Table 1

Biological Process

cellular macromolecular complex assembly	C=336;O=32;E=13.19;R=2.43;rawP=3.34e-06;adjP=0.0012
macromolecular complex subunit organization	C=741;O=55;E=29.09;R=1.89;rawP=3.63e-06;adjP=0.0012
macromolecular complex assembly	C=672;O=51;E=26.38;R=1.93;rawP=4.49e-06;adjP=0.0012
organelle organization	C=1339;O=87;E=52.57;R=1.66;rawP=1.38e-06;adjP=0.0012
cellular macromolecular complex subunit organization	C=396;O=36;E=15.55;R=2.32;rawP=2.42e-06;adjP=0.0012
protein complex assembly	C=520;O=42;E=20.41;R=2.06;rawP=7.28e-06;adjP=0.0014
protein complex biogenesis	C=520;O=42;E=20.41;R=2.06;rawP=7.28e-06;adjP=0.0014
cellular protein complex assembly	C=184;O=21;E=7.22;R=2.91;rawP=1.12e-05;adjP=0.0019
proteasomal ubiquitin-dependent protein catabolic process	C=107;O=15;E=4.20;R=3.57;rawP=1.79e-05;adjP=0.0022
proteasomal protein catabolic process	C=107;O=15;E=4.20;R=3.57;rawP=1.79e-05;adjP=0.0022
interspecies interaction between organisms	C=280;O=27;E=10.99;R=2.46;rawP=1.56e-05;adjP=0.0022
cell cycle	C=895;O=60;E=35.14;R=1.71;rawP=2.98e-05;adjP=0.0033
amine biosynthetic process	C=78;O=12;E=3.06;R=3.92;rawP=4.83e-05;adjP=0.0050
positive regulation of ubiquitin-protein ligase activity	C=67;O=11;E=2.63;R=4.18;rawP=5.37e-05;adjP=0.0052
positive regulation of ligase activity	C=70;O=11;E=2.75;R=4.00;rawP=8.13e-05;adjP=0.0073
chromatin organization	C=364;O=30;E=14.29;R=2.10;rawP=0.0001;adjP=0.0084