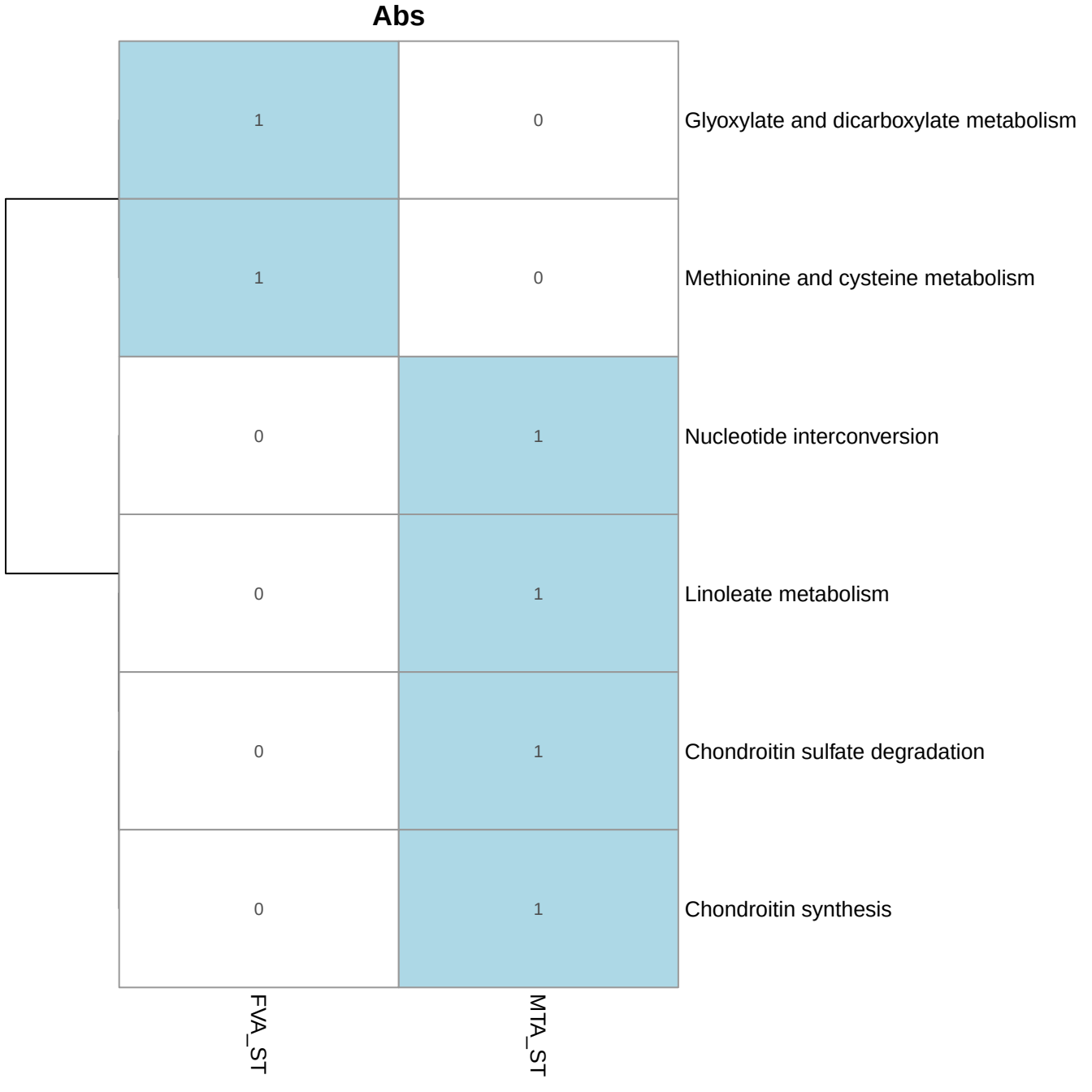
