

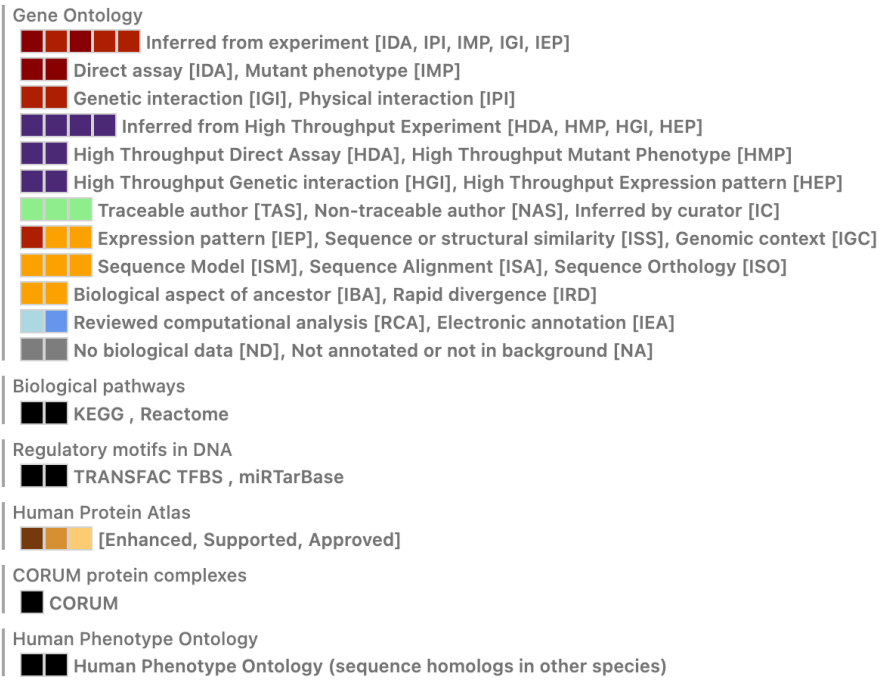
GO:BP		stats				TP53	PIK3CA	KMT2D	PTEN	L7R
Term name	Term ID	P <sub>adj</sub>	-log <sub>10</sub> (P <sub>adj</sub> )		0	16				
regulation of lymphocyte apoptotic process	GO:0070228	1.574×10 <sup>-4</sup>								
lymphocyte apoptotic process	GO:0070227	3.727×10 <sup>-4</sup>								
regulation of fibroblast apoptotic process	GO:2000269	4.882×10 <sup>-4</sup>								
regulation of leukocyte apoptotic process	GO:2000106	5.210×10 <sup>-4</sup>								
developmental growth	GO:0048589	6.375×10 <sup>-4</sup>								
fibroblast apoptotic process	GO:0044346	6.820×10 <sup>-4</sup>								
leukocyte apoptotic process	GO:0071887	1.158×10 <sup>-3</sup>								
positive regulation of signal transduction	GO:0009967	1.432×10 <sup>-3</sup>								
negative regulation of macroautophagy	GO:0016242	2.136×10 <sup>-3</sup>								
regulation of cell population proliferation	GO:0042127	2.333×10 <sup>-3</sup>								
positive regulation of cell communication	GO:0010647	2.638×10 <sup>-3</sup>								
positive regulation of signaling	GO:0023056	2.670×10 <sup>-3</sup>								
growth	GO:0040007	3.300×10 <sup>-3</sup>								
cell population proliferation	GO:0008283	5.279×10 <sup>-3</sup>								
positive regulation of response to stimulus	GO:0048584	8.056×10 <sup>-3</sup>								
peptidyl-amino acid modification	GO:0018193	1.041×10 <sup>-2</sup>								
cellular response to environmental stimulus	GO:0104004	1.075×10 <sup>-2</sup>								
cellular response to abiotic stimulus	GO:0071214	1.075×10 <sup>-2</sup>								
negative regulation of apoptotic process	GO:0043066	1.097×10 <sup>-2</sup>								
negative regulation of programmed cell death	GO:0043069	1.185×10 <sup>-2</sup>								
negative regulation of autophagy	GO:0010507	1.251×10 <sup>-2</sup>								
positive regulation of cell population proliferation	GO:0008284	1.425×10 <sup>-2</sup>								
negative regulation of cell death	GO:0060548	1.727×10 <sup>-2</sup>								
chromatin organization	GO:0006325	1.816×10 <sup>-2</sup>								
circadian behavior	GO:0048512	1.940×10 <sup>-2</sup>								
rhythmic behavior	GO:0007622	2.126×10 <sup>-2</sup>								
regulation of T cell apoptotic process	GO:0070232	2.254×10 <sup>-2</sup>								
regulation of cell aging	GO:0090342	2.422×10 <sup>-2</sup>								
regulation of generation of precursor metabolites and e...	GO:0043467	3.057×10 <sup>-2</sup>								
regulation of cellular component size	GO:0032535	3.692×10 <sup>-2</sup>								
regulation of DNA metabolic process	GO:0051052	3.723×10 <sup>-2</sup>								
negative regulation of pentose-phosphate shunt	GO:1905856	3.774×10 <sup>-2</sup>								
negative regulation of G1 to G0 transition	GO:1903451	3.774×10 <sup>-2</sup>								
regulation of transcription from RNA polymerase II prom...	GO:1990248	3.774×10 <sup>-2</sup>								
multicellular organism growth	GO:0035264	3.958×10 <sup>-2</sup>								
regulation of macroautophagy	GO:0016241	4.065×10 <sup>-2</sup>								
regulation of signal transduction	GO:0009966	4.148×10 <sup>-2</sup>								
cellular response to chemical stimulus	GO:0070887	4.739×10 <sup>-2</sup>								
T cell apoptotic process	GO:0070231	4.922×10 <sup>-2</sup>								

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REAC		stats				TP53	PIK3CA	KMT2D	PTEN	L7R
Term name	Term ID	P <sub>adj</sub>	-log <sub>10</sub> (P <sub>adj</sub> )		0	16				
PIP3 activates AKT signaling	REAC:R-HSA-1...	9.140×10 <sup>-3</sup>								
Ovarian tumor domain proteases	REAC:R-HSA-5...	1.329×10 <sup>-2</sup>								
Intracellular signaling by second messengers	REAC:R-HSA-9...	1.420×10 <sup>-2</sup>								
Synthesis of PIPs at the plasma membrane	REAC:R-HSA-1...	2.502×10 <sup>-2</sup>								
Regulation of TP53 Expression	REAC:R-HSA-6...	3.404×10 <sup>-2</sup>								

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The colors for different evidence codes in the table:



The colors for log scale:

