| 156/156 structural molecule | |
|--|-----------------|
| 116/116 structural constituent of ribosome | p < 0.001 |
| 21/21 extracellular matrix structural constituent | p < 0.01 |
| 42/42 translation regulator, nucleic acid binding | p < 0.05 |
| 8/8 translation elongation factor | |
| 16/17 oxidoreductase, acting on the CH–CH group of donors | |
| 8/8 acyl-CoA dehydrogenase | |
| 303/307 oxidoreductase | |
| 5/5 3-hydroxyacyl-CoA dehydrogenase | |
| 37/37 oxidoreductase, acting on CH-OH group of donors | |
| 25/25 oxidoreductase, acting on the CH-OH group of donors, NAD or NADP as acceptor | |
| 43/43 antioxidant | |
| 21/21 catalase | |
| 19/19 oxidoreductase, acting on NAD(P)H | |
| 30/30 electron transfer | |
| 10/10 heme-copper terminal oxidase | |
| 17/17 proton transmembrane transporter | |
| 28/28 carbon-oxygen lyase | |
| 15/15 carbonate dehydratase | |
| 12/12 threonine–type peptidase | |
| 8/8 glutathione transferase | |
| 6/6 pyridoxal phosphate binding | |
| 55/55 hydrolase, acting on glycosyl bonds | |
| 24/24 carboxylic ester hydrolase | |
| 8/8 phospholipase A2 | |
| 31/31 enzyme inhibitor | |
| 19/20 scavenger receptor | |
| 8/8 O-methyltransferase | |