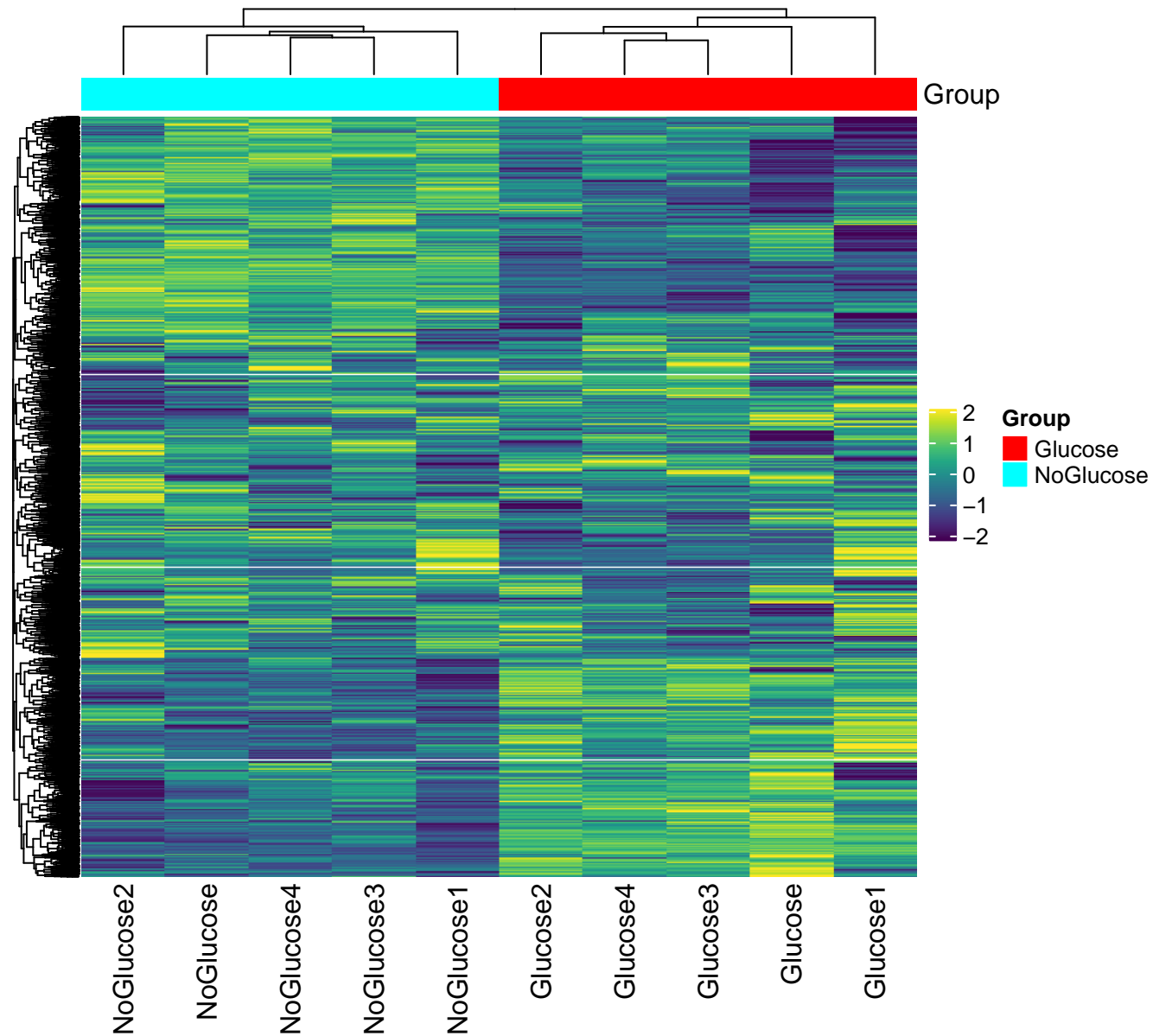
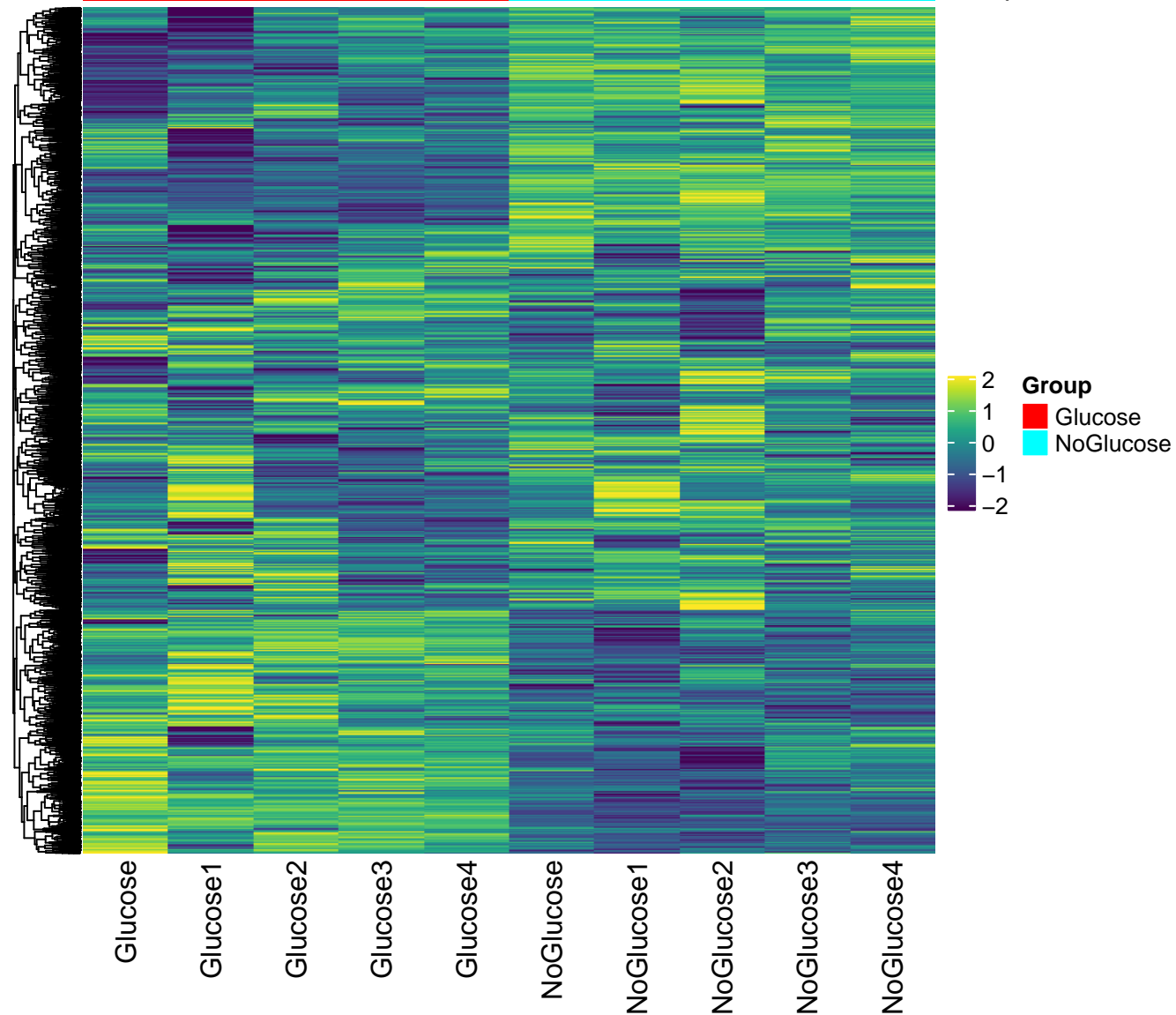


