Filtered Proximal High vs Low Enriched in High Enriched in Low WP RETINOBLASTOMA GENE IN CANCER-**GOMF POLYSACCHARIDE BINDING** GOBP RIBOSOME BIOGENESIS **BIOCARTA INFLAM PATHWAY** GOBP RIBONUCLEOPROTEIN COMPLEX BIOGENESIS GOBP IMMUNOLOGICAL MEMORY FORMATION PROCESS GOBP RRNA METABOLIC PROCESS-GOBP IMMUNOLOGICAL MEMORY PROCESS GOBP NCRNA PROCESSING GOBP MUSCLE CONTRACTION GOBP DNA REPAIR -WP GLUCURONIDATION -GOBP DNA DEPENDENT DNA REPLICATION -**GOBP MUSCLE SYSTEM PROCESS** Count GOBP NCRNA METABOLIC PROCESS GOBP URONIC ACID METABOLIC PROCESS GOBP SISTER CHROMATID SEGREGATION -REACTOME GLUCURONIDATION -200 GOBP DNA REPLICATION -**REACTOME PD 1 SIGNALING** p.adjust GOBP NUCLEAR CHROMOSOME SEGREGATION -KEGG ASCORBATE AND ALDARATE METABOLISM 0.0125 0.0100 GOCC CHROMOSOMAL REGION-**GOMF HORMONE RECEPTOR BINDING** 0.0075 0.0050 GOBP CHROMOSOME SEGREGATION -**BIOCARTA TCRA PATHWAY** 0.0025 GOBP MITOTIC NUCLEAR DIVISION -**BIOCARTA MHC PATHWAY** REACTOME CELL CYCLE CHECKPOINTS -GOMF GLUCURONOSYLTRANSFERASE ACTIVITY GOBP REGULATION OF CELL CYCLE PHASE TRANSITION-KEGG PENTOSE AND GLUCURONATE INTERCONVERSIONS: WP PATHOGENESIS OF SARSCOV2 MEDIATED BY GOBP B CELL ACTIVATION -NSP9NSP10 COMPLEX **GOBP ORGANELLE FISSION-**BIOCARTA CSK PATHWAY GOBP CELL CYCLE PHASE TRANSITION -BIOCARTA CTLA4 PATHWAY GOBP INCLUSION BODY ASSEMBLY KEGG DRUG METABOLISM CYTOCHROME P450 0.2 0.3 0.4 0.5 0.6 0.2 0.3 0.4 0.5 0.6 GeneRatio