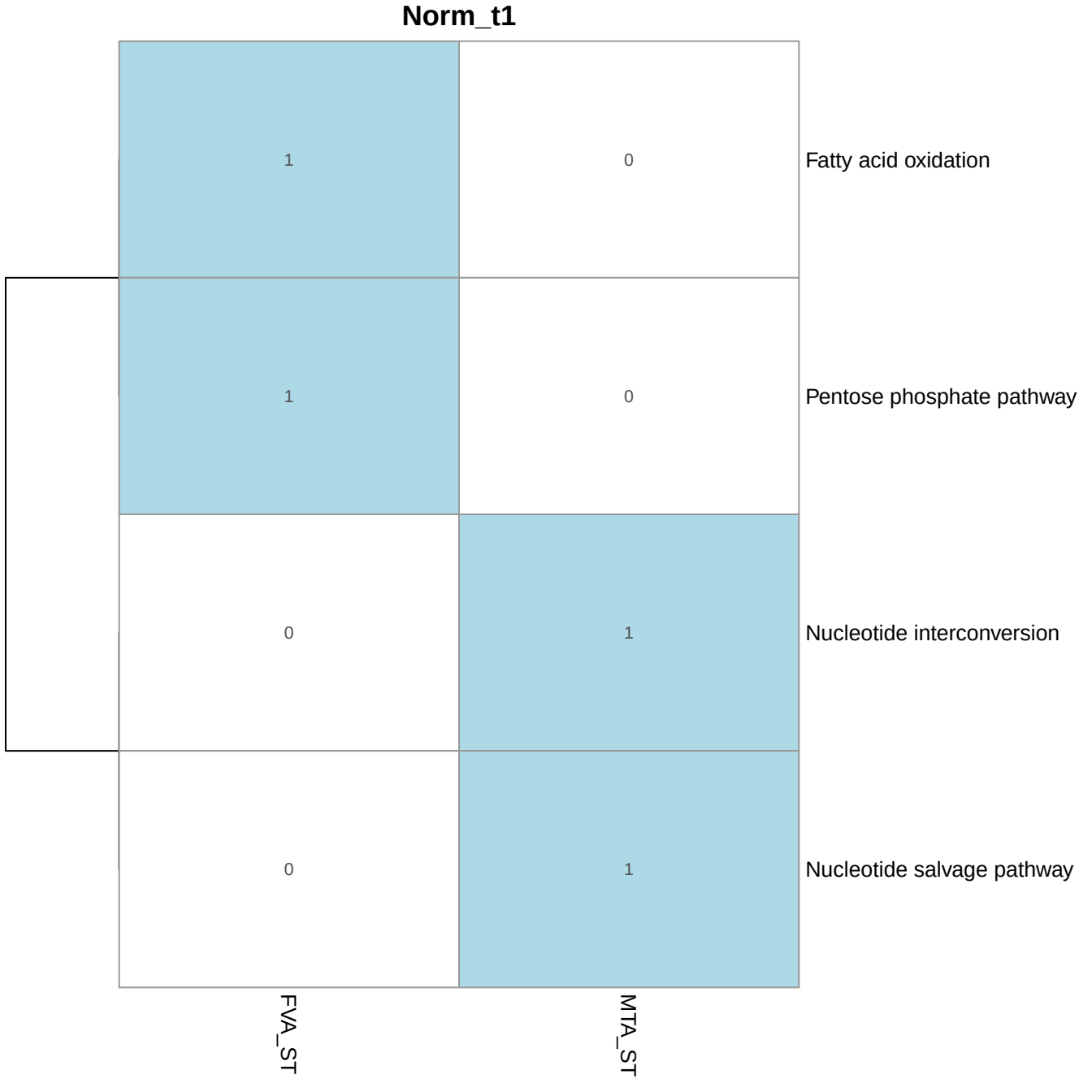
Abs − ST vs Li+

