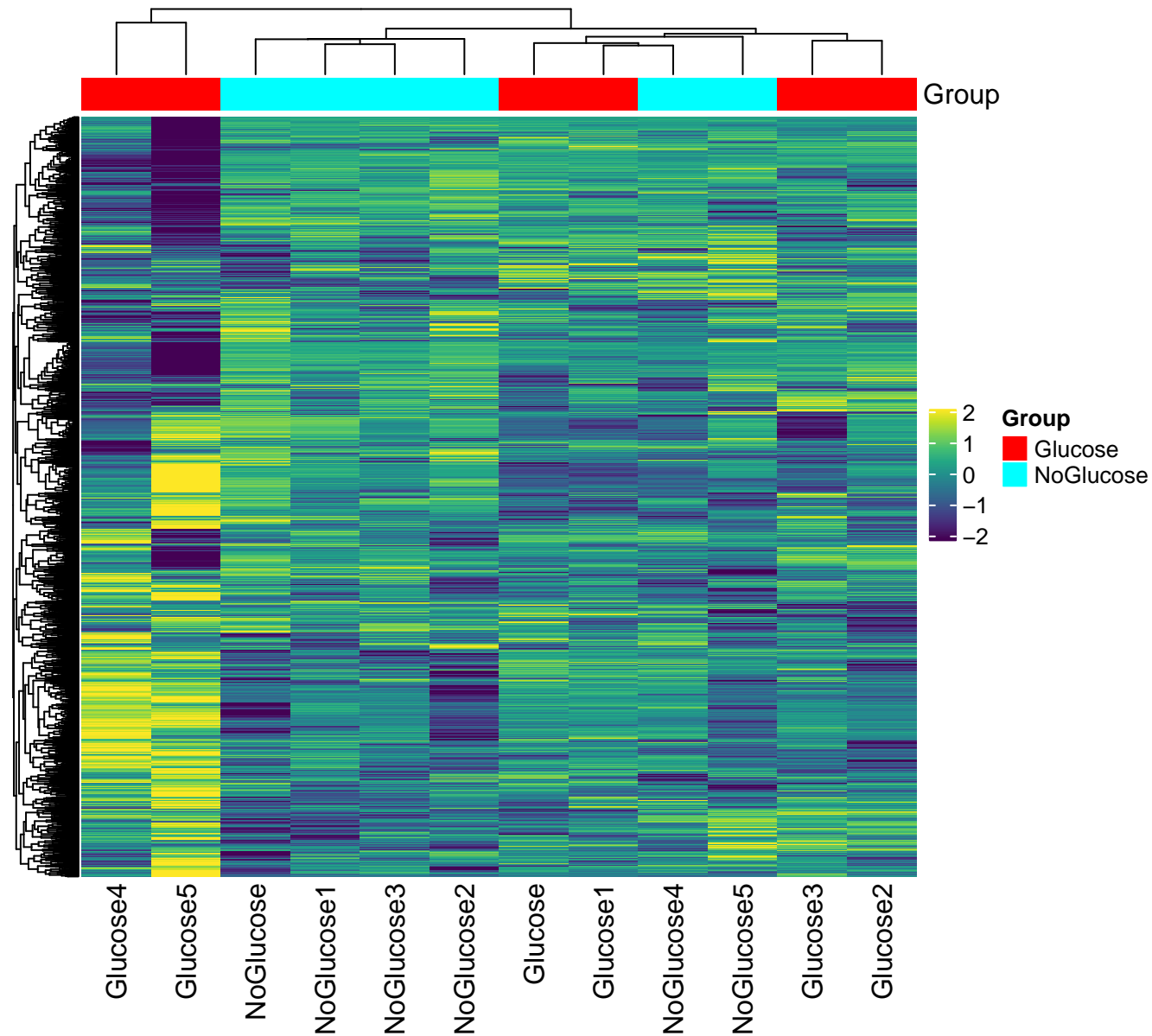
