DNA-binding transcription factor binding

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positive regulation of cytokice production transcription coregulator activite stablishment	unation of cellular component biogenesis of organelle localization	
transcription obligatator activity	cellular component disassembly	
noacyl–tR © A biosynthesis regulation of int ac ellular positiv establishment of p		-log(p.adjust, 10)
	positive regulation of cellular catabolic p	20
proteasebinding	histone modification ubiquitin protein iga proteasomal protein catabolic process	1
of Fibrin Clot (Clotting Cascade)	proteasomal protein catabolic process	class
rix <mark>structural constituent</mark> Platelet degranulation		metabolomics
acute inflammatory response	mRNA processing	oproteomics
kage to MARK bipleginal the respiny of vector intersection of the control of the	action with symposportse to endoplasinic r	-o- transcriptome
Diseases of signal transduction by growth factor receptors and tra		
protein activation Castalement cascade	ABC transport	jaccard_index
cytolysis	7.20 1.50 1.50	0.2
Plasma lipopr <mark>ote</mark> in remodeling Chylomicr or assembly	GTPase requiator activi	0.3
	nine, tyrosine and tryptophan biosynthes	0.4
HDL remodeling	Caffeine netabolism	
regulation of hagocytosis	Phenylalanine metabolism	
D-Amino ad metabolism		
phosphata	se binding	