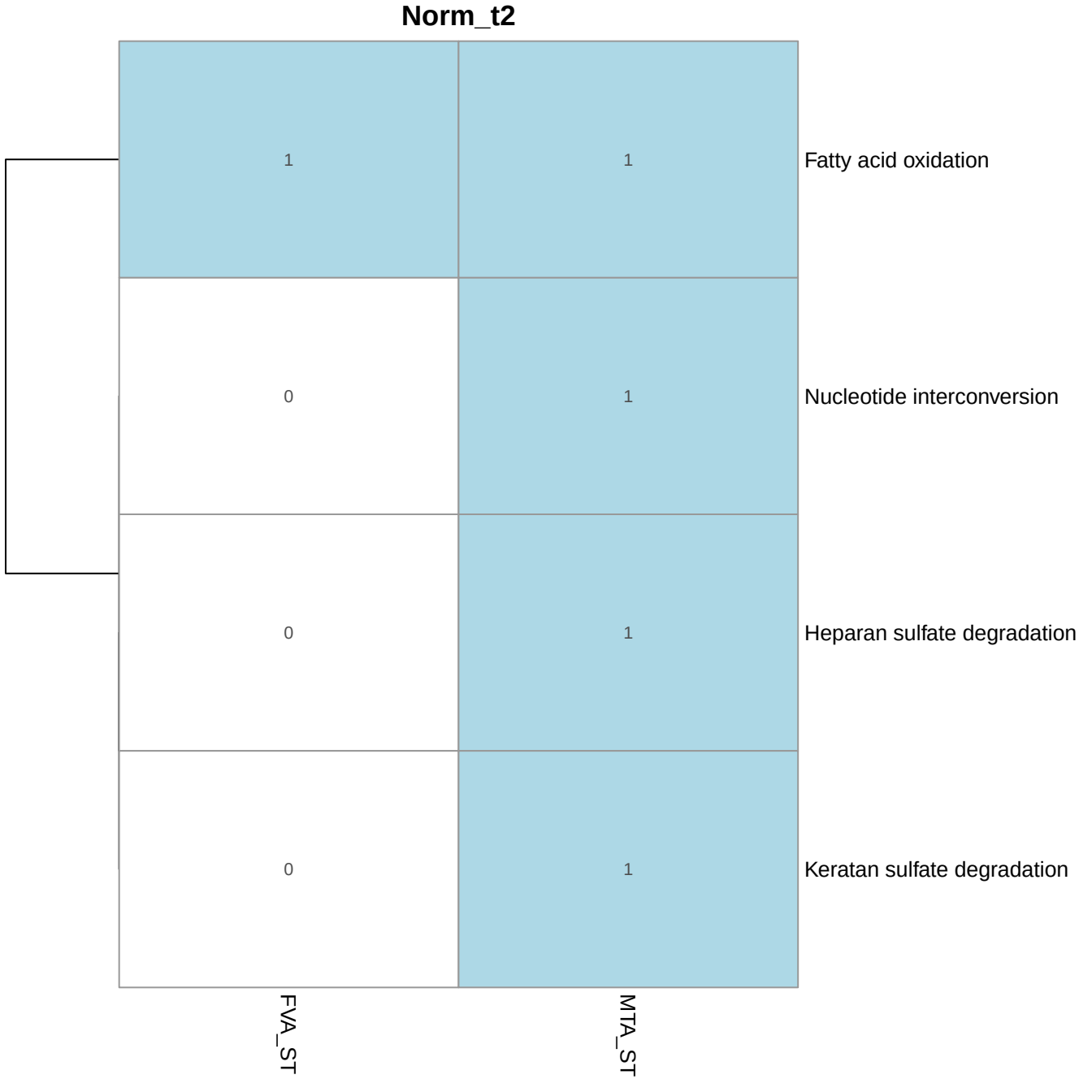

NA NA NA NA NA Lanz
1.0e+00 NA NA NA NA Rivera
9.8e−25 1.1e−02 NA NA NA Akkouh
3.1e−02 1.0e+00 7.7e−01 NA NA FVA_ST
3.9e−01 1.0e+00 8.9e−01 1.0e+00 NA MTA_ST
Lanz
Rivera
Akkouh
FVA_ST
MTA_ST

