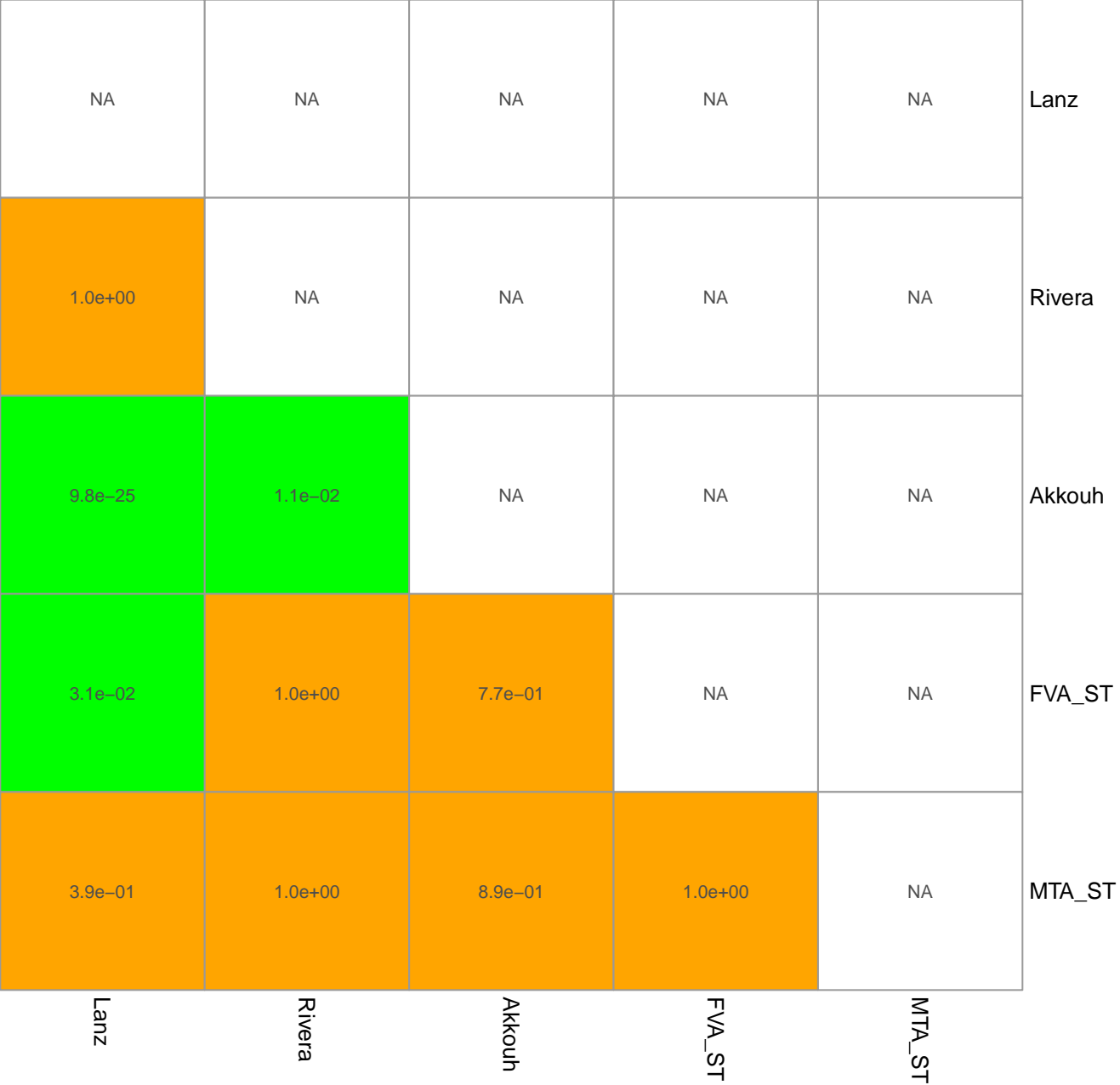
