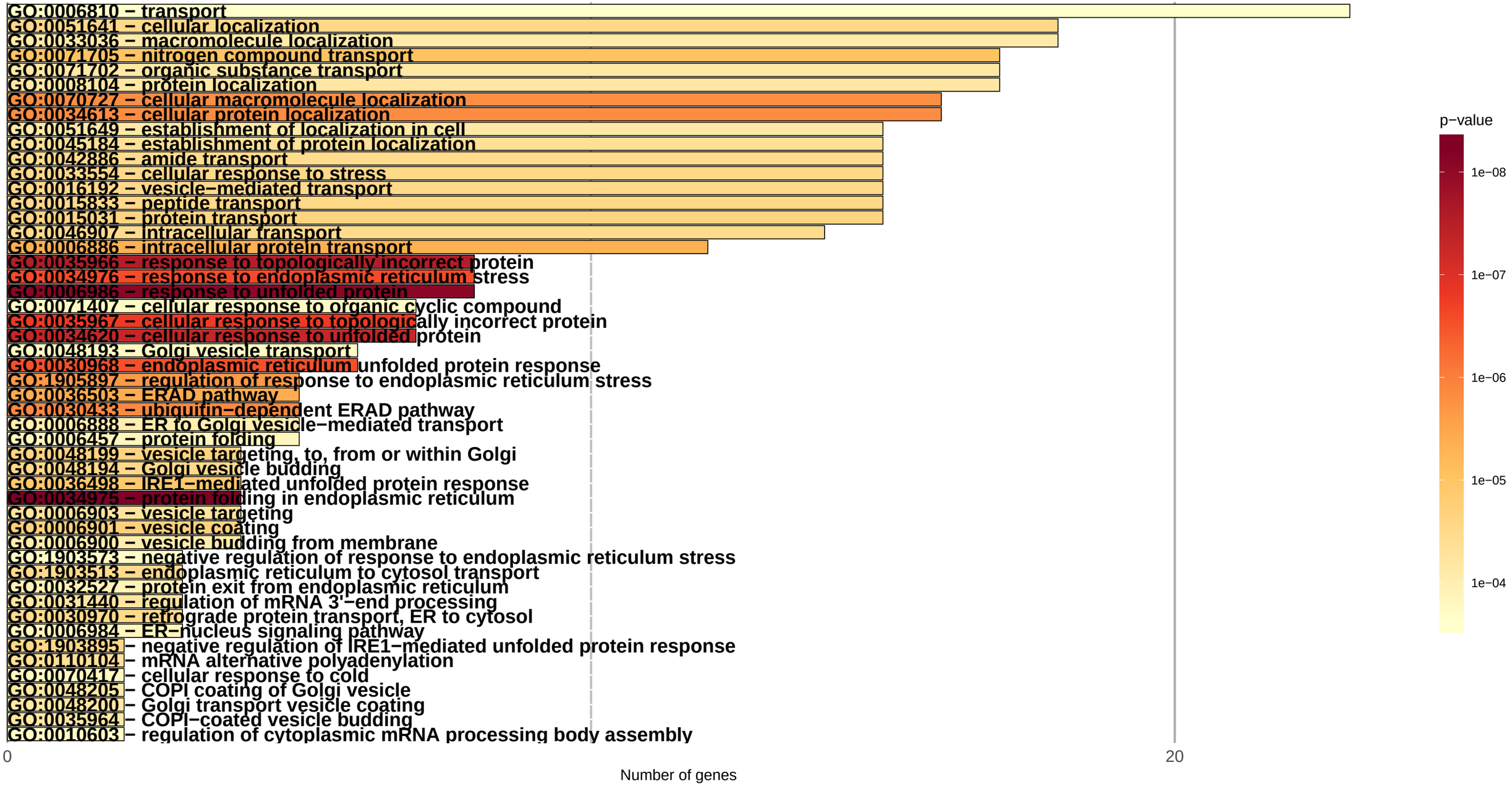
GO:0006810 − transport
GO:0051641 − cellular localization
GO:0033036 − macromolecule localization
GO:0071705 − nitrogen compound transport
GO:0071702 − organic substance transport
GO:0008104 − protein localization
GO:0070727 − cellular macromolecule localization
GO:0034613 − cellular protein localization
GO:0051649 − establishment of localization in cell
GO:0045184 − establishment of protein localization
GO:0042886 − amide transport
GO:0033554 − cellular response to stress
GO:0016192 − vesicle−mediated transport
GO:0015833 − peptide transport
GO:0015031 − protein transport
GO:0046907 − intracellular transport
GO:0006886 − intracellular protein transport
GO:0035966 − response to topologically incorrect protein
GO:0034976 − response to endoplasmic reticulum stress
GO:0006986 − response to unfolded protein
GO:0071407 − cellular response to organic cyclic compound
GO:0035967 − cellular response to topologically incorrect protein
GO:0034620 − cellular response to unfolded protein
GO:0048193 − Golgi vesicle transport
GO:0030968 − endoplasmic reticulum unfolded protein response
GO:1905897 − regulation of response to endoplasmic reticulum stress
GO:0036503 − ERAD pathway
GO:0030433 − ubiquitin−dependent ERAD pathway
GO:0006888 − ER to Golgi vesicle−mediated transport
GO:0006457 − protein folding
GO:0048199 − vesicle targeting, to, from or within Golgi
GO:0048194 − Golgi vesicle budding
GO:0036498 − IRE1−mediated unfolded protein response
GO:0034975 − protein folding in endoplasmic reticulum
GO:0006903 − vesicle targeting
GO:0006901 − vesicle coating
GO:0006900 − vesicle budding from membrane
GO:1903573 − negative regulation of response to endoplasmic reticulum stress
GO:1903513 − endoplasmic reticulum to cytosol transport
GO:0032527 − protein exit from endoplasmic reticulum
GO:0031440 − regulation of mRNA 3'−end processing
GO:0030970 − retrograde protein transport, ER to cytosol
GO:0006984 − ER−nucleus signaling pathway
GO:1903895 − negative regulation of IRE1−mediated unfolded protein response
GO:0110104 − mRNA alternative polyadenylation
GO:0070417 − cellular response to cold
GO:0048205 − COPI coating of Golgi vesicle
GO:0048200 − Golgi transport vesicle coating
GO:0035964 − COPI−coated vesicle budding
GO:0010603 − regulation of cytoplasmic mRNA processing body assembly
0
20
Number of genes
p−value
1e−08
1e−07
1e−06
1e−05
1e−04