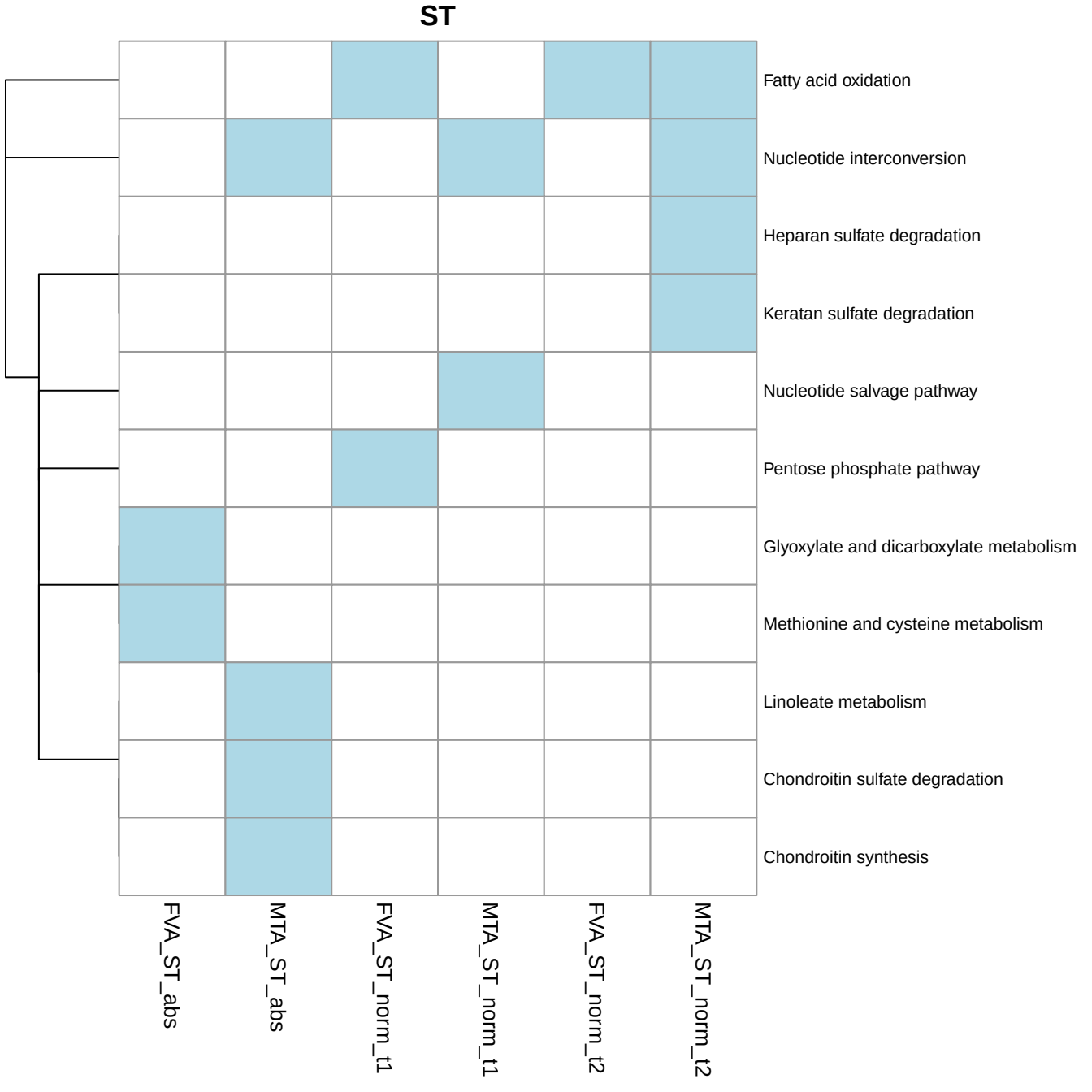

Norm_t1 − ST vs Li+
NA
NA
NA
NA
NA
Lanz
1.0e+00
NA
NA
NA
NA
Rivera
3.8e−29
5.5e−03
NA
NA
NA
Akkouh
1.5e−03
1.0e+00
3.0e−01
NA
NA
FVA_ST
2.5e−01
1.0e+00
1.0e+00
1.0e+00
NA
MTA_ST
Lanz
Rivera
Akkouh
FVA_ST
MTA_ST