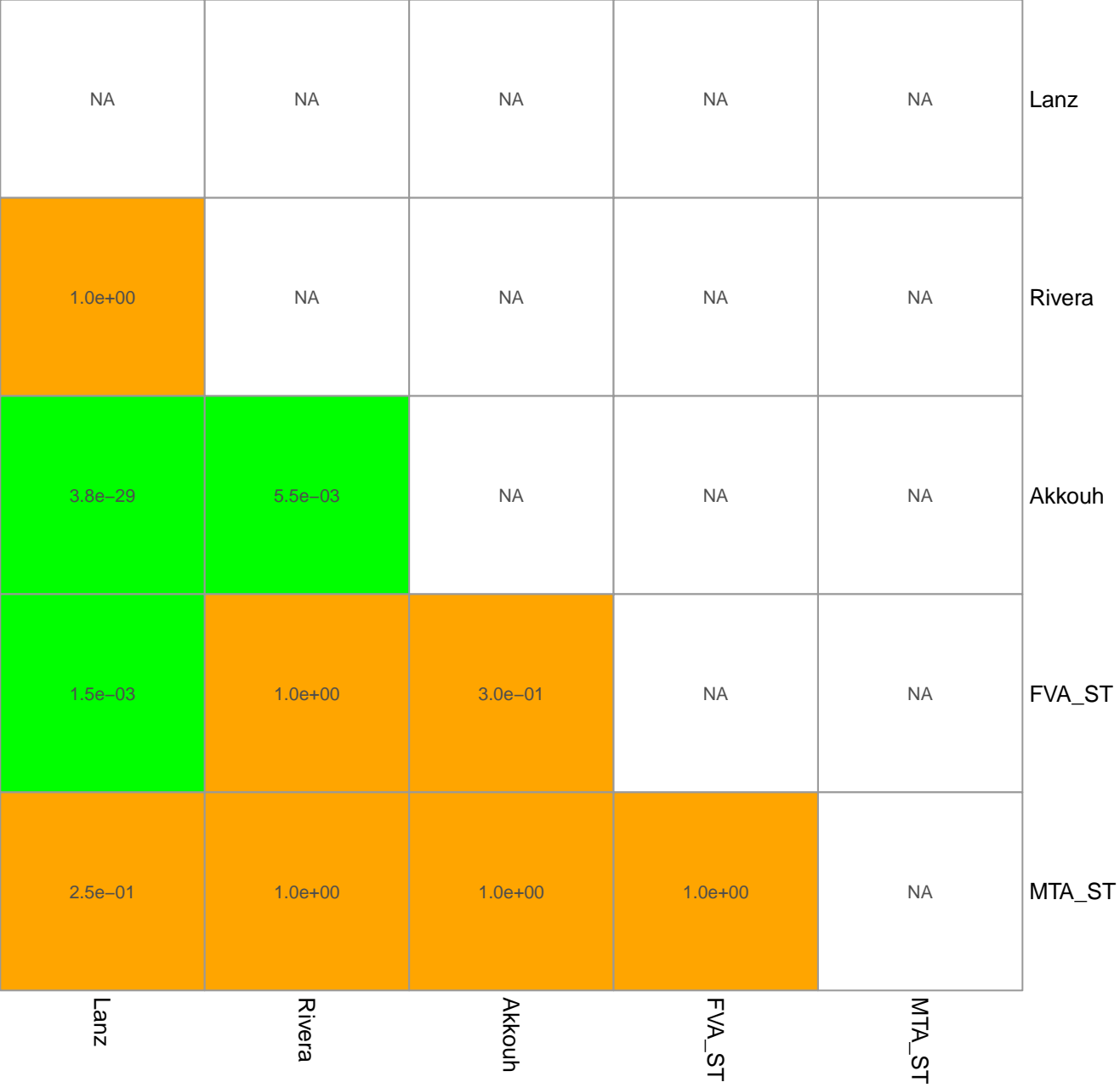