Phospho:
All features, row z score



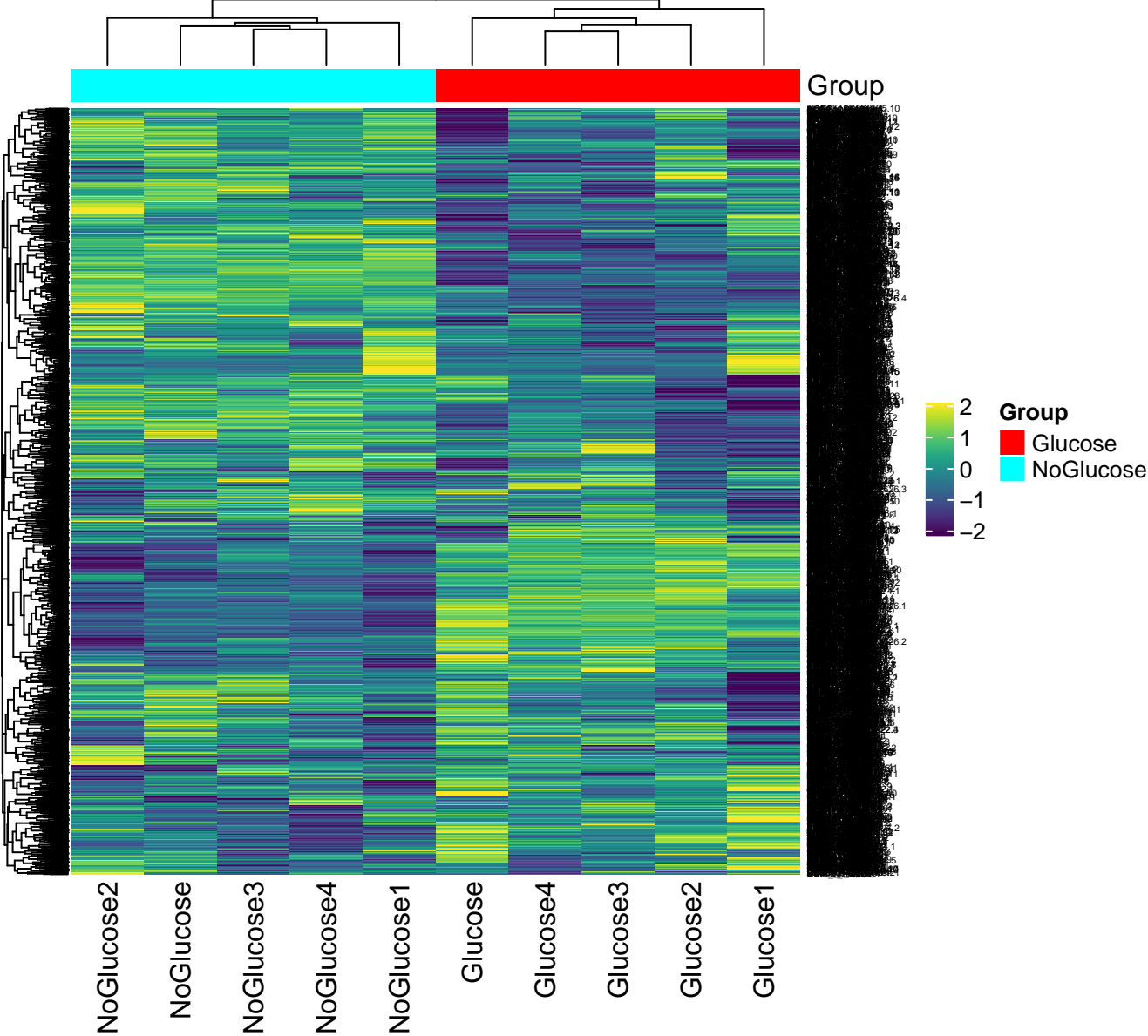
All features, row z score

Group



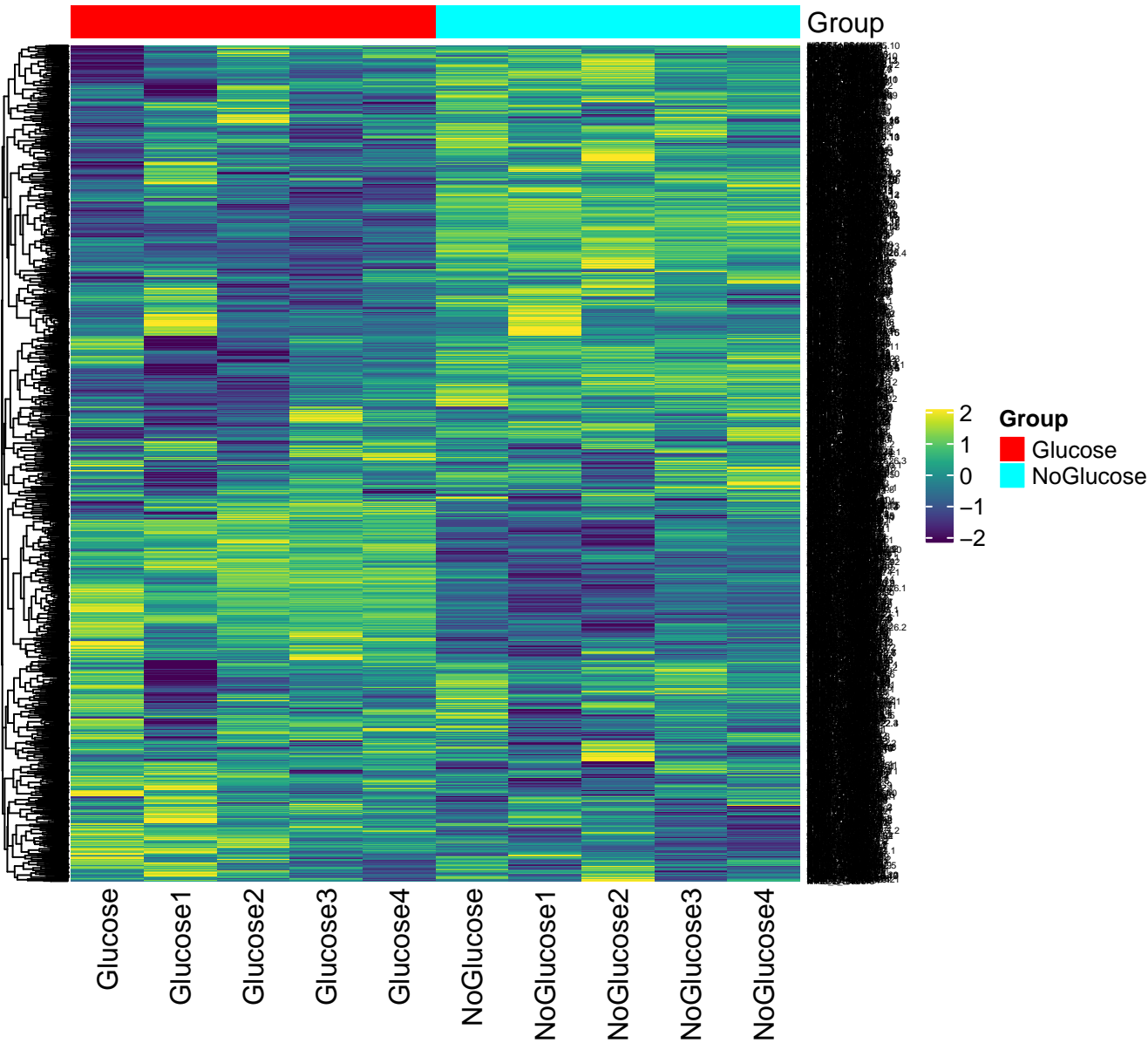
Phospho: human1_gene1_expanded

Subset, row z score



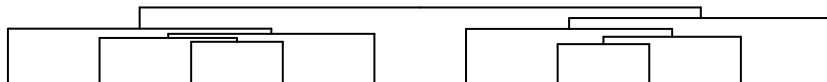
Phospho: human1_gene1_expanded

Subset, row z score

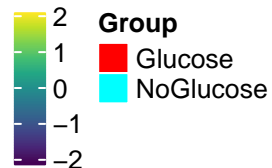


Phospho: human1_gene1_expanded

Subset, row z score



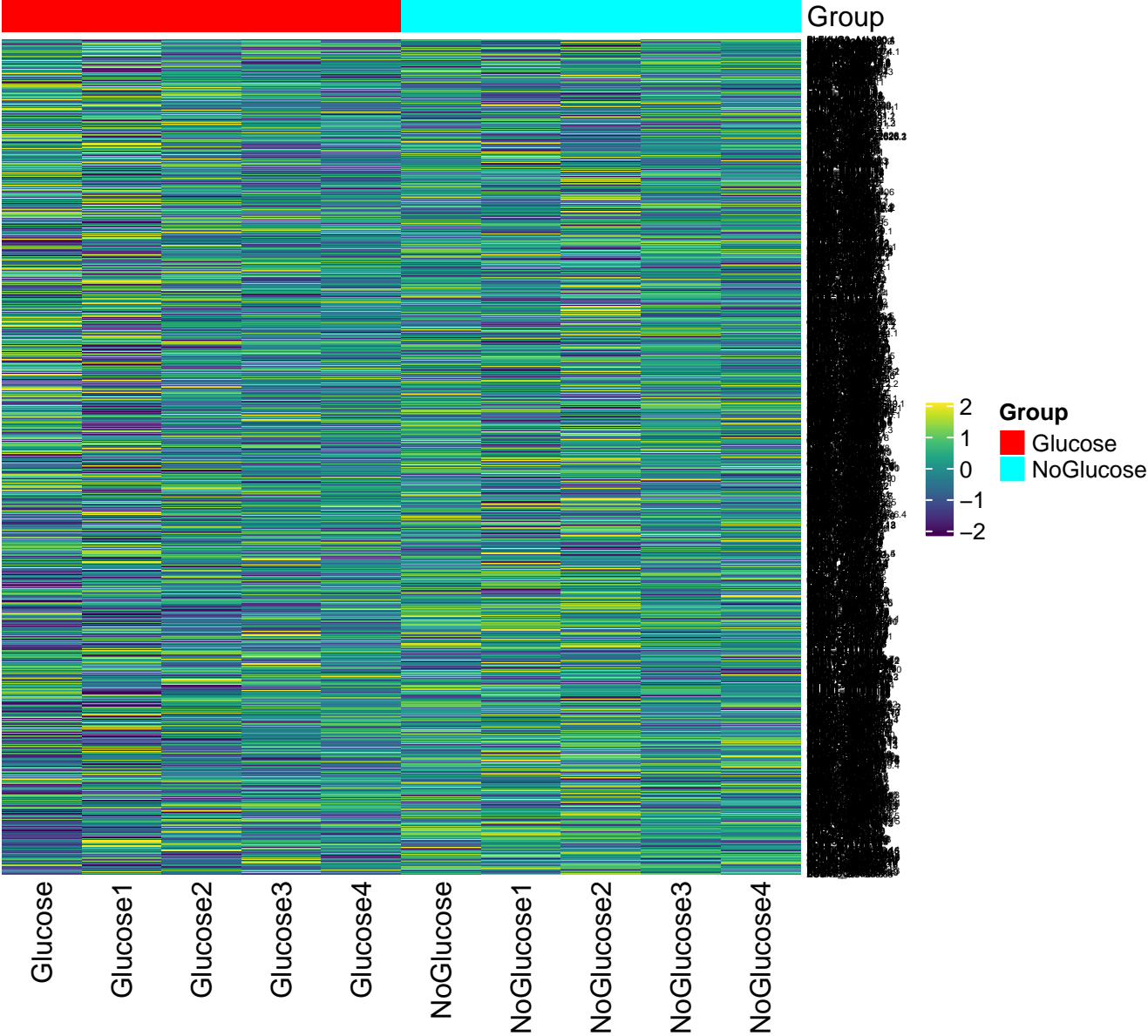
Group



NoGlucose2
NoGlucose
NoGlucose3
NoGlucose4
NoGlucose1
Glucose
Glucose4
Glucose3
Glucose2
Glucose1

