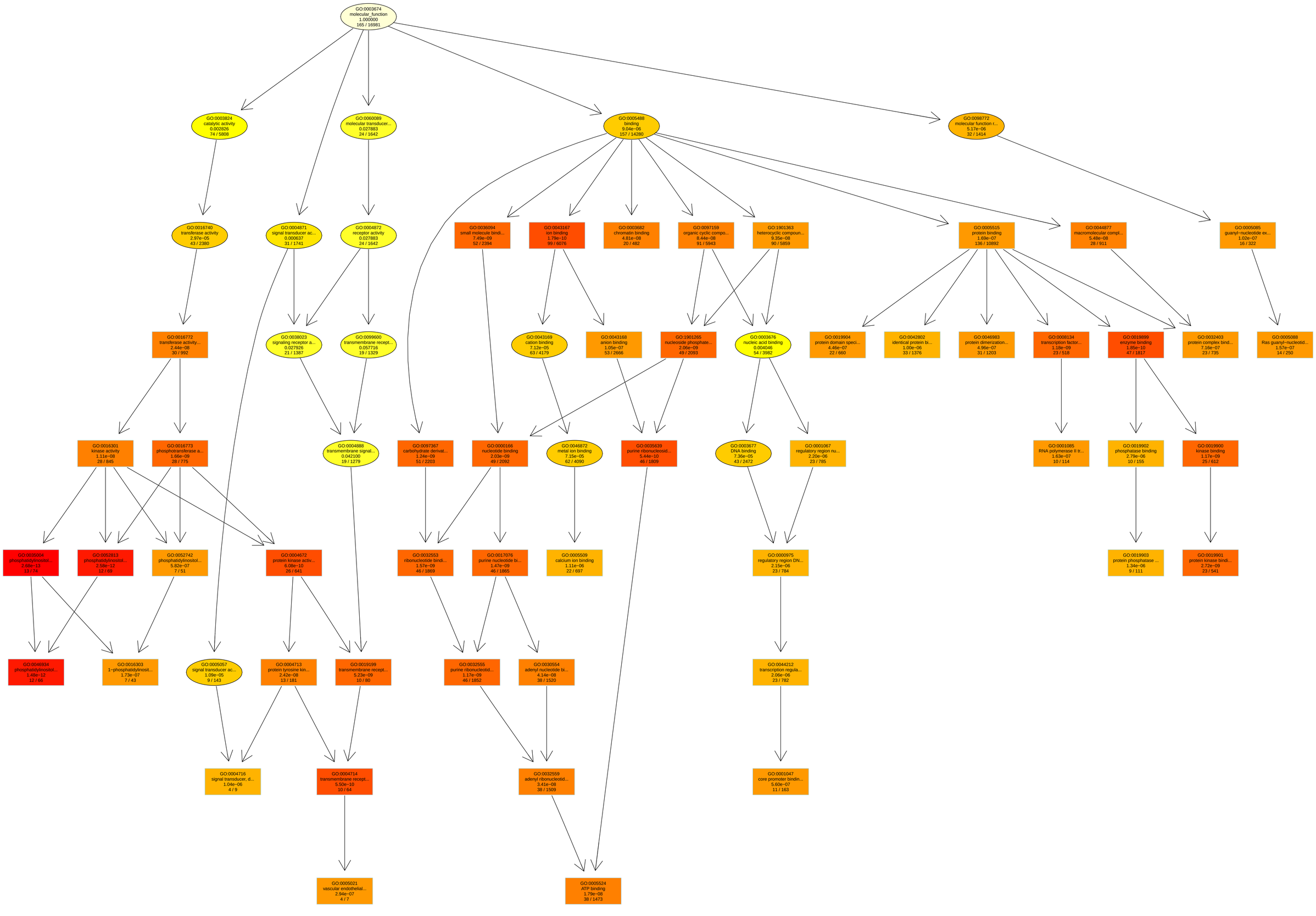
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GO:0003824 catalytic activity 0.002826 74 / 5808
GO:0060089 molecular transducer... 0.027883 24 / 1642
GO:0005488 binding 9.04e−06 157 / 14280
GO:0098772 molecular function r... 5.17e−06 32 / 1414
GO:0016740 transferase activity 2.97e−05 43 / 2380
GO:0004871 signal transducer ac... 0.000637 31 / 1741
GO:0004872 receptor activity 0.027883 24 / 1642
GO:0036094 small molecule bindi... 7.49e−09 52 / 2394
GO:0043167 ion binding 1.79e−10 99 / 6076
GO:0003682 chromatin binding 4.81e−08 20 / 482
GO:0097159 organic cyclic compo... 8.44e−08 91 / 5943
GO:1901363 heterocyclic compoun... 9.35e−08 90 / 5859
GO:0005515 protein binding 1.69e−07 136 / 10892
GO:0044877 macromolecular compl... 5.48e−08 28 / 911
GO:0005085 guanyl−nucleotide ex... 1.02e−07 16 / 322
GO:0016772 transferase activity,... 2.44e−08 30 / 992
GO:0038023 signaling receptor a... 0.027926 21 / 1387
GO:0099600 transmembrane recept... 0.057716 19 / 1329
GO:0043169 cation binding 7.12e−05 63 / 4179
GO:0043168 anion binding 1.05e−07 53 / 2666
GO:1901265 nucleoside phosphate... 2.06e−09 49 / 2093
GO:0003676 nucleic acid binding 0.004046 54 / 3982
GO:0019904 protein domain speci... 4.46e−07 22 / 660
GO:0042802 identical protein bi... 1.00e−06 33 / 1376
GO:0046983 protein dimerization... 4.96e−07 31 / 1203
GO:0008134 transcription factor... 1.18e−09 23 / 518
GO:0019899 enzyme binding 1.85e−10 47 / 1817
GO:0032403 protein complex bind... 7.16e−07 23 / 735
GO:0005088 Ras guanyl−nucleotid... 1.57e−07 14 / 250
GO:0016301 kinase activity 1.11e−08 28 / 845
GO:0016773 phosphotransferase a... 1.66e−09 28 / 775