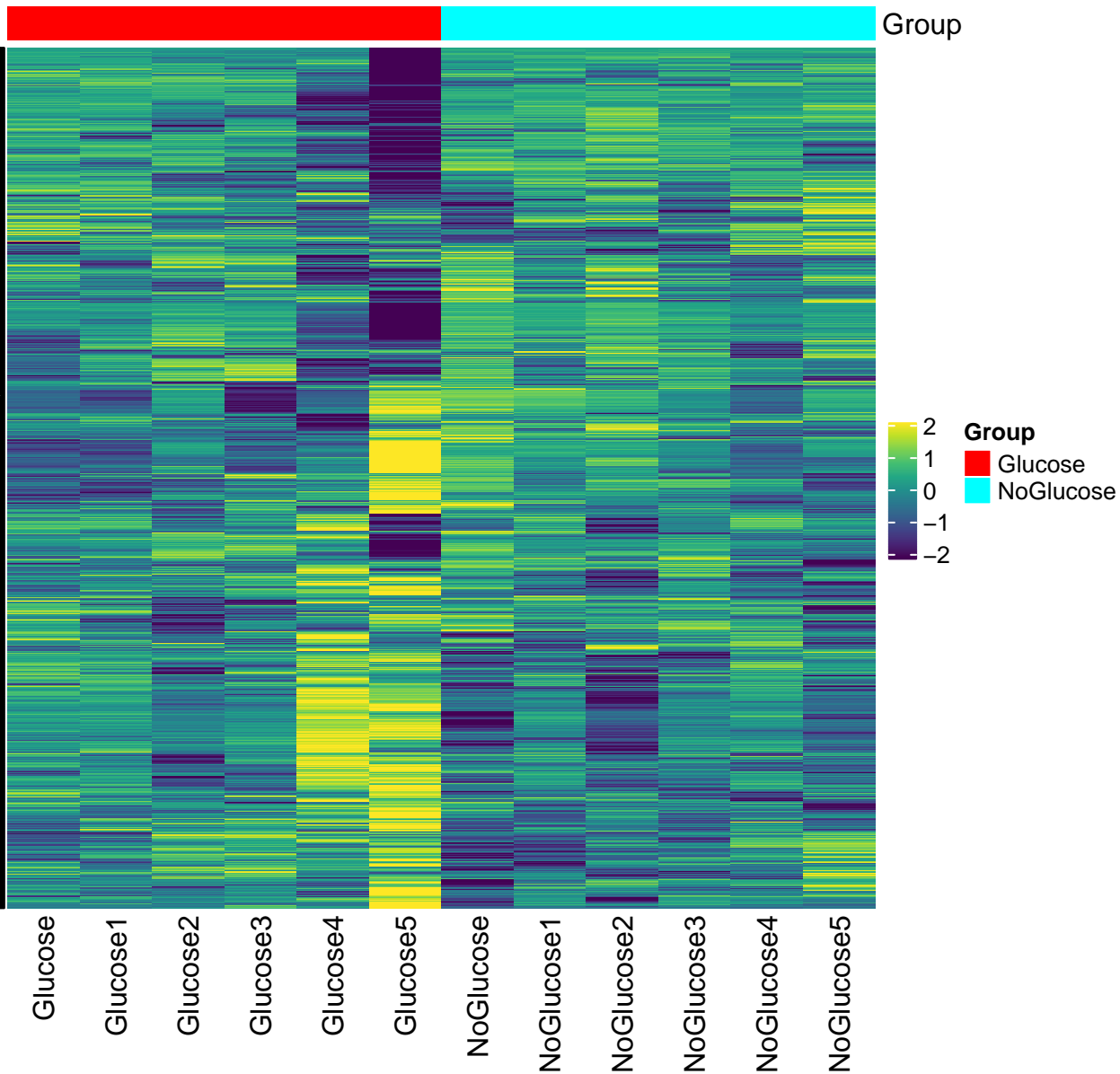


