NA NA NA NA Lanz

1.0e+00 NA NA NA NA Rivera

Abs - ST vs Li+

NA

7.7e-01

8.9e-01

Akkouh

9.8e-25

3.1e-02

3.9e-01

Lanz

1.1e-02

1.0e+00

1.0e+00

Rivera

NA

NA

1.0e+00

Akkouh

FVA\_ST

MTA\_ST

NA

NA

NA

NA NA NA NA NA Lanz 1.0e+00 NA NA NA NA Rivera

Norm\_t1 - ST vs Li+

NA

3.0e-01

1.0e+00

Akkouh

3.8e-29

1.5e-03

2.5e-01

Lanz

5.5e-03

1.0e+00

1.0e+00

Rivera

NA NA

1.0e+00

Akkouh NA NA

NA

FVA\_ST MTA\_ST NA NA NA NA Lanz

1.0e+00 NA NA NA NA NA Rivera

Norm\_t2 - ST vs Li+

NA

3.0e-01

4.5e-02

Akkouh

1.9e-31

1.2e-06

8.9e-02

Lanz

4.2e-03

1.0e+00

1.0e+00

Rivera

NA NA

1.0e+00

FVA\_ST

Akkouh

FVA\_ST

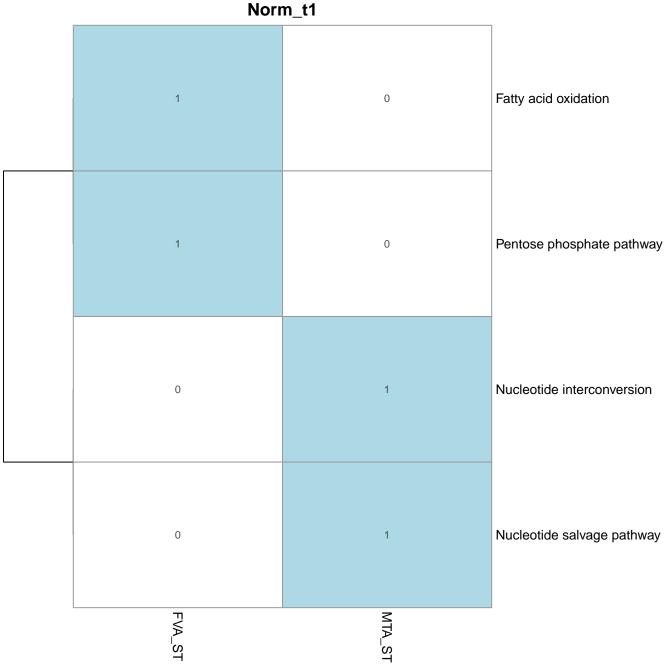
MTA\_ST

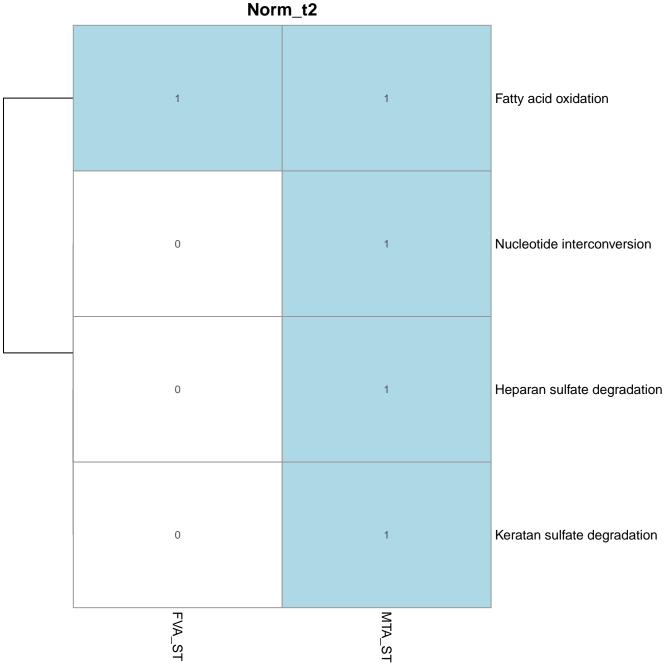
NA

NA

NA

**Abs** Glyoxylate and dicarboxylate metabolism 0 Methionine and cysteine metabolism Nucleotide interconversion 0 Linoleate metabolism 0 0 Chondroitin sulfate degradation Chondroitin synthesis 0





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 MTA\_ST\_norm\_t2

