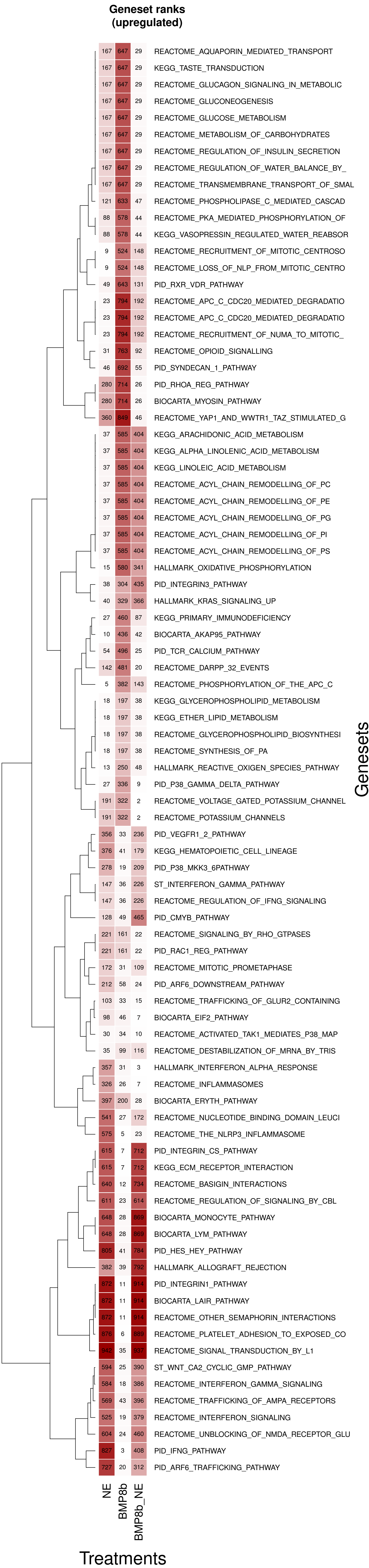
Geneset ranks
(upregulated)
167 647 29 REACTOME_AQUAPORIN_MEDIATED_TRANSPORT
167 647 29 KEGG_TASTE_TRANSDUCTION
167 647 29 REACTOME_GLUCAGON_SIGNALING_IN_METABOLIC
167 647 29 REACTOME_GLUCONEOGENESIS
167 647 29 REACTOME_GLUCOSE_METABOLISM
167 647 29 REACTOME_METABOLISM_OF_CARBOHYDRATES
167 647 29 REACTOME_REGULATION_OF_INSULIN_SECRETION
167 647 29 REACTOME_REGULATION_OF_WATER_BALANCE_BY_
167 647 29 REACTOME_TRANSMEMBRANE_TRANSPORT_OF_SMAL
121 633 47 REACTOME_PHOSPHOLIPASE_C_MEDIATED_CASCAD
88 578 44 REACTOME_PKA_MEDIATED_PHOSPHORYLATION_OF
88 578 44 KEGG_VASOPRESSIN_REGULATED_WATER_REABSOR
9 524 148 REACTOME_RECRUITMENT_OF_MITOTIC_CENTROSO
9 524 148 REACTOME_LOSS_OF_NLP_FROM_MITOTIC_CENTRO
49 643 131 PID_RXR_VDR_PATHWAY
23 794 192 REACTOME_APC_C_CDC20_MEDIATED_DEGRADATIO
23 794 192 REACTOME_APC_C_CDC20_MEDIATED_DEGRADATIO
23 794 192 REACTOME_RECRUITMENT_OF_NUMA_TO_MITOTIC_
31 763 92 REACTOME_OPIOID_SIGNALLING
46 692 55 PID_SYNDECAN_1_PATHWAY
280 714 26 PID_RHOA_REG_PATHWAY
280 714 26 BIOCARTA_MYOSIN_PATHWAY
360 849 46 REACTOME_YAP1_AND_WWTR1_TAZ_STIMULATED_G
37 585 404 KEGG_ARACHIDONIC_ACID_METABOLISM
37 585 404 KEGG_ALPHA_LINOLENIC_ACID_METABOLISM
37 585 404 KEGG_LINOLEIC_ACID_METABOLISM
37 585 404 REACTOME_ACYL_CHAIN_REMODELLING_OF_PC
37 585 404 REACTOME_ACYL_CHAIN_REMODELLING_OF_PE
37 585 404 REACTOME_ACYL_CHAIN_REMODELLING_OF_PG
37 585 404 REACTOME_ACYL_CHAIN_REMODELLING_OF_PI
37 585 404 REACTOME_ACYL_CHAIN_REMODELLING_OF_PS
15 580 341 HALLMARK_OXIDATIVE_PHOSPHORYLATION
38 304 435 PID_INTEGRIN3_PATHWAY
40 329 366 HALLMARK_KRAS_SIGNALING_UP
27 460 87 KEGG_PRIMARY_IMMUNODEFICIENCY
10 436 42 BIOCARTA_AKAP95_PATHWAY
54 496 25 PID_TCR_CALCIUM_PATHWAY
142 481 20 REACTOME_DARPP_32_EVENTS
5 382 143 REACTOME_PHOSPHORYLATION_OF_THE_APC_C
18 197 38 KEGG_GLYCEROPHOSPHOLIPID_METABOLISM
18 197 38 KEGG_ETHER_LIPID_METABOLISM