Met_Neg:
All features, row z score



Met_Neg:
All features, row z score



Met_Neg: human1_gene1_expanded

Subset, row z score

Group

Glucose
NoGlucose

Glucose

Glucose1

Glucose2

Glucose3

Glucose4

Glucose5

NoGlucose

NoGlucose1

NoGlucose2

NoGlucose3

NoGlucose4

NoGlucose5

Met_Neg: human1_gene1_expanded

Subset, row z score

Group

Glucose
NoGlucose

Glucose

Glucose1

Glucose2

Glucose3

Glucose4

Glucose5

NoGlucose

NoGlucose1

NoGlucose2

NoGlucose3

NoGlucose4

NoGlucose5

Met_Neg: human1_gene1_expanded

Subset, row z score

Group

Glucose
NoGlucose

Glucose

Glucose1

Glucose2

Glucose3

Glucose4

Glucose5

NoGlucose

NoGlucose1

NoGlucose2

NoGlucose3

NoGlucose4

NoGlucose5

Met_Neg: human1_gene1_expanded

Subset, row z score

Group

Glucose
NoGlucose

Glucose

Glucose1

Glucose2

Glucose3

Glucose4

Glucose5

NoGlucose

NoGlucose1

NoGlucose2

NoGlucose3

NoGlucose4

NoGlucose5

Met_Neg: human2_gene1_expanded

Subset, row z score

Group

Glucose
NoGlucose

Glucose

Glucose1

Glucose2

Glucose3

Glucose4

Glucose5

NoGlucose

NoGlucose1

NoGlucose2

NoGlucose3

NoGlucose4

NoGlucose5

Met_Neg: human2_gene1_expanded

Subset, row z score

Group

Glucose
NoGlucose

Glucose

Glucose1

Glucose2

Glucose3

Glucose4

Glucose5

NoGlucose

NoGlucose1

NoGlucose2

NoGlucose3

NoGlucose4

NoGlucose5

Met_Neg: human2_gene1_expanded

