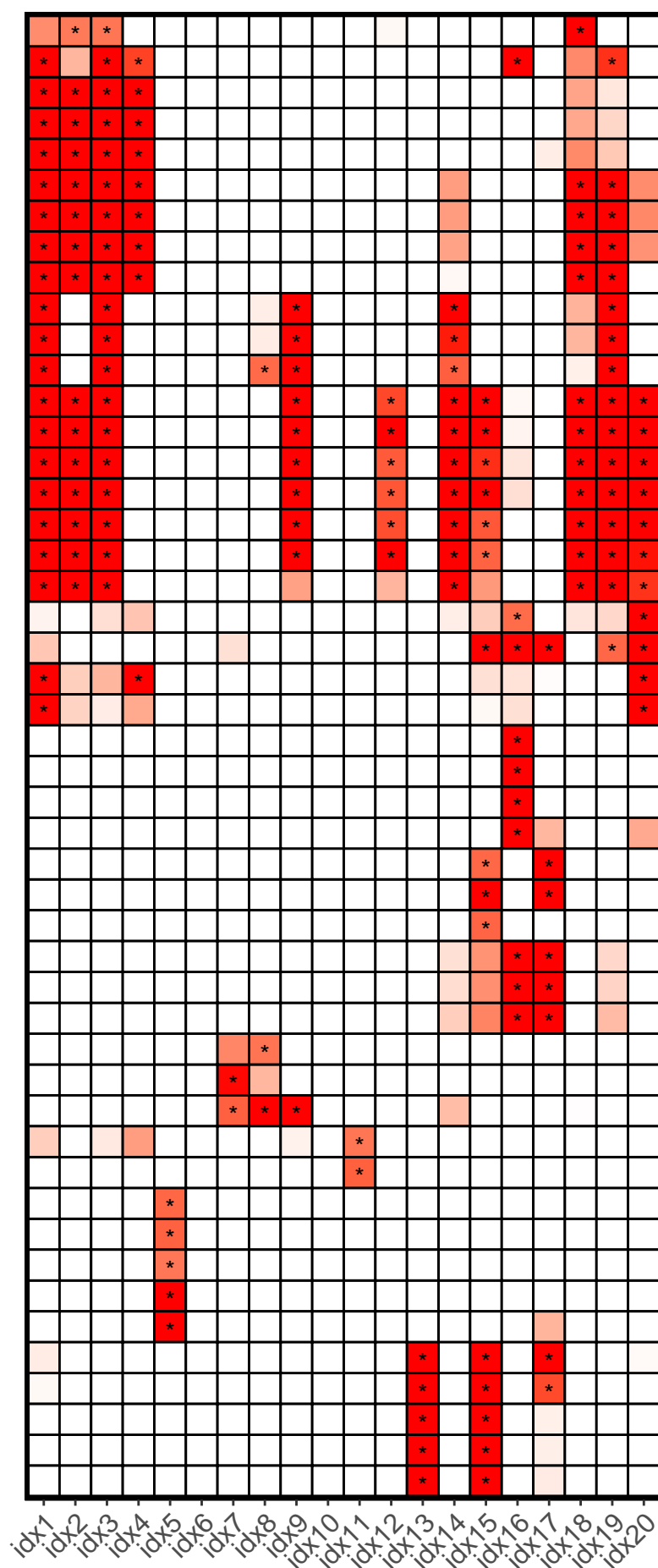
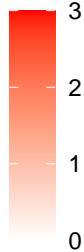


– Log P–value



GO:0071496:cellular response to external stimulus

GO:0022604:regulation of cell morphogenesis

GO:2000147:positive regulation of cell motility

GO:0030335:positive regulation of cell migration

GO:0009888:tissue development

GO:0043062:extracellular structure organization

GO:0030198:extracellular matrix organization

GO:0045229:external encapsulating structure organization

GO:0030199:collagen fibril organization

GO:0016072:rRNA metabolic process

GO:0006364:rRNA processing

GO:0042254:ribosome biogenesis

GO:0070972:protein localization to endoplasmic reticulum

GO:0006613:cotranslational protein targeting to membrane

GO:0045047:protein targeting to ER

GO:0072599:establishment of protein localization to endoplasmic reticulum

GO:0006614:SRP–dependent cotranslational protein targeting to membrane

GO:0000184:nuclear–transcribed mRNA catabolic process, nonsense–mediated decay

GO:0019080:viral gene expression

GO:0009952:anterior/posterior pattern specification

GO:0007399:nervous system development

GO:0007389:pattern specification process

GO:0003002:regionalization

GO:0044264:cellular polysaccharide metabolic process

GO:0016358:dendrite development

GO:0005976:polysaccharide metabolic process

GO:0050808:synapse organization

GO:0098742:cell–cell adhesion via plasma–membrane adhesion molecules

GO:0007156:homophilic cell adhesion via plasma membrane adhesion molecules

GO:0048668:collateral sprouting

GO:0048858:cell projection morphogenesis

GO:0120039:plasma membrane bounded cell projection morphogenesis

GO:0032990:cell part morphogenesis

GO:0046485:ether lipid metabolic process

GO:1902001:fatty acid transmembrane transport

GO:0090407:organophosphate biosynthetic process

GO:0007159:leukocyte cell–cell adhesion

GO:0006040:amino sugar metabolic process

GO:1904970:brush border assembly

GO:0098655:cation transmembrane transport

GO:0070268:cornification

GO:0034220:ion transmembrane transport

GO:0099601:regulation of neurotransmitter receptor activity

GO:0006357:regulation of transcription by RNA polymerase II

GO:0006366:transcription by RNA polymerase II

GO:0097659:nucleic acid–templated transcription

GO:0006351:transcription, DNA–templated

GO:0032774:RNA biosynthetic process