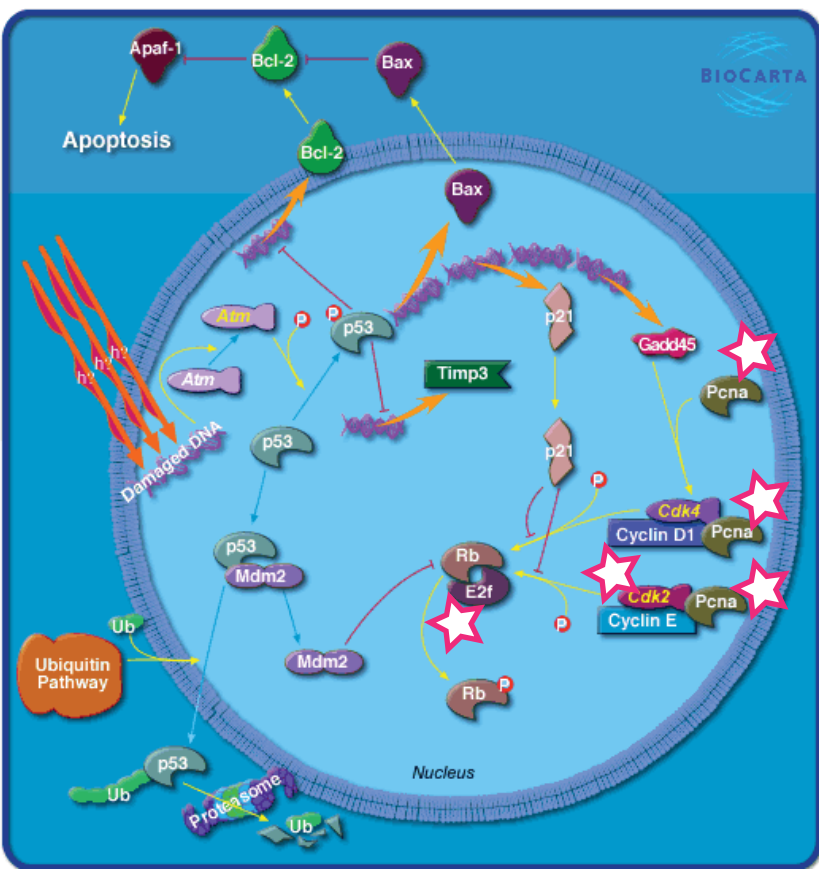


A

Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
<input type="checkbox"/>	KEGG_PATHWAY	DNA replication	RT		4	9.5	5.7E-5	1.7E-3
<input type="checkbox"/>	BIOCARTA	p53 Signaling Pathway	RT		3	7.1	1.8E-4	2.4E-3
<input type="checkbox"/>	KEGG_PATHWAY	Mismatch repair	RT		3	7.1	1.1E-3	1.6E-2
<input type="checkbox"/>	KEGG_PATHWAY	Cell cycle	RT		4	9.5	2.6E-3	2.6E-2
<input type="checkbox"/>	KEGG_PATHWAY	Nucleotide excision repair	RT		3	7.1	4.1E-3	3.0E-2
<input type="checkbox"/>	BIOCARTA	Cyclin E Destruction Pathway	RT		2	4.8	1.5E-2	1.0E-1
<input type="checkbox"/>	BIOCARTA	E2F1 Destruction Pathway	RT		2	4.8	1.7E-2	7.7E-2
<input type="checkbox"/>	BIOCARTA	Regulation of p27 Phosphorylation during Cell Cycle Progression	RT		2	4.8	2.0E-2	7.0E-2
<input type="checkbox"/>	BIOCARTA	Cyclins and Cell Cycle Regulation	RT		2	4.8	3.9E-2	1.1E-1
<input type="checkbox"/>	BIOCARTA	Influence of Ras and Rho proteins on G1 to S Transition	RT		2	4.8	4.1E-2	9.2E-2
<input type="checkbox"/>	BIOCARTA	Cell Cycle	RT		2	4.8	4.2E-2	8.3E-2
<input type="checkbox"/>	PANTHER_PATHWAY	P00017:DNA replication	RT		2	4.8	4.5E-2	3.1E-1
<input type="checkbox"/>	KEGG_PATHWAY	Purine metabolism	RT		3	7.1	4.8E-2	2.5E-1
<input type="checkbox"/>	KEGG_PATHWAY	Base excision repair	RT		2	4.8	8.7E-2	3.7E-1
<input type="checkbox"/>	PANTHER_PATHWAY	P00025:Hedgehog signaling pathway	RT		2	4.8	9.0E-2	3.1E-1

B



C

Trawler



Jaspar

Logo for MA0033.1