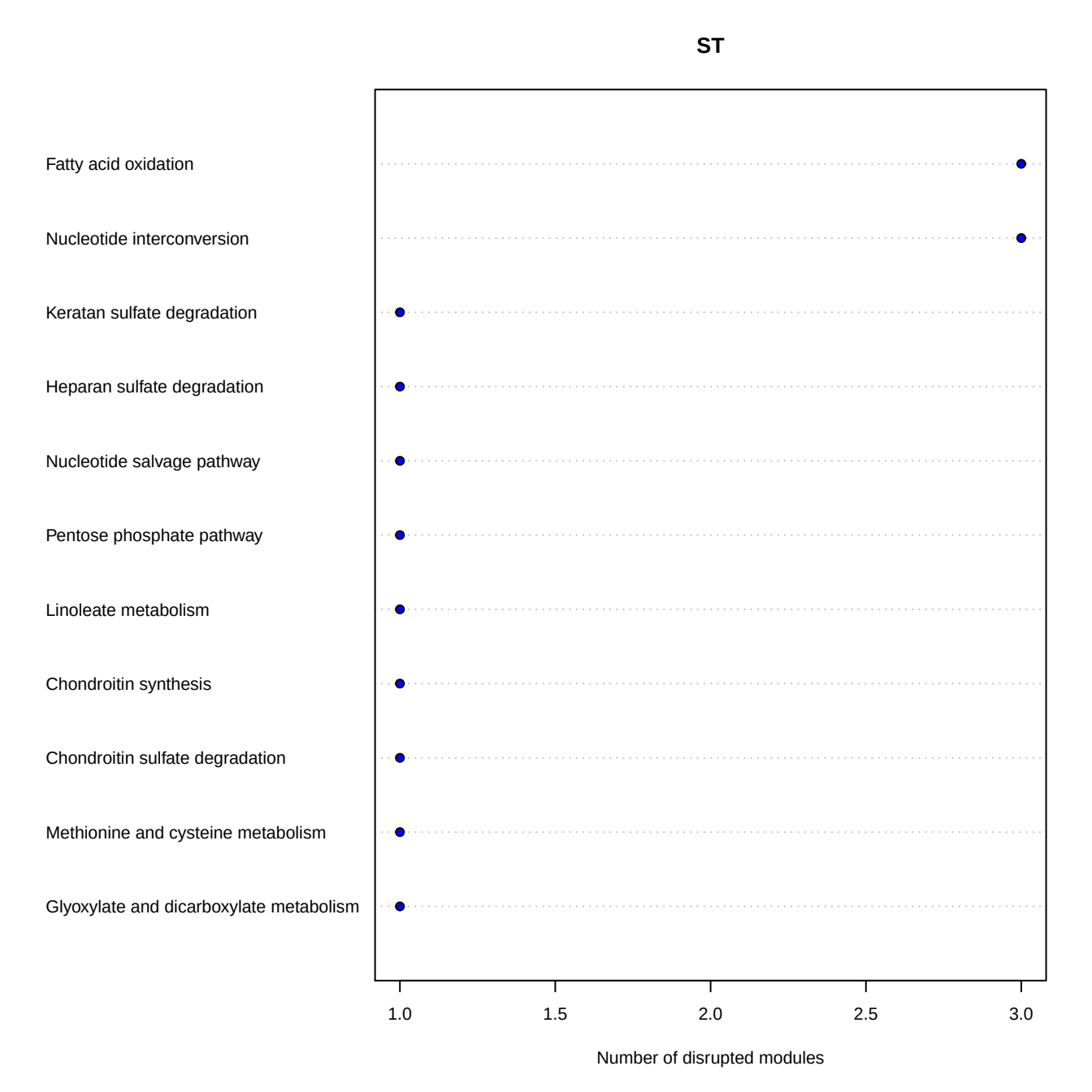

Norm_t2 − ST vs Li+
NA
NA
NA
NA
NA
Lanz
1.0e+00
NA
NA
NA
NA
Rivera
1.9e−31
4.2e−03
NA
NA
NA
Akkouh
1.2e−06
1.0e+00
3.0e−01
NA
NA
FVA_ST
8.9e−02
1.0e+00
4.5e−02
1.0e+00
NA
MTA_ST
Lanz
Rivera
Akkouh
FVA_ST
MTA_ST

