enriched KEGG pathway heterocyclic compound binding -Gene number organic cyclic compound binding 10 structural molecule activity -20 30 structural constituent of ribosome 40 KEGG_pathway 50 RNA binding $-\log_{10}(pvalue)$ transition metal ion binding rRNA binding -2.5 2.0 peptidase activity -1.5 peptidase activity, acting on _L-amino acid peptides 1.0 4 iron, 4 sulfur cluster binding 0.15 0.05 0.10 0.20 0.25 GeneRatio