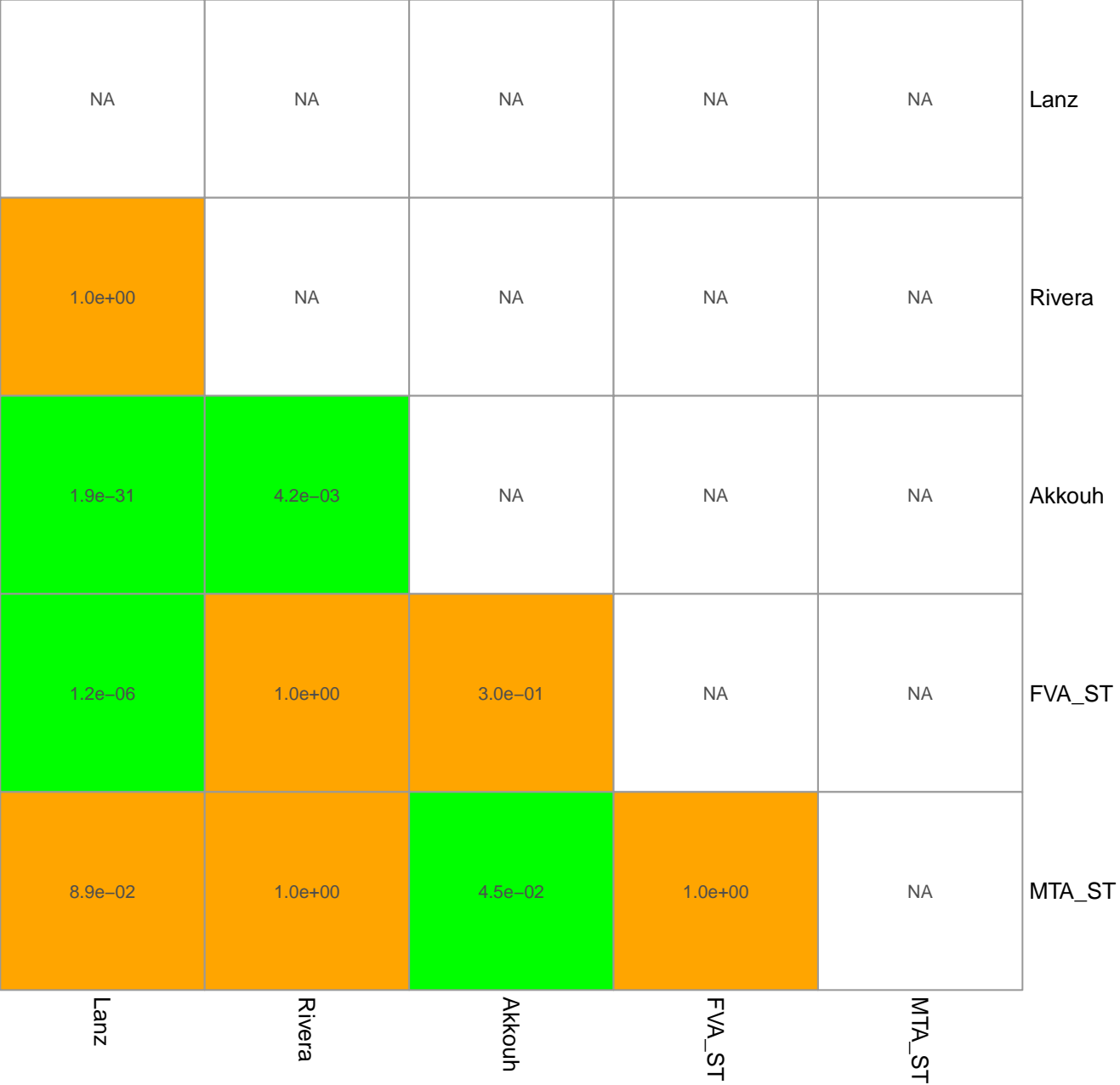
Abs - ST vs Li+



Norm_t1 - ST vs Li+



Norm_t2 - ST vs Li+



Abs

	1	0
	1	0
	0	1
	0	1
	0	1
	0	1

FVA_ST

MTA_ST

Glyoxylate and dicarboxylate metabolism

Methionine and cysteine metabolism

