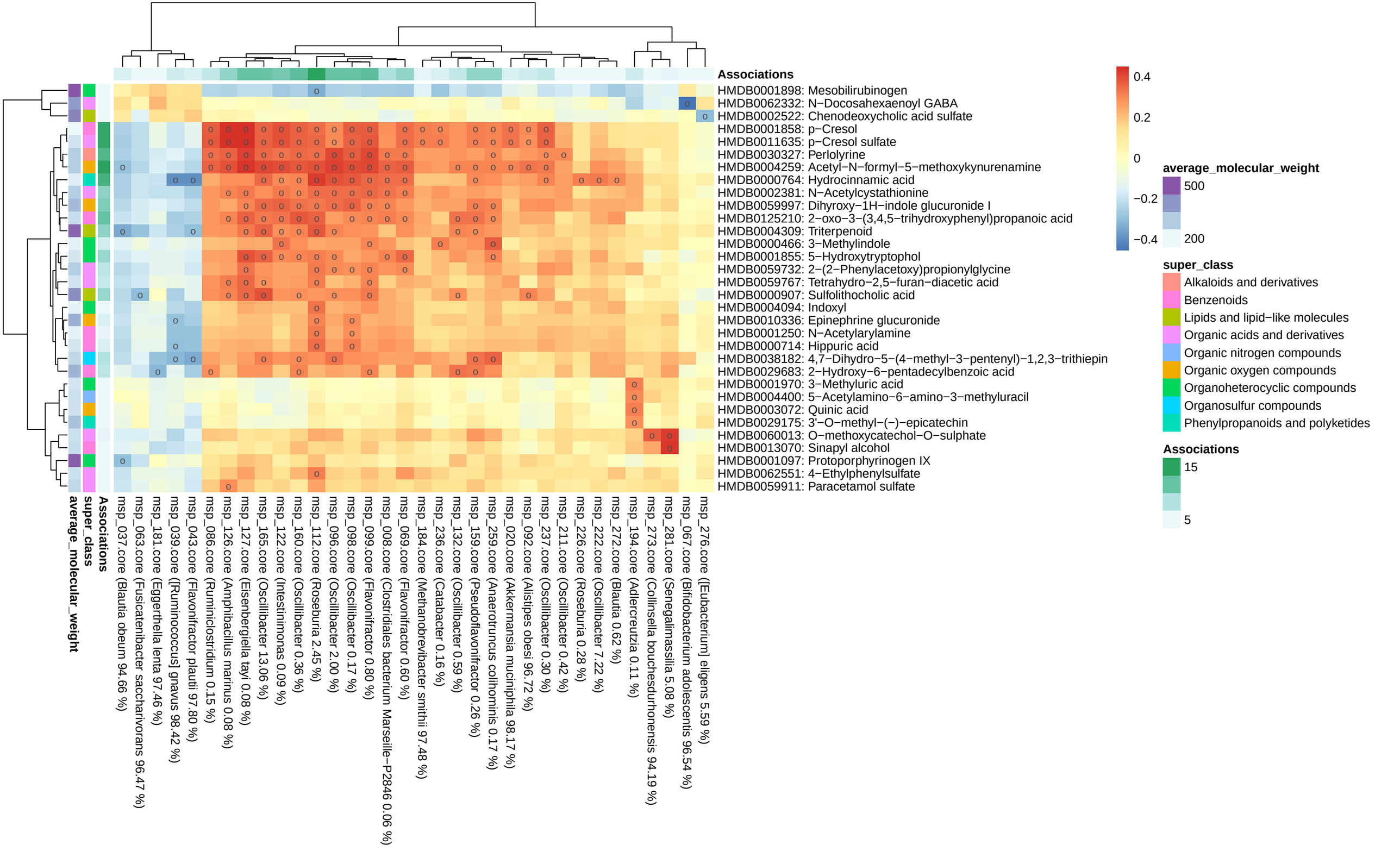
Associations
HMDB0001898: Mesobilirubinogen
HMDB0062332: N−Docosahexaenoyl GABA
HMDB0002522: Chenodeoxycholic acid sulfate
HMDB0001858: p−Cresol
HMDB0011635: p−Cresol sulfate
HMDB0030327: Perlolyrine
HMDB0004259: Acetyl−N−formyl−5−methoxykynurenamine
HMDB0000764: Hydrocinnamic acid
HMDB0002381: N−Acetylcystathionine
HMDB0059997: Dihyroxy−1H−indole glucuronide I
HMDB0125210: 2−oxo−3−(3,4,5−trihydroxyphenyl)propanoic acid
HMDB0004309: Triterpenoid
HMDB0000466: 3−Methylindole
HMDB0001855: 5−Hydroxytryptophol
HMDB0059732: 2−(2−Phenylacetoxy)propionylglycine
HMDB0059767: Tetrahydro−2,5−furan−diacetic acid
HMDB0000907: Sulfolithocholic acid
HMDB0004094: Indoxyl
HMDB0010336: Epinephrine glucuronide
HMDB0001250: N−Acetylarylamine
HMDB0000714: Hippuric acid
HMDB0038182: 4,7−Dihydro−5−(4−methyl−3−pentenyl)−1,2,3−trithiepin
HMDB0029683: 2−Hydroxy−6−pentadecylbenzoic acid
HMDB0001970: 3−Methyluric acid
HMDB0004400: 5−Acetylamino−6−amino−3−methyluracil
HMDB0003072: Quinic acid
HMDB0029175: 3'−O−methyl−(−)−epicatechin
HMDB0060013: O−methoxycatechol−O−sulphate
HMDB0013070: Sinapyl alcohol
HMDB0001097: Protoporphyrinogen IX
HMDB0062551: 4−Ethylphenylsulfate
HMDB0059911: Paracetamol sulfate
msp_276.core ([Eubacterium] eligens 5.59 %)
msp_067.core (Bifidobacterium adolescentis 96.54 %)
msp_281.core (Senegalimassilia 5.08 %)
msp_273.core (Collinsella bouchesdurhonensis 94.19 %)
msp_194.core (Adlercreutzia 0.11 %)
msp_272.core (Blautia 0.62 %)