18 197 38 REACTOME_GLYCEROPHOSPHOLIPID_BIOSYNTHESI
18 197 38 REACTOME_SYNTHESIS_OF_PA
13 250 48 HALLMARK_REACTIVE_OXIGEN_SPECIES_PATHWAY
27 336 9 PID_P38_GAMMA_DELTA_PATHWAY
191 322 2 REACTOME_VOLTAGE_GATED_POTASSIUM_CHANNEL
191 322 2 REACTOME_POTASSIUM_CHANNELS
356 33 236 PID_VEGFR1_2_PATHWAY
376 41 179 KEGG_HEMATOPOIETIC_CELL_LINEAGE
278 19 209 PID_P38_MKK3_6PATHWAY
147 36 226 ST_INTERFERON_GAMMA_PATHWAY
147 36 226 REACTOME_REGULATION_OF_IFNG_SIGNALING
128 49 465 PID_CMYB_PATHWAY
221 161 22 REACTOME_SIGNALING_BY_RHO_GTPASES
221 161 22 PID_RAC1_REG_PATHWAY
172 31 109 REACTOME_MITOTIC_PROMETAPHASE
212 58 24 PID_ARF6_DOWNSTREAM_PATHWAY
103 33 15 REACTOME_TRAFFICKING_OF_GLUR2_CONTAINING
98 46 7 BIOCARTA_EIF2_PATHWAY
30 34 10 REACTOME_ACTIVATED_TAK1_MEDIATES_P38_MAP
35 99 116 REACTOME_DESTABILIZATION_OF_MRNA_BY_TRIS
357 31 3 HALLMARK_INTERFERON_ALPHA_RESPONSE
326 26 7 REACTOME_INFLAMMASOMES
397 200 28 BIOCARTA_ERYTH_PATHWAY
541 27 172 REACTOME_NUCLEOTIDE_BINDING_DOMAIN_LEUCI
575 5 23 REACTOME_THE_NLRP3_INFLAMMASOME
615 7 712 PID_INTEGRIN_CS_PATHWAY
615 7 712 KEGG_ECM_RECEPTOR_INTERACTION
640 12 734 REACTOME_BASIGIN_INTERACTIONS
611 23 614 REACTOME_REGULATION_OF_SIGNALING_BY_CBL
648 28 869 BIOCARTA_MONOCYTE_PATHWAY
648 28 869 BIOCARTA_LYM_PATHWAY
805 41 784 PID_HES_HEY_PATHWAY
382 39 792 HALLMARK_ALLOGRAFT_REJECTION
872 11 914 PID_INTEGRIN1_PATHWAY
872 11 914 BIOCARTA_LAIR_PATHWAY
872 11 914 REACTOME_OTHER_SEMAPHORIN_INTERACTIONS
876 6 889 REACTOME_PLATELET_ADHESION_TO_EXPOSED_CO
942 35 937 REACTOME_SIGNAL_TRANSDUCTION_BY_L1
594 25 390 ST_WNT_CA2_CYCLIC_GMP_PATHWAY
584 18 386 REACTOME_INTERFERON_GAMMA_SIGNALING
569 43 396 REACTOME_TRAFFICKING_OF_AMPA_RECEPTORS
525 19 379 REACTOME_INTERFERON_SIGNALING
604 24 460 REACTOME_UNBLOCKING_OF_NMDA_RECEPTOR_GLU
827 3 408 PID_IFNG_PATHWAY
727 20 312 PID_ARF6_TRAFFICKING_PATHWAY
Genesets
NE
BMP8b
BMP8b_NE
Treatments