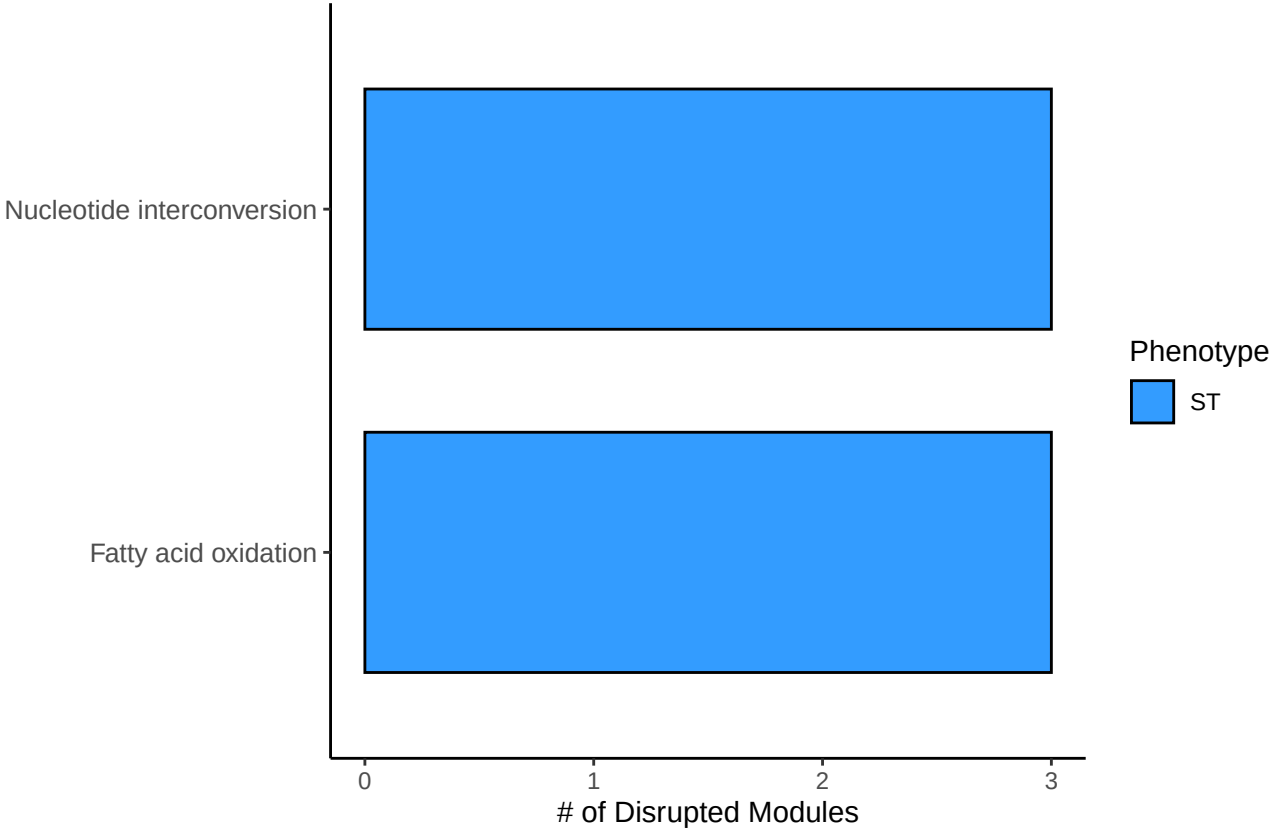

Abs
1
0
Glyoxylate and dicarboxylate metabolism
1
0
Methionine and cysteine metabolism
0
1
Nucleotide interconversion
0
1
Linoleate metabolism
0
1
Chondroitin sulfate degradation
0
1
Chondroitin synthesis
FVA_ST
MTA_ST

