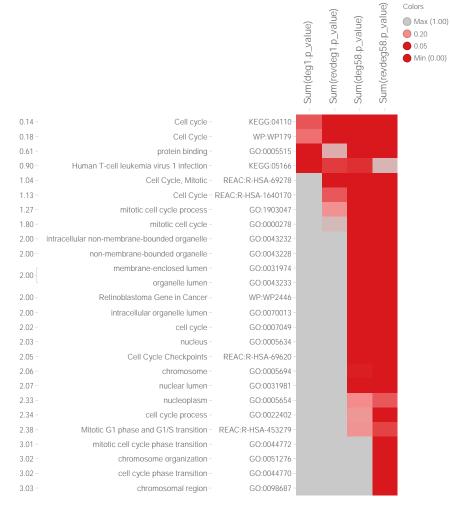# g1rev_gpro_heat

deg1.p_value, revdeg1.p_value, deg58.p_value, revdeg58.p_value per p_sum, term_name, term_id

Data table:
g1rev_settests_hitsumma

Colors
- Max (1.00)
- 0.20
- 0.05
- Min (0.00)

p_sum	term_name	term_id	Sum(deg1.p_value)	Sum(revdeg1.p_value)	Sum(deg58.p_value)	Sum(revdeg58.p_value)
0.14	Cell cycle	KEGG:04110				
0.18	Cell Cycle	WP:WP179				
0.61	protein binding	GO:0005515				
0.90	Human T-cell leukemia virus 1 infection	KEGG:05166				
1.04	Cell Cycle, Mitotic	REAC:R-HSA-69278				
1.13	Cell Cycle	REAC:R-HSA-1640170				
1.27	mitotic cell cycle process	GO:1903047				
1.80	mitotic cell cycle	GO:0000278				
2.00	intracellular non-membrane-bounded organelle	GO:0043232				
2.00	non-membrane-bounded organelle	GO:0043228				
2.00	membrane-enclosed lumen	GO:0031974				
2.00	organelle lumen	GO:0043233				
2.00	Retinoblastoma Gene in Cancer	WP:WP2446				
2.00	intracellular organelle lumen	GO:0070013				
2.02	cell cycle	GO:0007049				
2.03	nucleus	GO:0005634				
2.05	Cell Cycle Checkpoints	REAC:R-HSA-69620				
2.06	chromosome	GO:0005694				
2.07	nuclear lumen	GO:0031981				
2.33	nucleoplasm	GO:0005654				
2.34	cell cycle process	GO:0022402				
2.38	Mitotic G1 phase and G1/S transition	REAC:R-HSA-453279				
3.01	mitotic cell cycle phase transition	GO:0044772				
3.02	chromosome organization	GO:0051276				
3.02	cell cycle phase transition	GO:0044770				
3.03	chromosomal region	GO:0098687				

Filter settings

g1rev_settests_hitsummary_gprofiler
- min_p: (0.00 <= min_p <= 0.20)