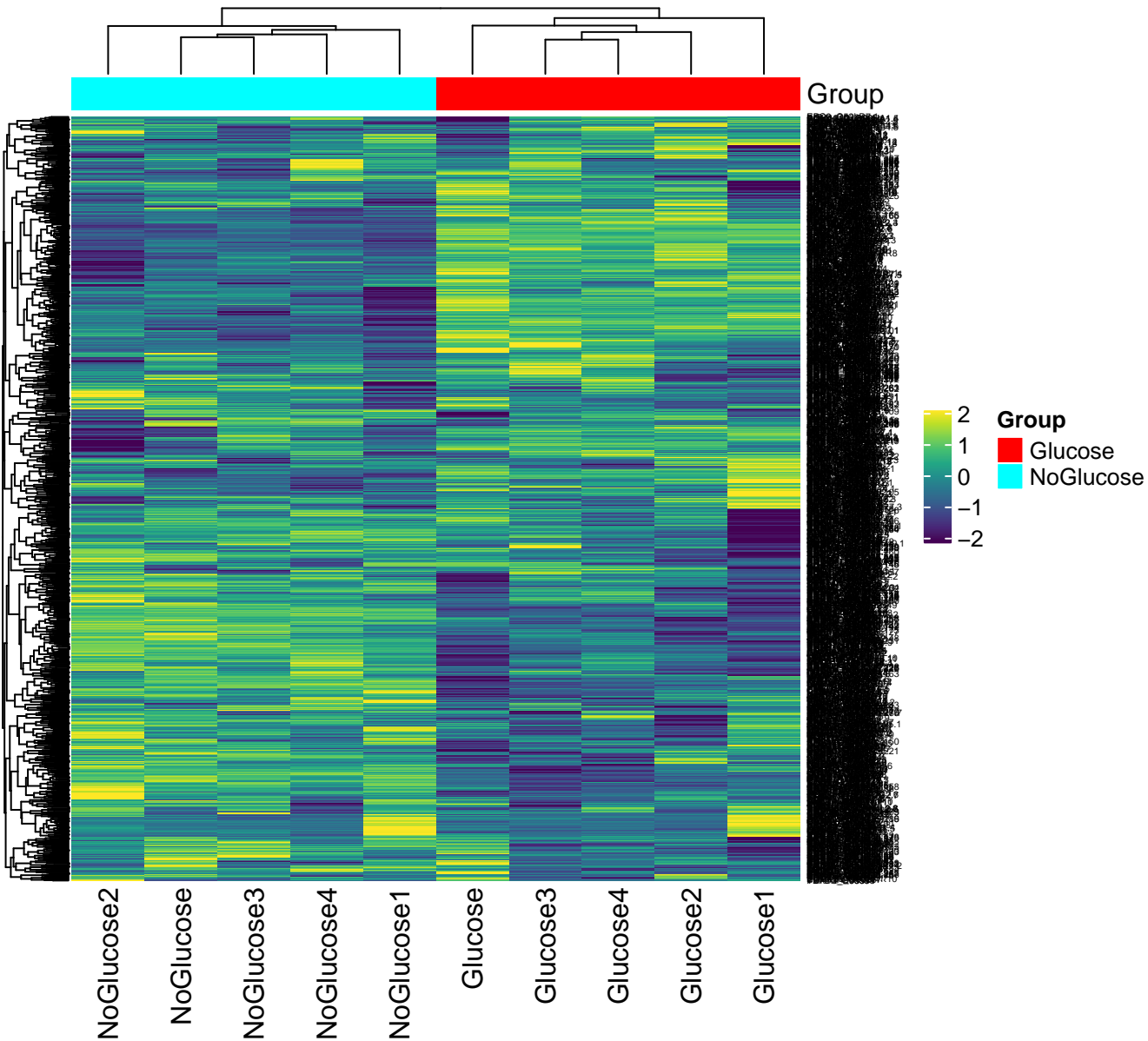
Phospho: human1_gene1_expanded

Subset, row z score



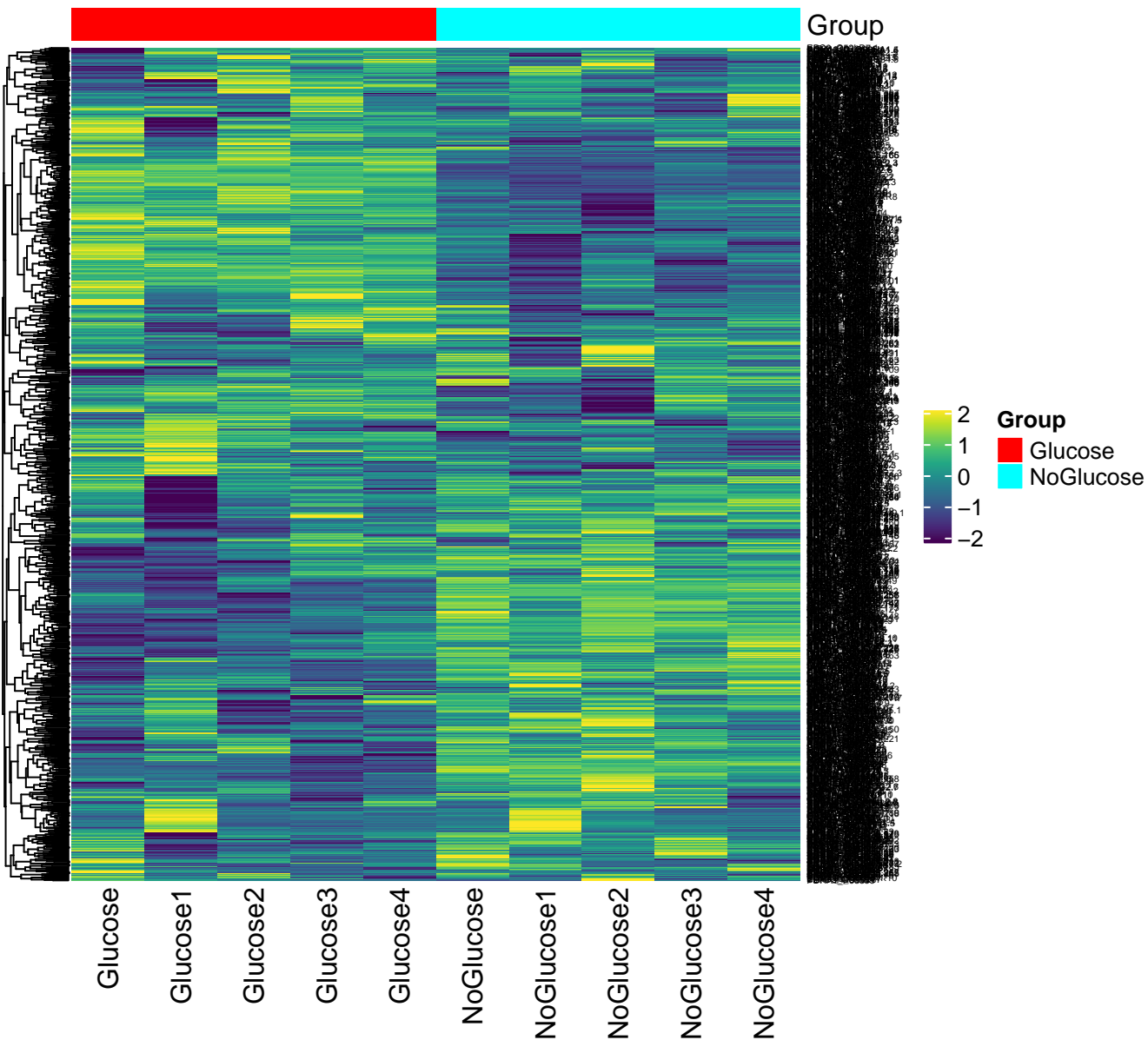
Phospho: human2_gene1_expanded

Subset, row z score



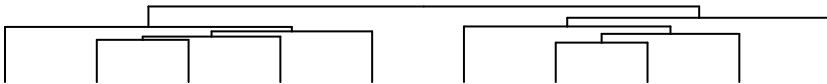
Phospho: human2_gene1_expanded

Subset, row z score

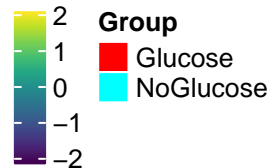
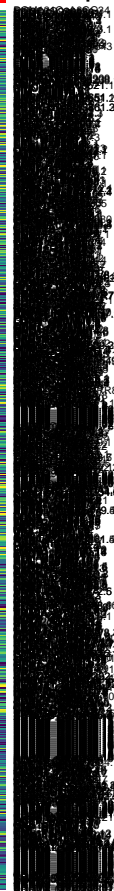