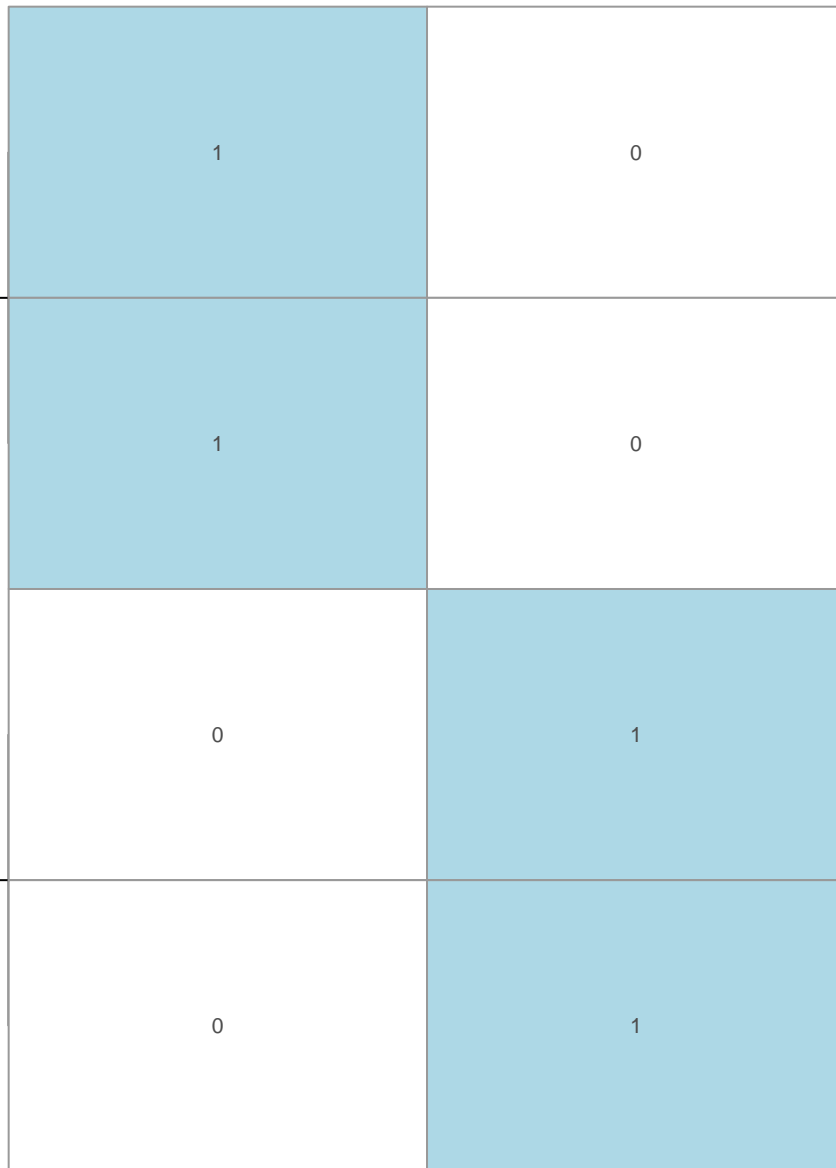
Nucleotide interconversion

Linoleate metabolism

Chondroitin sulfate degradation

Chondroitin synthesis

Norm_t1



1

0

Fatty acid oxidation

1

0

Pentose phosphate pathway

0