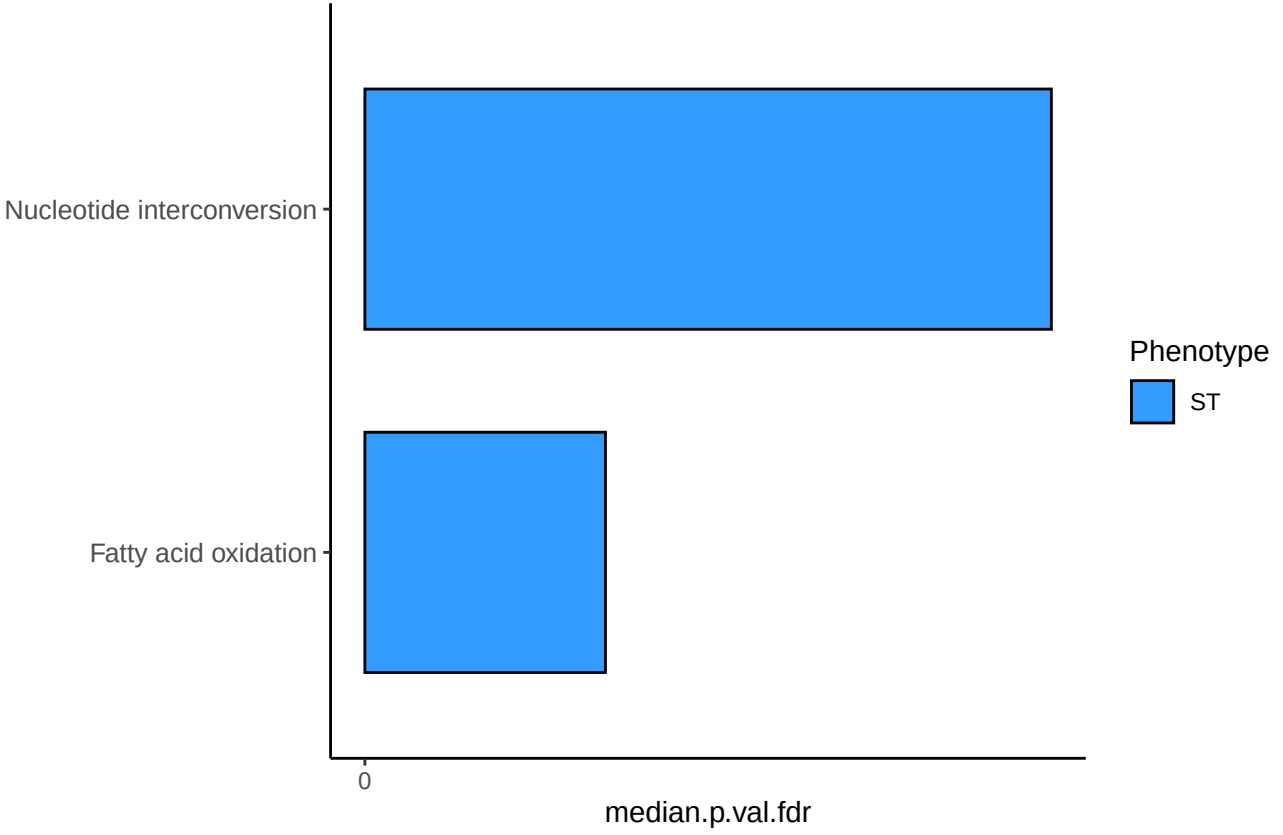

Norm_t1
1
0
Fatty acid oxidation
1
0
Pentose phosphate pathway
0
1
Nucleotide interconversion
0
1
Nucleotide salvage pathway
FVA_ST
MTA_ST


Norm_t2
1
1
Fatty acid oxidation
0
1
Nucleotide interconversion
0
1
Heparan sulfate degradation
0
1
Keratan sulfate degradation
FVA_ST
MTA_ST


ST
Fatty acid oxidation
Nucleotide interconversion
Heparan sulfate degradation
Keratan sulfate degradation
Nucleotide salvage pathway
Pentose phosphate pathway
Glyoxylate and dicarboxylate metabolism
Methionine and cysteine metabolism
Linoleate metabolism
Chondroitin sulfate degradation
Chondroitin synthesis
FVA_ST_abs
MTA_ST_abs
FVA_ST_norm_t1
MTA_ST_norm_t1
FVA_ST_norm_t2
MTA_ST_norm_t2


ST
Fatty acid oxidation
Nucleotide interconversion
Keratan sulfate degradation
Heparan sulfate degradation
Nucleotide salvage pathway
Pentose phosphate pathway
Linoleate metabolism
Chondroitin synthesis
Chondroitin sulfate degradation
Methionine and cysteine metabolism
Glyoxylate and dicarboxylate metabolism
1.0
1.5
2.0
2.5
3.0
Number of disrupted modules


Nucleotide interconversion
Fatty acid oxidation
0
1
2
3
# of Disrupted Modules
Phenotype
ST


Nucleotide interconversion
Fatty acid oxidation
0
median.p.val.fdr
Phenotype
ST