Subset, row z score

Group

Glucose
NoGlucose

Glucose

Glucose1

Glucose2

Glucose3

Glucose4

Glucose5

NoGlucose

NoGlucose1

NoGlucose2

NoGlucose3

NoGlucose4

NoGlucose5

Met_Neg: human2_gene1_expanded

Subset, row z score

Group

Glucose
NoGlucose

Glucose

Glucose1

Glucose2

Glucose3

Glucose4

Glucose5

NoGlucose

NoGlucose1

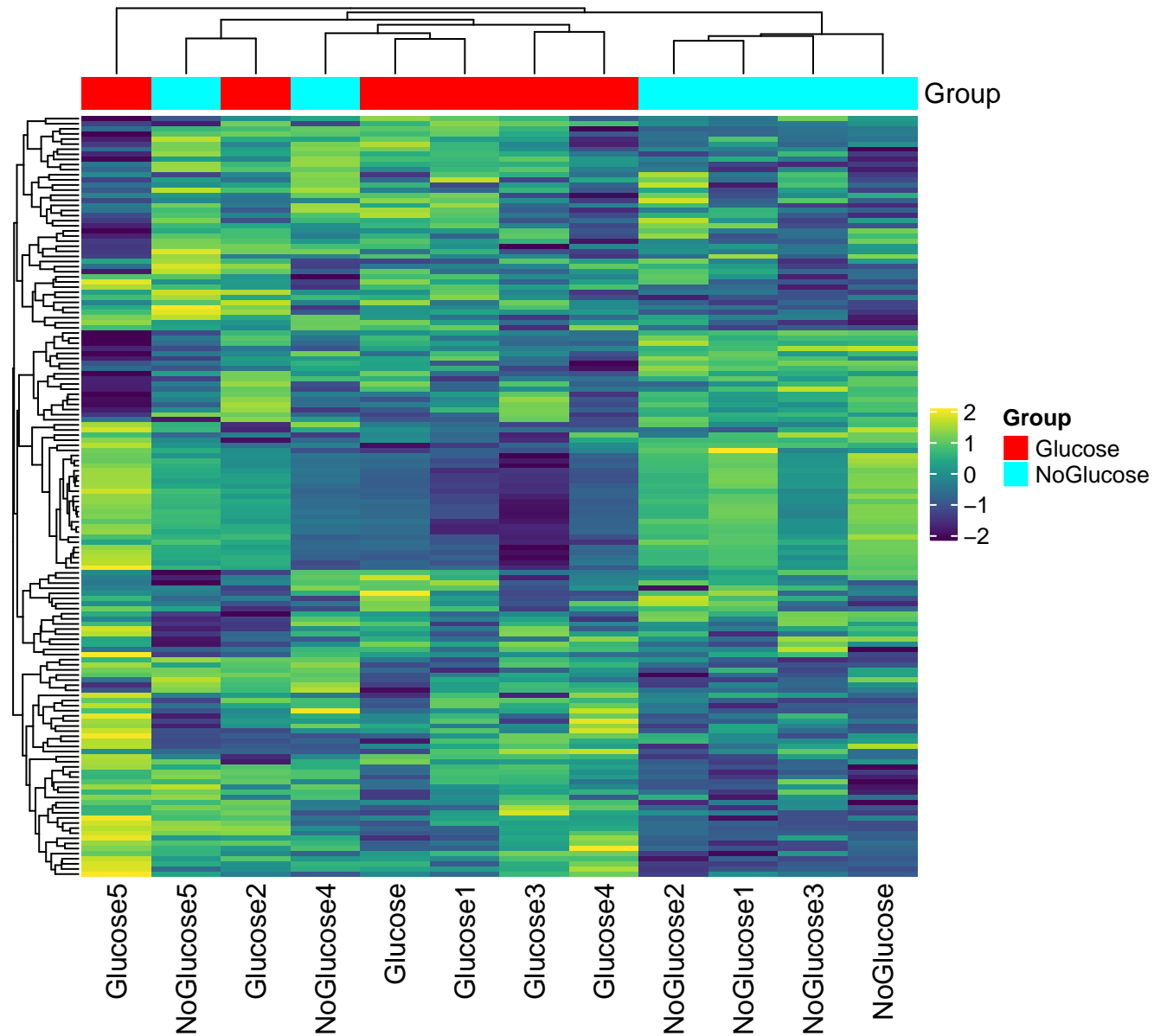
NoGlucose2

NoGlucose3

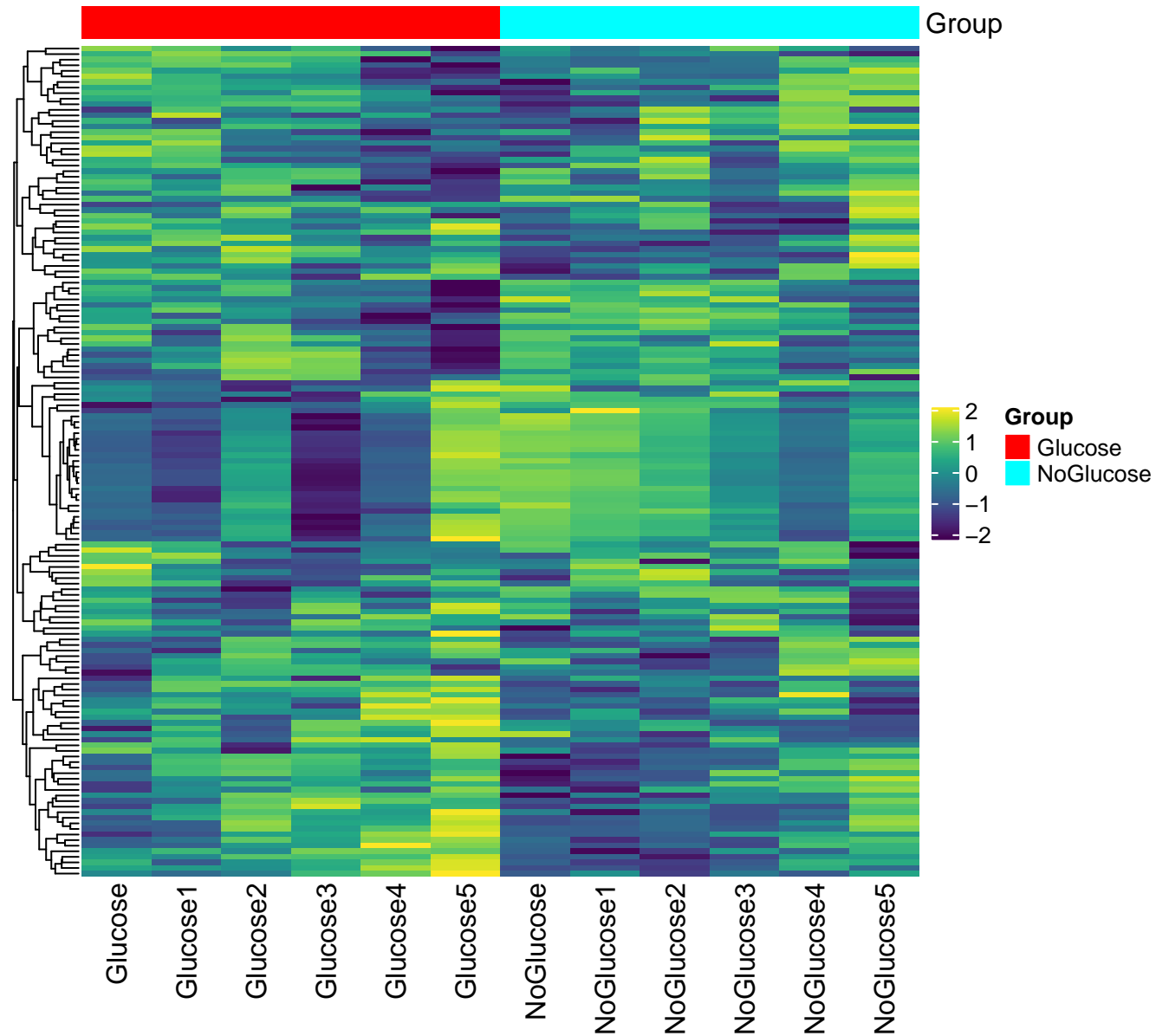
NoGlucose4

NoGlucose5

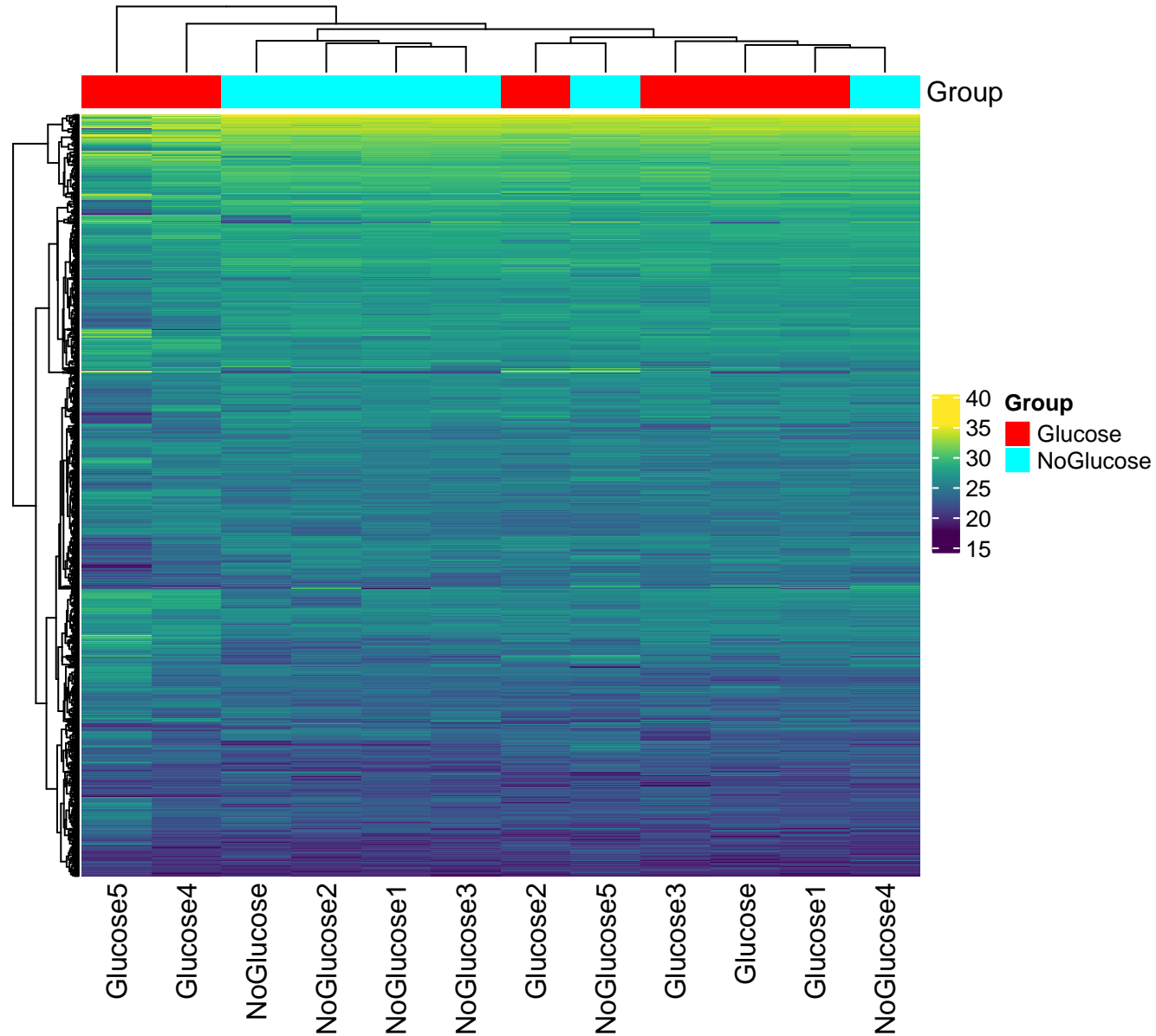
Met_Neg:
Highest variation, row z score



Met_Neg:
Highest variation, row z score



Met_Neg:
All features, log2 Intensity



Met_Neg:
All features, log2 Intensity

Group

