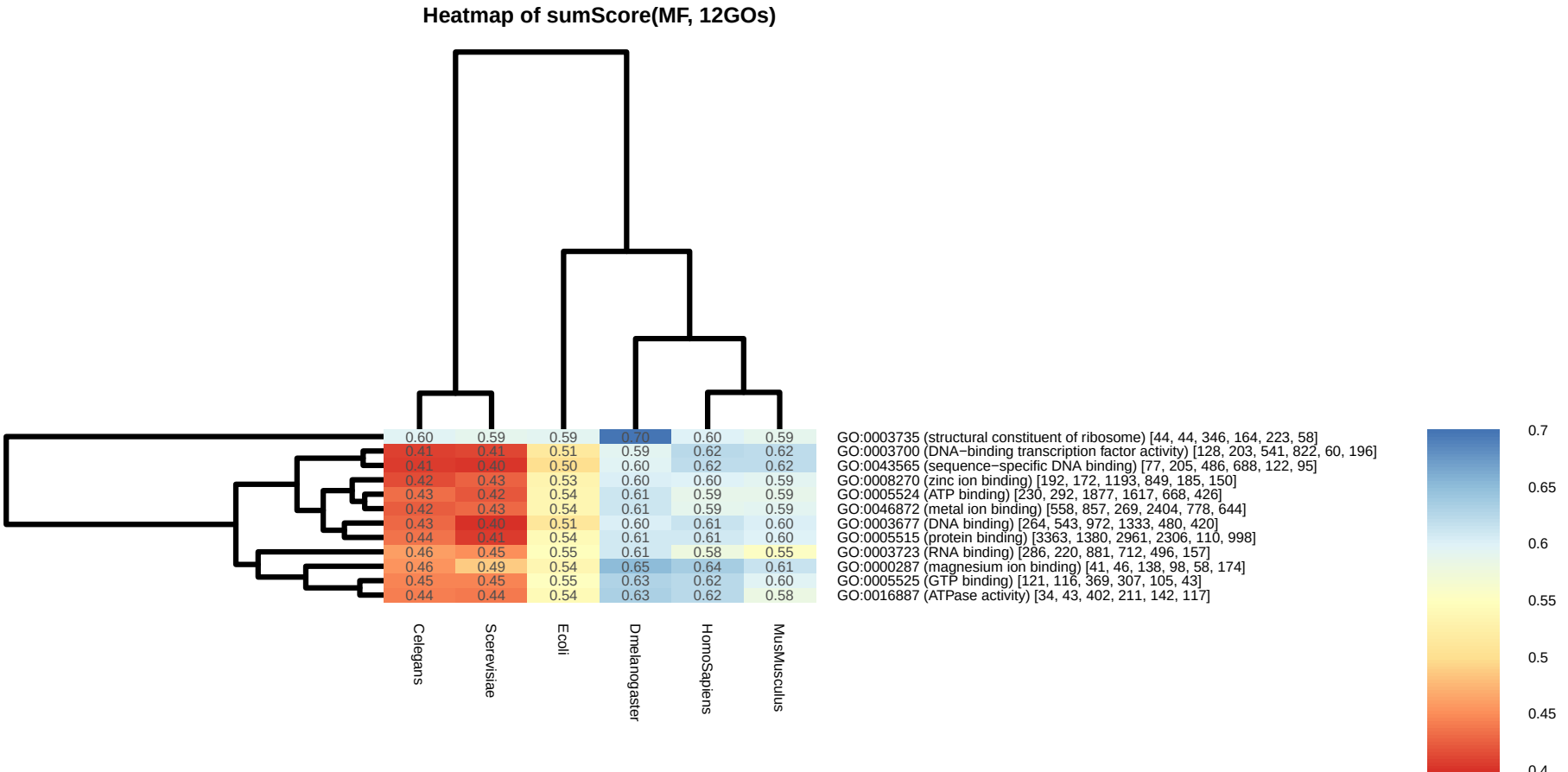
Heatmap of sumScore(MF, 12GOs)
0.60 0.59 0.59 0.70 0.60 0.59 GO:0003735 (structural constituent of ribosome) [44, 44, 346, 164, 223, 58]
0.41 0.41 0.51 0.59 0.62 0.62 GO:0003700 (DNA−binding transcription factor activity) [128, 203, 541, 822, 60, 196]
0.41 0.40 0.50 0.60 0.62 0.62 GO:0043565 (sequence−specific DNA binding) [77, 205, 486, 688, 122, 95]
0.42 0.43 0.53 0.60 0.60 0.59 GO:0008270 (zinc ion binding) [192, 172, 1193, 849, 185, 150]
0.43 0.42 0.54 0.61 0.59 0.59 GO:0005524 (ATP binding) [230, 292, 1877, 1617, 668, 426]
0.42 0.43 0.54 0.61 0.59 0.59 GO:0046872 (metal ion binding) [558, 857, 269, 2404, 778, 644]
0.43 0.40 0.51 0.60 0.61 0.60 GO:0003677 (DNA binding) [264, 543, 972, 1333, 480, 420]
0.44 0.41 0.54 0.61 0.61 0.60 GO:0005515 (protein binding) [3363, 1380, 2961, 2306, 110, 998]
0.46 0.45 0.55 0.61 0.58 0.55 GO:0003723 (RNA binding) [286, 220, 881, 712, 496, 157]
0.46 0.49 0.54 0.65 0.64 0.61 GO:0000287 (magnesium ion binding) [41, 46, 138, 98, 58, 174]
0.45 0.45 0.55 0.63 0.62 0.60 GO:0005525 (GTP binding) [121, 116, 369, 307, 105, 43]
0.44 0.44 0.54 0.63 0.62 0.58 GO:0016887 (ATPase activity) [34, 43, 402, 211, 142, 117]
Celegans
Scerevisiae
Ecoli
Dmelanogaster
HomoSapiens
MusMusculus
0.7
0.65
0.6
0.55
0.5
0.45
0.4