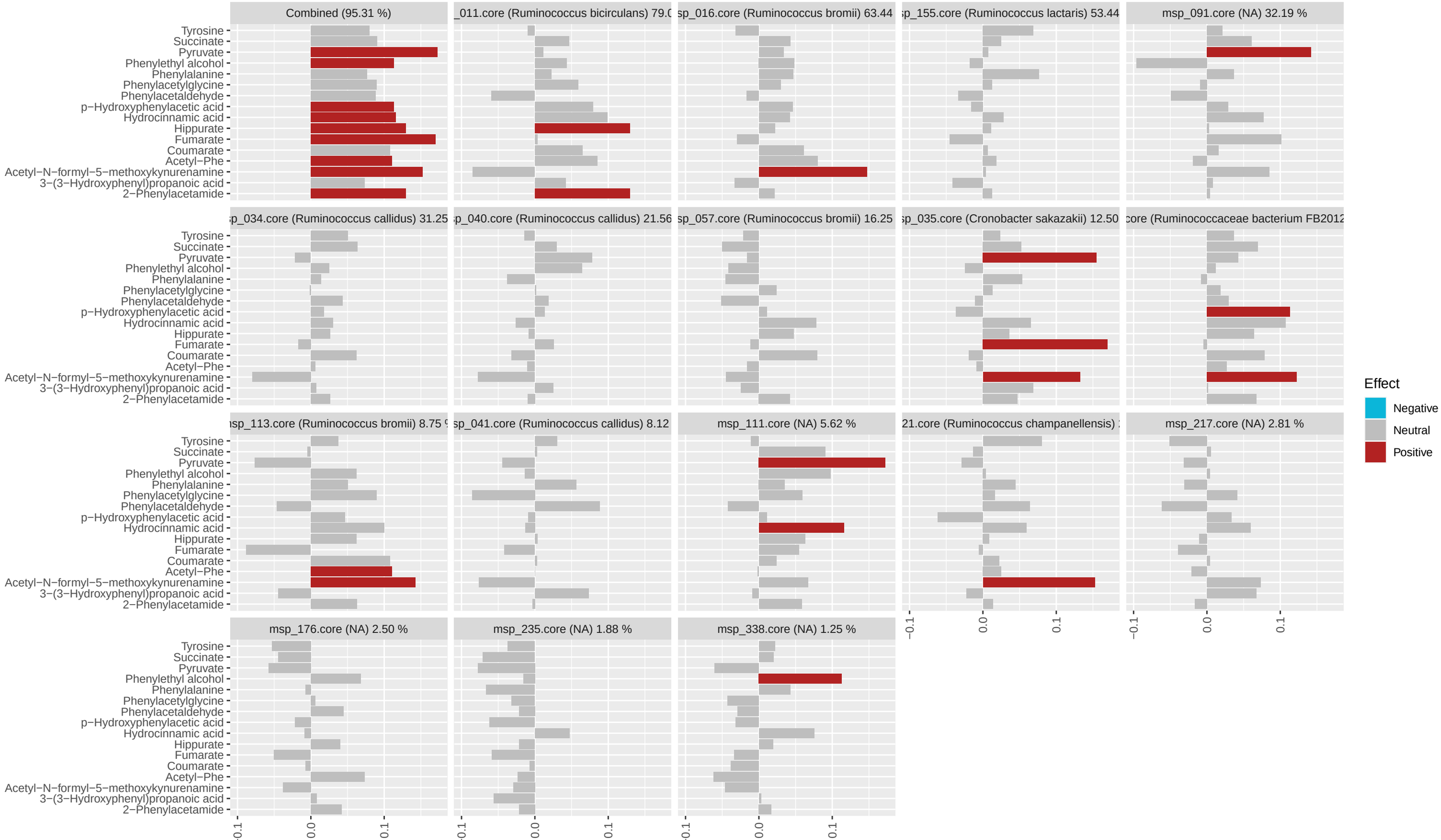
Combined (95.31 %)
_011.core (Ruminococcus bicirculans) 79.0
sp_016.core (Ruminococcus bromii) 63.44
sp_155.core (Ruminococcus lactaris) 53.44
msp_091.core (NA) 32.19 %
sp_034.core (Ruminococcus callidus) 31.25
p_040.core (Ruminococcus callidus) 21.56
sp_057.core (Ruminococcus bromii) 16.25
sp_035.core (Cronobacter sakazakii) 12.50
core (Ruminococcaceae bacterium FB2012
msp_113.core (Ruminococcus bromii) 8.75 %
sp_041.core (Ruminococcus callidus) 8.12
msp_111.core (NA) 5.62 %
21.core (Ruminococcus champanellensis)
msp_217.core (NA) 2.81 %
msp_176.core (NA) 2.50 %
msp_235.core (NA) 1.88 %
msp_338.core (NA) 1.25 %
Tyrosine
Succinate
Pyruvate
Phenylethyl alcohol
Phenylalanine
Phenylacetylglycine
Phenylacetaldehyde
p−Hydroxyphenylacetic acid
Hydrocinnamic acid
Hippurate
Fumarate
Coumarate
Acetyl−Phe
Acetyl−N−formyl−5−methoxykynurenamine
3−(3−Hydroxyphenyl)propanoic acid
2−Phenylacetamide
−0.1
0.0
0.1
Effect
Negative
Neutral
Positive