msp_222.core (Oscillibacter 7.22 %)
msp_226.core (Roseburia 0.28 %)
msp_211.core (Oscillibacter 0.42 %)
msp_237.core (Oscillibacter 0.30 %)
msp_092.core (Alistipes obesi 96.72 %)
msp_020.core (Akkermansia muciniphila 98.17 %)
msp_259.core (Anaerotruncus colihominis 0.17 %)
msp_159.core (Pseudoflavonifractor 0.26 %)
msp_132.core (Oscillibacter 0.59 %)
msp_236.core (Catabacter 0.16 %)
msp_184.core (Methanobrevibacter smithii 97.48 %)
msp_069.core (Flavonifractor 0.60 %)
msp_008.core (Clostridiales bacterium Marseille−P2846 0.06 %)
msp_099.core (Flavonifractor 0.80 %)
msp_098.core (Oscillibacter 0.17 %)
msp_096.core (Oscillibacter 2.00 %)
msp_112.core (Roseburia 2.45 %)
msp_160.core (Oscillibacter 0.36 %)
msp_122.core (Intestinimonas 0.09 %)
msp_165.core (Oscillibacter 13.06 %)
msp_127.core (Eisenbergiella tayi 0.08 %)
msp_126.core (Amphibacillus marinus 0.08 %)
msp_086.core (Ruminiclostridium 0.15 %)
msp_043.core (Flavonifractor plautii 97.80 %)
msp_039.core ([Ruminococcus] gnavus 98.42 %)
msp_181.core (Eggerthella lenta 97.46 %)
msp_063.core (Fusicatenibacter saccharivorans 96.47 %)
msp_037.core (Blautia obeum 94.66 %)
Associations
super_class
average_molecular_weight
0.4
0.2
0
−0.2
−0.4
average_molecular_weight
500
200
super_class
Alkaloids and derivatives
Benzenoids
Lipids and lipid−like molecules
Organic acids and derivatives
Organic nitrogen compounds
Organic oxygen compounds
Organoheterocyclic compounds
Organosulfur compounds
Phenylpropanoids and polyketides
Associations
15
5