GO enrichment vesicle-mediated transport structural molecule activity regulation of transcription by RNA polymerase II receptor-mediated endocytosis protein catabolic process --1 \* log10(Qvalue) proteasome regulatory particle, lid subcomplex phospholipid translocation nucleosome disassembly nucleolus -3 GO term intracellular protein transport histone-lysine N-methyltransferase activity histone methyltransferase activity (H3-K36 specific) -Genes cytosol -100 cytoplasm -200 clathrin light chain binding -300 clathrin complex clathrin coat of trans-Golgi network vesicle clathrin coat of coated pit -ATPase-coupled intramembrane lipid transporter activity -ATP-dependent chromatin remodeling -5 4 8 -log10(Pvalue)