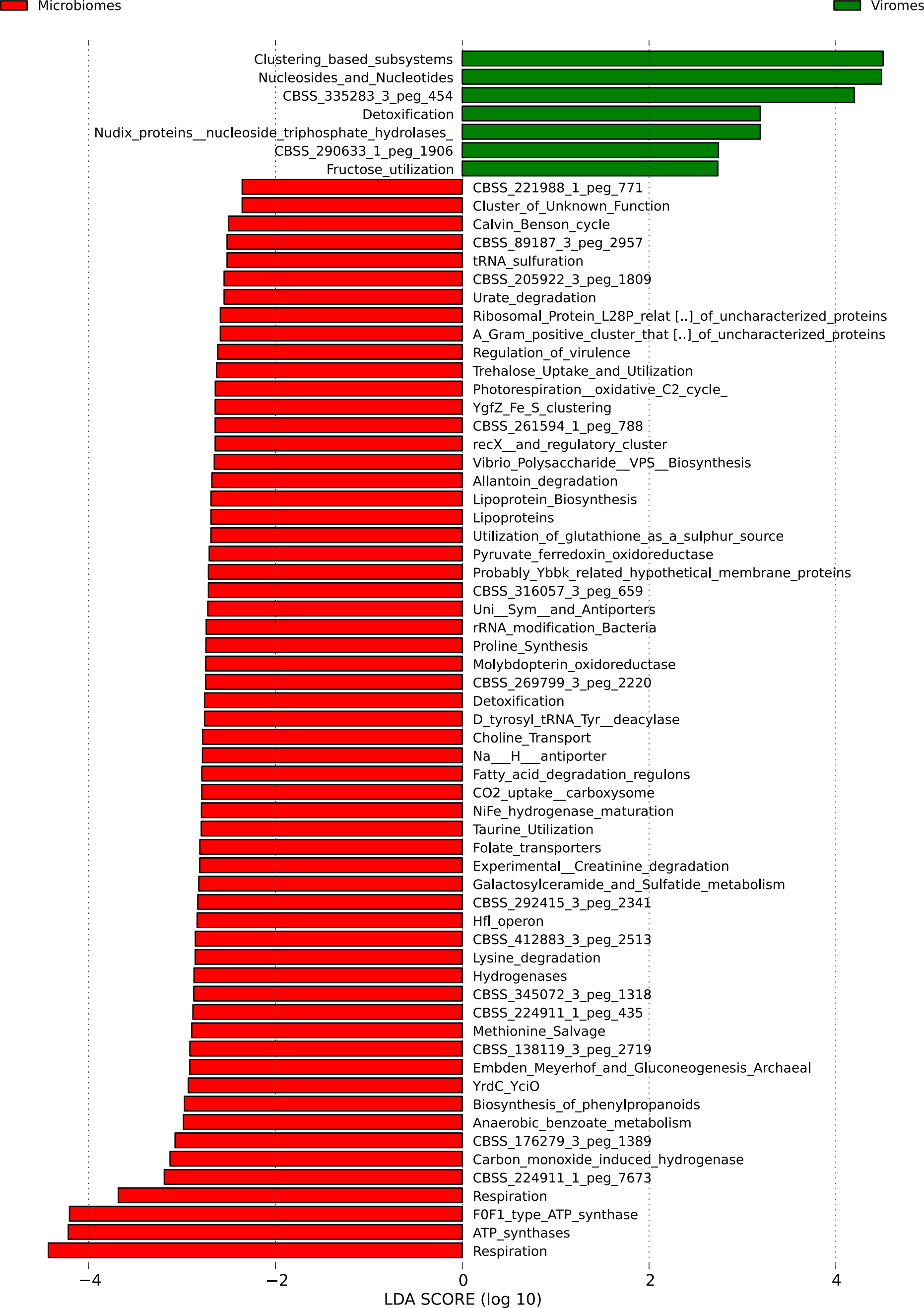
Microbiomes
Viromes
Clustering_based_subsystems
Nucleosides_and_Nucleotides
CBSS_335283_3_peg_454
Detoxification
Nudix_proteins__nucleoside_triphosphate_hydrolases_
CBSS_290633_1_peg_1906
Fructose_utilization
CBSS_221988_1_peg_771
Cluster_of_Unknown_Function
Calvin_Benson_cycle
CBSS_89187_3_peg_2957
tRNA_sulfuration
CBSS_205922_3_peg_1809
Urate_degradation
Ribosomal_Protein_L28P_relat [..]_of_uncharacterized_proteins
A_Gram_positive_cluster_that [..]_of_uncharacterized_proteins
Regulation_of_virulence
Trehalose_Uptake_and_Utilization
Photorespiration__oxidative_C2_cycle_
YgfZ_Fe_S_clustering
CBSS_261594_1_peg_788
recX__and_regulatory_cluster
Vibrio_Polysaccharide__VPS__Biosynthesis
Allantoin_degradation
Lipoprotein_Biosynthesis
Lipoproteins
Utilization_of_glutathione_as_a_sulphur_source
Pyruvate_ferredoxin_oxidoreductase
Probably_Ybbk_related_hypothetical_membrane_proteins
CBSS_316057_3_peg_659
Uni__Sym__and_Antiporters
rRNA_modification_Bacteria
Proline_Synthesis
Molybdopterin_oxidoreductase
CBSS_269799_3_peg_2220
Detoxification
D_tyrosyl_tRNA_Tyr__deacylase
Choline_Transport
Na___H___antiporter
Fatty_acid_degradation_regulons
CO2_uptake__carboxysome
NiFe_hydrogenase_maturation
Taurine_Utilization
Folate_transporters
Experimental__Creatinine_degradation
Galactosylceramide_and_Sulfatide_metabolism
CBSS_292415_3_peg_2341
Hfl_operon
CBSS_412883_3_peg_2513
Lysine_degradation
Hydrogenases
CBSS_345072_3_peg_1318
CBSS_224911_1_peg_435
Methionine_Salvage
CBSS_138119_3_peg_2719
Embden_Meyerhof_and_Gluconeogenesis_Archaeal
YrdC_YciO
Biosynthesis_of_phenylpropanoids
Anaerobic_benzoate_metabolism
CBSS_176279_3_peg_1389
Carbon_monoxide_induced_hydrogenase
CBSS_224911_1_peg_7673
Respiration
F0F1_type_ATP_synthase
ATP_synthases
Respiration
−4
−2
0
2
4
LDA SCORE (log 10)