Phospho: human2_gene1_expanded

Subset, row z score



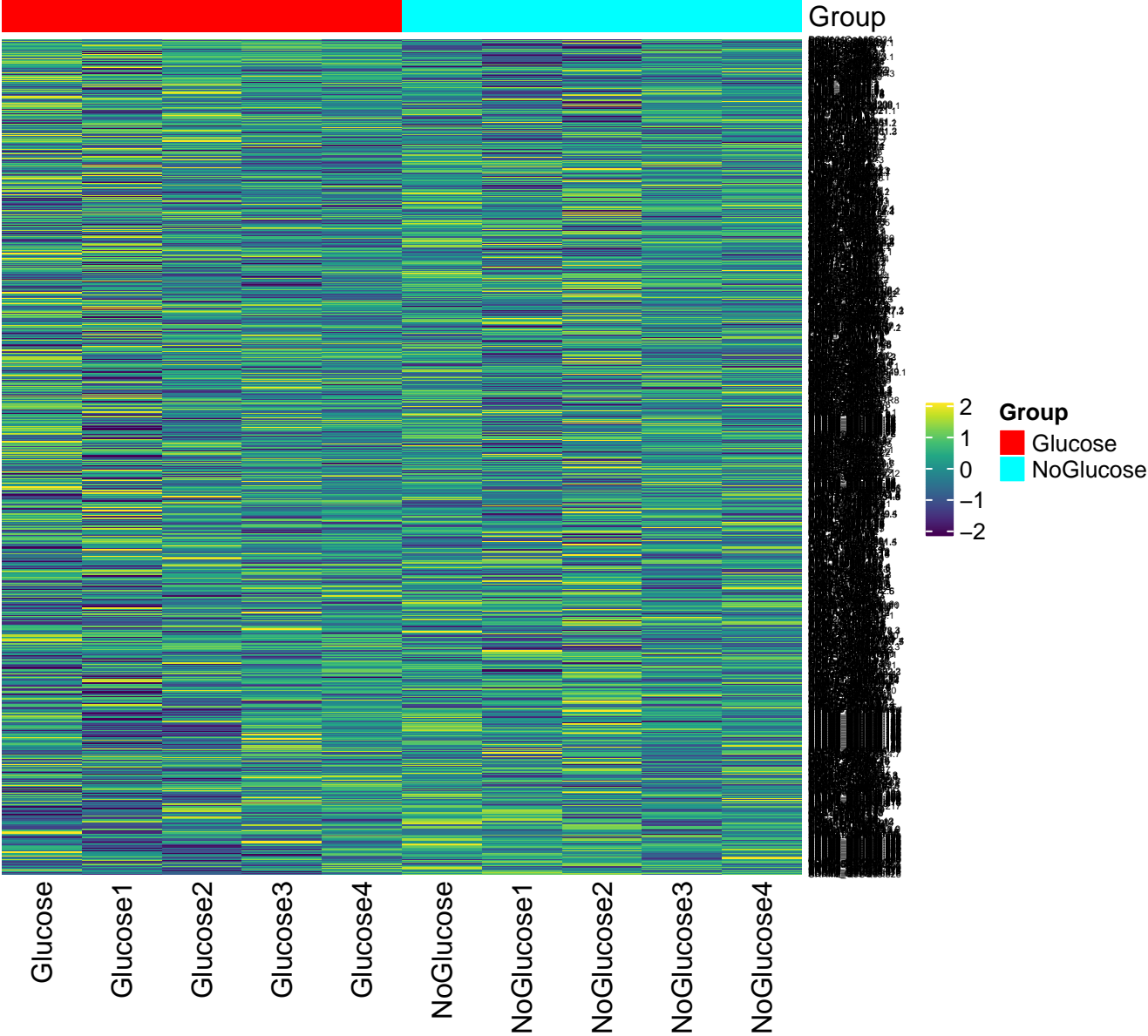
Group



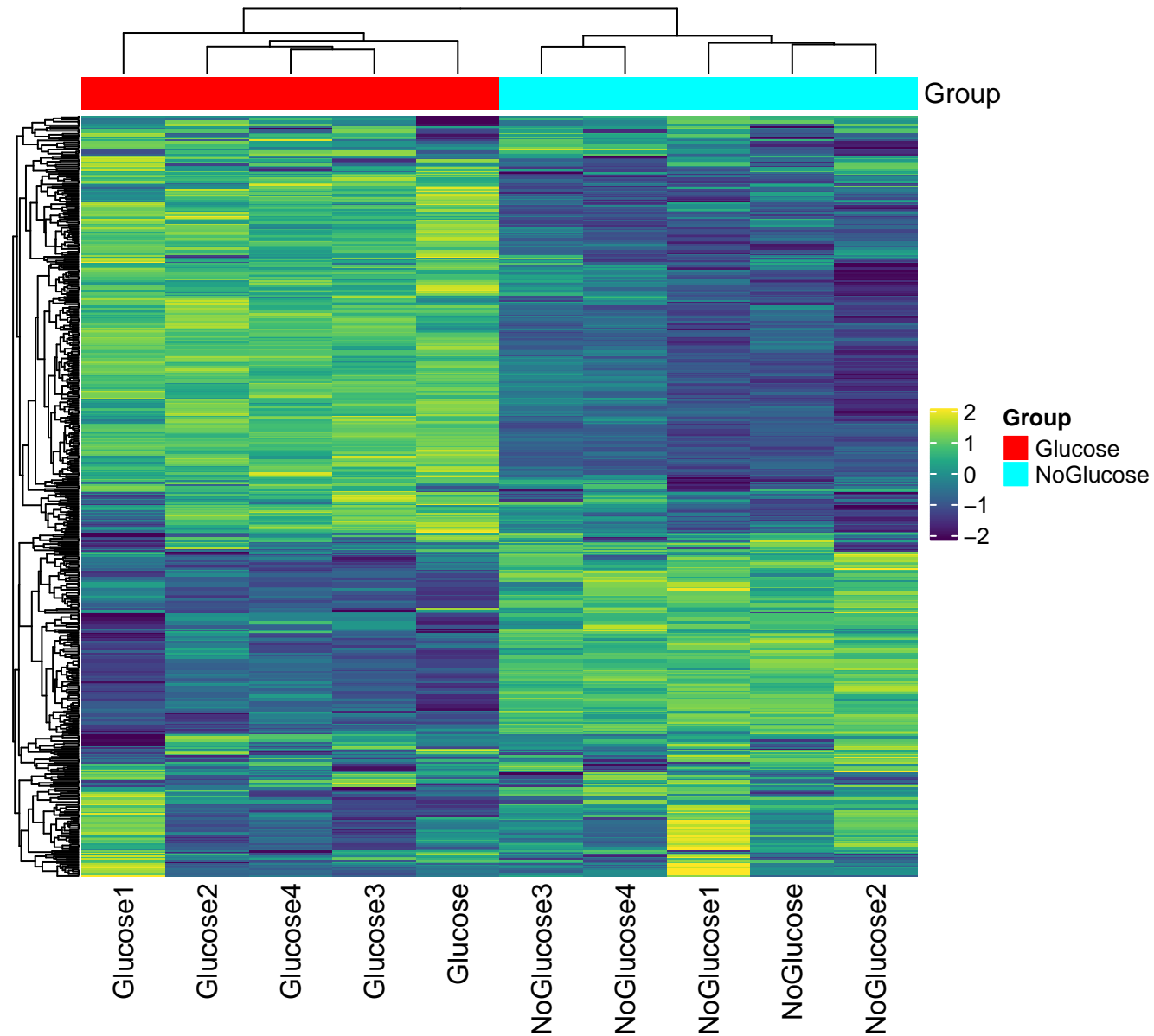
NoGlucose2	NoGlucose3	NoGlucose4	NoGlucose1	Glucose	Glucose3	Glucose4	Glucose2	Glucose1
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Phospho: human2_gene1_expanded

