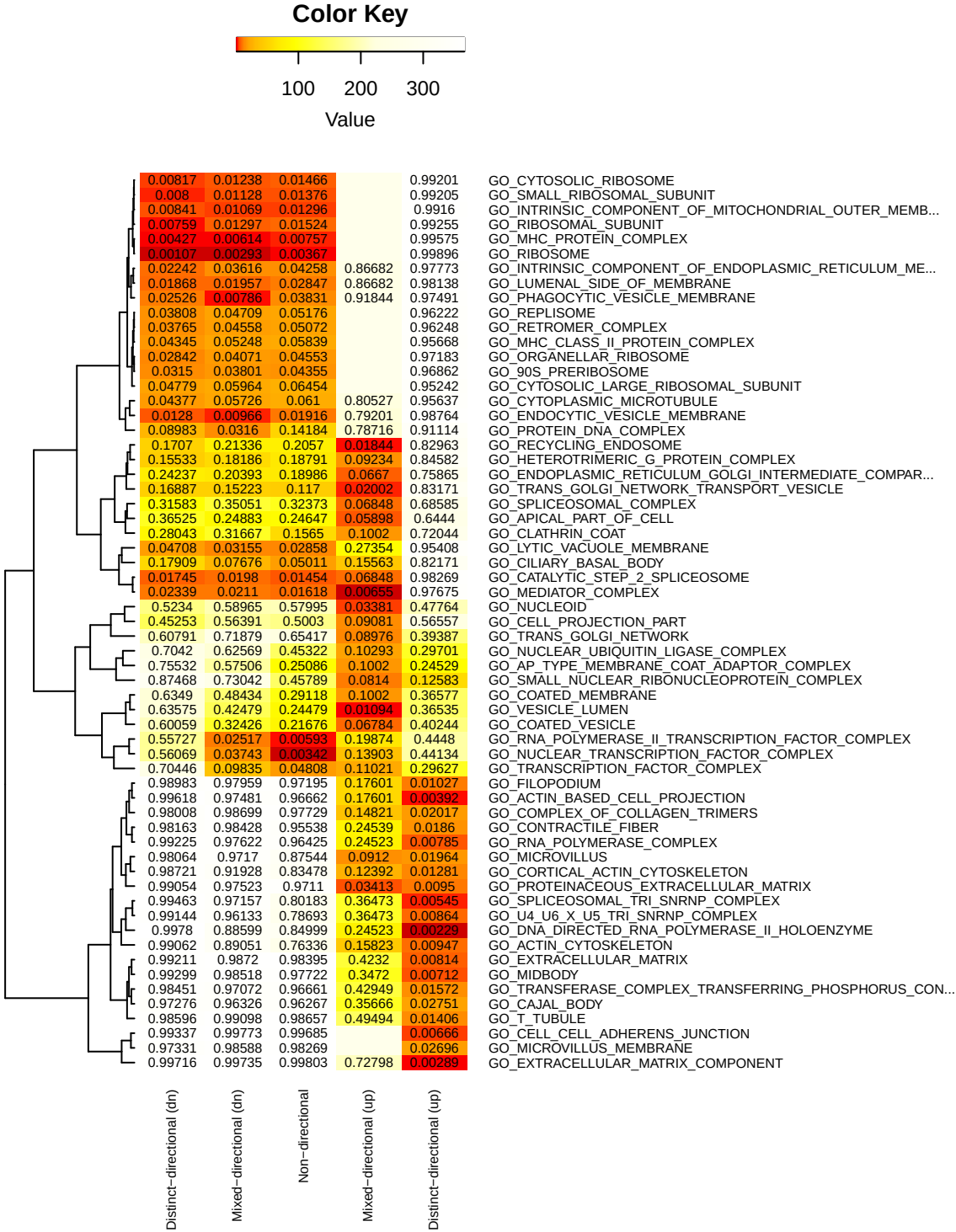
Color Key
100
200
300
Value
Distinct−directional (dn)
Mixed−directional (dn)
Non−directional
Mixed−directional (up)
Distinct−directional (up)
0.00817 0.01238 0.01466 0.99201 GO_CYTOSOLIC_RIBOSOME
0.008 0.01128 0.01376 0.99205 GO_SMALL_RIBOSOMAL_SUBUNIT
0.00841 0.01069 0.01296 0.9916 GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_OUTER_MEMB...
0.00759 0.01297 0.01524 0.99255 GO_RIBOSOMAL_SUBUNIT
0.00427 0.00614 0.00757 0.99575 GO_MHC_PROTEIN_COMPLEX
0.00107 0.00293 0.00367 0.99896 GO_RIBOSOME
0.02242 0.03616 0.04258 0.86682 0.97773 GO_INTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_ME...
