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

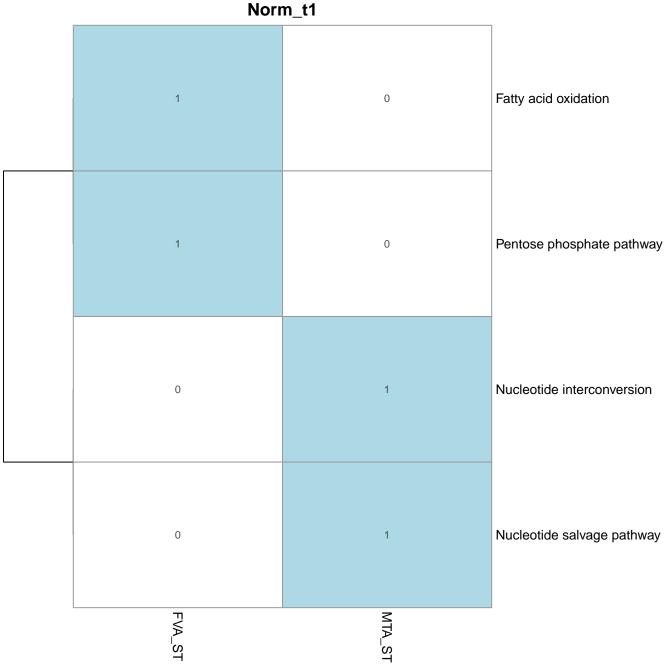
MTA_ST

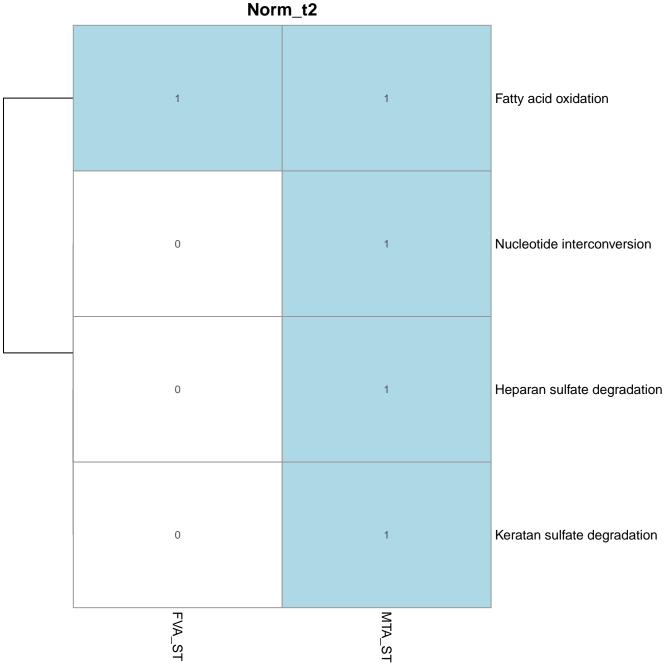
NA

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