GO:0004888 transmembrane signal... 0.042100 19 / 1279
GO:0097367 carbohydrate derivat... 1.24e−09 51 / 2203
GO:0000166 nucleotide binding 2.03e−09 49 / 2092
GO:0046872 metal ion binding 7.15e−05 62 / 4090
GO:0035639 purine ribonucleosid... 5.44e−10 46 / 1809
GO:0003677 DNA binding 7.36e−05 43 / 2472
GO:0001067 regulatory region nu... 2.20e−06 23 / 785
GO:0001085 RNA polymerase II tr... 1.63e−07 10 / 114
GO:0019902 phosphatase binding 2.79e−06 10 / 155
GO:0019900 kinase binding 1.17e−09 25 / 612
GO:0035004 phosphatidylinositol... 2.68e−13 13 / 74
GO:0052813 phosphatidylinositol... 2.58e−12 12 / 69
GO:0052742 phosphatidylinositol... 5.82e−07 7 / 51
GO:0004672 protein kinase activ... 6.08e−10 26 / 641
GO:0032553 ribonucleotide bindi... 1.57e−09 46 / 1869
GO:0017076 purine nucleotide bi... 1.47e−09 46 / 1865
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GO:0000975 regulatory region DN... 2.15e−06 23 / 784
GO:0019903 protein phosphatase ... 1.34e−06 9 / 111
GO:0019901 protein kinase bindi... 2.72e−09 23 / 541
GO:0046934 phosphatidylinositol... 1.48e−12 12 / 66
GO:0016303 1−phosphatidylinosit... 1.73e−07 7 / 43
GO:0005057 signal transducer ac... 1.09e−05 9 / 143
GO:0004713 protein tyrosine kin... 2.42e−08 13 / 181
GO:0019199 transmembrane recept... 5.23e−09 10 / 80
GO:0032555 purine ribonucleotid... 1.17e−09 46 / 1852
GO:0030554 adenyl nucleotide bi... 4.14e−08 38 / 1520
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GO:0032559 adenyl ribonucleotid... 3.41e−08 38 / 1509
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GO:0005021 vascular endothelial... 2.94e−07 4 / 7
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