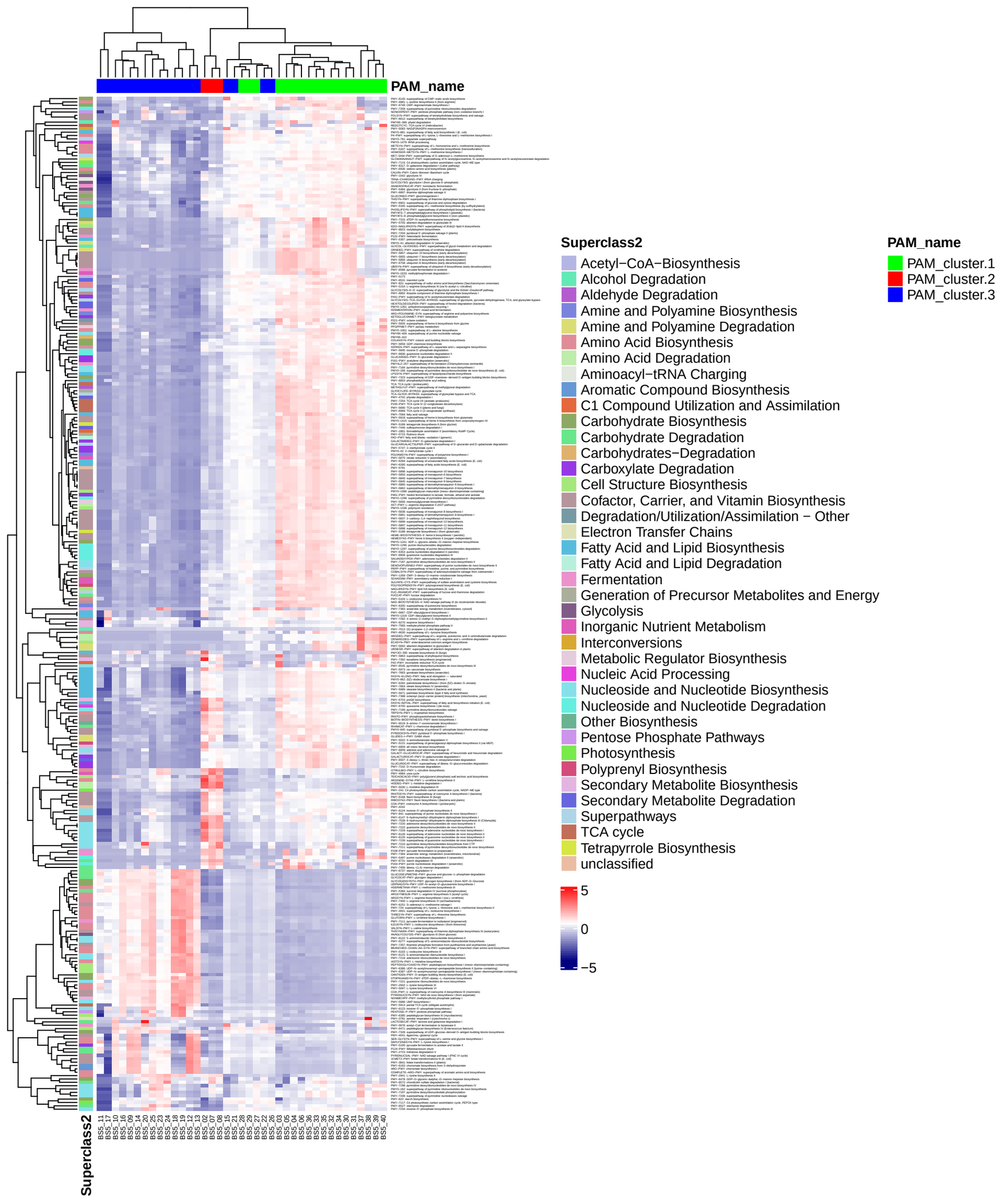
PAM_name
Superclass2
BS5_11
BS5_17
BS5_10
BS5_16
BS5_09
BS5_14
BS5_20
BS5_25
BS5_23
BS5_24
BS5_18
BS5_19
BS5_12
BS5_13
BS5_02
BS5_07
BS5_08
BS5_15
BS5_21
BS5_28
BS5_29
BS5_27
BS5_22
BS5_26
BS5_03
BS5_05
BS5_04
BS5_06
BS5_36
BS5_33
BS5_35
BS5_32
BS5_34
BS5_30
BS5_31
BS5_37
BS5_38
BS5_39
BS5_40
Superclass2
Acetyl−CoA−Biosynthesis
Alcohol Degradation
Aldehyde Degradation
Amine and Polyamine Biosynthesis
Amine and Polyamine Degradation
Amino Acid Biosynthesis
Amino Acid Degradation
Aminoacyl−tRNA Charging
Aromatic Compound Biosynthesis
C1 Compound Utilization and Assimilation
Carbohydrate Biosynthesis
Carbohydrate Degradation
Carbohydrates−Degradation
Carboxylate Degradation
Cell Structure Biosynthesis
Cofactor, Carrier, and Vitamin Biosynthesis
Degradation/Utilization/Assimilation − Other
Electron Transfer Chains
Fatty Acid and Lipid Biosynthesis
Fatty Acid and Lipid Degradation
Fermentation
Generation of Precursor Metabolites and Energy
Glycolysis
Inorganic Nutrient Metabolism
Interconversions
Metabolic Regulator Biosynthesis
Nucleic Acid Processing
Nucleoside and Nucleotide Biosynthesis
Nucleoside and Nucleotide Degradation
Other Biosynthesis
Pentose Phosphate Pathways
Photosynthesis
Polyprenyl Biosynthesis
Secondary Metabolite Biosynthesis
Secondary Metabolite Degradation
Superpathways
TCA cycle
Tetrapyrrole Biosynthesis
unclassified
5
0
−5
PAM_name
PAM_cluster.1
PAM_cluster.2
PAM_cluster.3