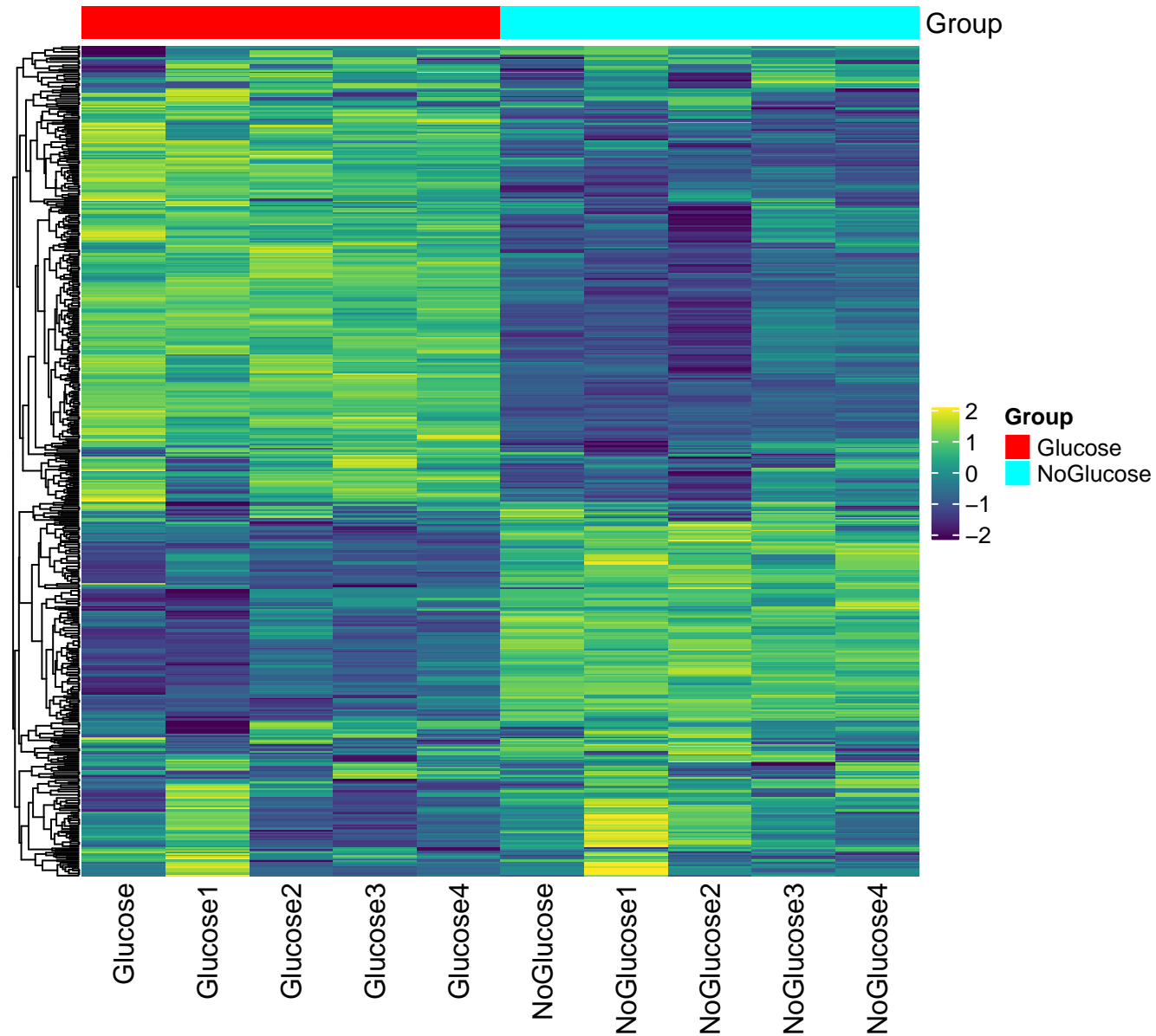
Subset, row z score



Phospho:
Highest variation, row z score

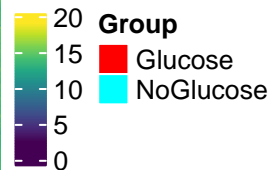


Phospho:
Highest variation, row z score



Phospho:
All features, log2 Intensity

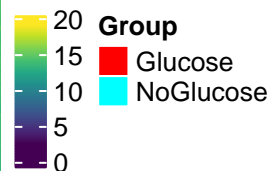
Group



Glucose1
Glucose
Glucose2
Glucose4
Glucose3
NoGlucose4
NoGlucose3
NoGlucose
NoGlucose1
NoGlucose2

Phospho:
All features, log₂ Intensity

Group



Glucose

Glucose1

Glucose2

Glucose3

Glucose4

NoGlucose

NoGlucose1

NoGlucose2

NoGlucose3

NoGlucose4