1

Nucleotide interconversion

0

1

Nucleotide salvage pathway

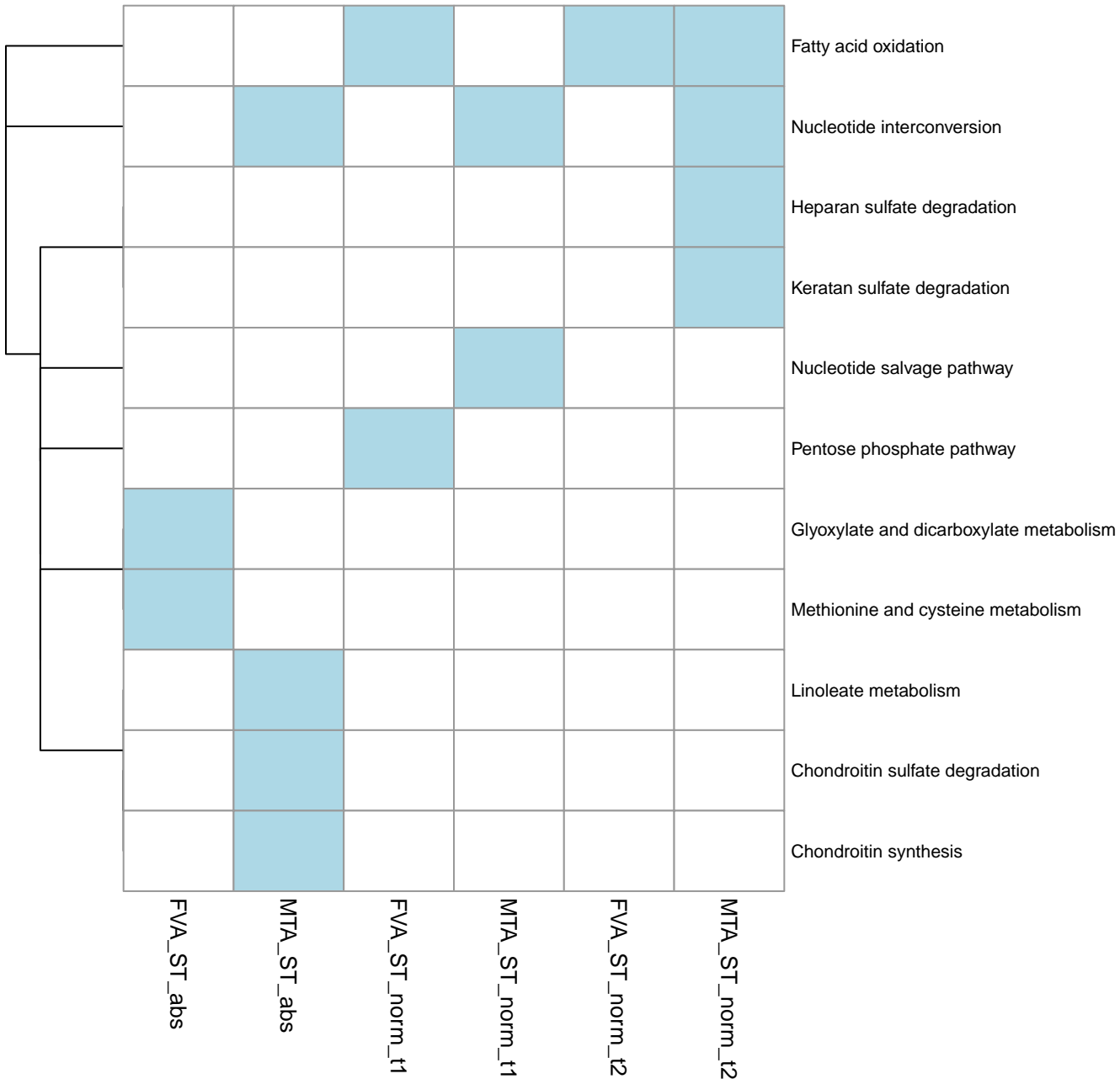
FVA_ST

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



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