0.01868 0.01957 0.02847 0.86682 0.98138 GO_LUMENAL_SIDE_OF_MEMBRANE
0.02526 0.00786 0.03831 0.91844 0.97491 GO_PHAGOCYTIC_VESICLE_MEMBRANE
0.03808 0.04709 0.05176 0.96222 GO_REPLISOME
0.03765 0.04558 0.05072 0.96248 GO_RETROMER_COMPLEX
0.04345 0.05248 0.05839 0.95668 GO_MHC_CLASS_II_PROTEIN_COMPLEX
0.02842 0.04071 0.04553 0.97183 GO_ORGANELLAR_RIBOSOME
0.0315 0.03801 0.04355 0.96862 GO_90S_PRERIBOSOME
0.04779 0.05964 0.06454 0.95242 GO_CYTOSOLIC_LARGE_RIBOSOMAL_SUBUNIT
0.04377 0.05726 0.061 0.80527 0.95637 GO_CYTOPLASMIC_MICROTUBULE
0.0128 0.00966 0.01916 0.79201 0.98764 GO_ENDOCYTIC_VESICLE_MEMBRANE
0.08983 0.0316 0.14184 0.78716 0.91114 GO_PROTEIN_DNA_COMPLEX
0.1707 0.21336 0.2057 0.01844 0.82963 GO_RECYCLING_ENDOSOME
0.15533 0.18186 0.18791 0.09234 0.84582 GO_HETEROTRIMERIC_G_PROTEIN_COMPLEX
0.24237 0.20393 0.18986 0.0667 0.75865 GO_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPAR...
0.16887 0.15223 0.117 0.02002 0.83171 GO_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE
0.31583 0.35051 0.32373 0.06848 0.68585 GO_SPLICEOSOMAL_COMPLEX
0.36525 0.24883 0.24647 0.05898 0.6444 GO_APICAL_PART_OF_CELL
0.28043 0.31667 0.1565 0.1002 0.72044 GO_CLATHRIN_COAT
0.04708 0.03155 0.02858 0.27354 0.95408 GO_LYTIC_VACUOLE_MEMBRANE
0.17909 0.07676 0.05011 0.15563 0.82171 GO_CILIARY_BASAL_BODY
0.01745 0.0198 0.01454 0.06848 0.98269 GO_CATALYTIC_STEP_2_SPLICEOSOME
0.02339 0.0211 0.01618 0.00655 0.97675 GO_MEDIATOR_COMPLEX
0.5234 0.58965 0.57995 0.03381 0.47764 GO_NUCLEOID
0.45253 0.56391 0.5003 0.09081 0.56557 GO_CELL_PROJECTION_PART
0.60791 0.71879 0.65417 0.08976 0.39387 GO_TRANS_GOLGI_NETWORK
0.7042 0.62569 0.45322 0.10293 0.29701 GO_NUCLEAR_UBIQUITIN_LIGASE_COMPLEX
0.75532 0.57506 0.25086 0.1002 0.24529 GO_AP_TYPE_MEMBRANE_COAT_ADAPTOR_COMPLEX
0.87468 0.73042 0.45789 0.0814 0.12583 GO_SMALL_NUCLEAR_RIBONUCLEOPROTEIN_COMPLEX
0.6349 0.48434 0.29118 0.1002 0.36577 GO_COATED_MEMBRANE
0.63575 0.42479 0.24479 0.01094 0.36535 GO_VESICLE_LUMEN
0.60059 0.32426 0.21676 0.06784 0.40244 GO_COATED_VESICLE
0.55727 0.02517 0.00593 0.19874 0.4448 GO_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_COMPLEX
0.56069 0.03743 0.00342 0.13903 0.44134 GO_NUCLEAR_TRANSCRIPTION_FACTOR_COMPLEX
0.70446 0.09835 0.04808 0.11021 0.29627 GO_TRANSCRIPTION_FACTOR_COMPLEX
0.98983 0.97959 0.97195 0.17601 0.01027 GO_FILOPODIUM
0.99618 0.97481 0.96662 0.17601 0.00392 GO_ACTIN_BASED_CELL_PROJECTION
0.98008 0.98699 0.97729 0.14821 0.02017 GO_COMPLEX_OF_COLLAGEN_TRIMERS
0.98163 0.98428 0.95538 0.24539 0.0186 GO_CONTRACTILE_FIBER
0.99225 0.97622 0.96425 0.24523 0.00785 GO_RNA_POLYMERASE_COMPLEX
0.98064 0.9717 0.87544 0.0912 0.01964 GO_MICROVILLUS
0.98721 0.91928 0.83478 0.12392 0.01281 GO_CORTICAL_ACTIN_CYTOSKELETON
0.99054 0.97523 0.9711 0.03413 0.0095 GO_PROTEINACEOUS_EXTRACELLULAR_MATRIX
0.99463 0.97157 0.80183 0.36473 0.00545 GO_SPLICEOSOMAL_TRI_SNRNP_COMPLEX
0.99144 0.96133 0.78693 0.36473 0.00864 GO_U4_U6_X_U5_TRI_SNRNP_COMPLEX
0.9978 0.88599 0.84999 0.24523 0.00229 GO_DNA_DIRECTED_RNA_POLYMERASE_II_HOLOENZYME
0.99062 0.89051 0.76336 0.15823 0.00947 GO_ACTIN_CYTOSKELETON
0.99211 0.9872 0.98395 0.4232 0.00814 GO_EXTRACELLULAR_MATRIX
0.99299 0.98518 0.97722 0.3472 0.00712 GO_MIDBODY
0.98451 0.97072 0.96661 0.42949 0.01572 GO_TRANSFERASE_COMPLEX_TRANSFERRING_PHOSPHORUS_CON...
0.97276 0.96326 0.96267 0.35666 0.02751 GO_CAJAL_BODY
0.98596 0.99098 0.98657 0.49494 0.01406 GO_T_TUBULE
0.99337 0.99773 0.99685 0.00666 GO_CELL_CELL_ADHERENS_JUNCTION
0.97331 0.98588 0.98269 0.02696 GO_MICROVILLUS_MEMBRANE
0.99716 0.99735 0.99803 0.72798 0.00289 GO_EXTRACELLULAR_MATRIX_COMPONENT