## -log10(adj.p) 0 2 4 6 8 10

## threshholded to number of paths c2

	threshholded to number of paths c2								
	84	29	30	8	37	13	17	9	REACTOME_METABOLISM_OF_RNA
	62	21	30	6	17	14	16	2	REACTOME_CELL_CYCLE
	50	16	23	4	13			2	REACTOME_CELL_CYCLE_MITOTIC
	11	8			2		1	0	REACTOME_CHROMOSOME_MAINTENANCE
	26	15	10	4	8	11	8		REACTOME_TRANSCRIPTIONAL_REGULATION_BY_TP53
	3	7	2	2	1	5			REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_DNA_REPAIR_GENES
	3	6	2	2	1				REACTOME_RNA_POLYMERASE_I_TRANSCRIPTION
	5	8	3		2				REACTOME_DUAL_INCISION_IN_TC_NER
	7	8	4		3				REACTOME_TRANSCRIPTION_COUPLED_NUCLEOTIDE_EXCISION_REPAIR_TC_NER
	4	6	3		1				REACTOME_TRANSCRIPTION_OF_THE_HIV_GENOME
	25	14	12	3	13	5		6	REACTOME_RRNA_PROCESSING
	22	15	10	3	11		1	6	REACTOME_EUKARYOTIC_TRANSLATION_INITIATION
	14	12	6	2	8			5	WP_CYTOPLASMIC_RIBOSOMAL_PROTEINS
	51	19	27	5	19		5	7	REACTOME_TRANSLATION
	39		11	3	17	5	11	3	REACTOME_PROCESSING_OF_CAPPED_INTRON_CONTAINING_PRE_MRNA
	78		25	7	35	17	18		REACTOME_RNA_POLYMERASE_II_TRANSCRIPTION
	16	4	2		14	3		1	WP_PROLACTIN_SIGNALING_PATHWAY
	13	3	1		12	2		1	WP_KIT_RECEPTOR_SIGNALING_PATHWAY
	14	2	1		13	1		1	PID_IL2_1PATHWAY
	6		6						PID_NFKAPPAB_CANONICAL_PATHWAY
	12	3	1		1	2	10	1	WP_16P112_PROXIMAL_DELETION_SYNDROME
	10	1	1		1	1	8		REACTOME_ASSOCIATION_OF_TRIC_CCT_WITH_TARGET_PROTEINS_DURING_BIOSYNTHESIS
	9		2				7		REACTOME_FORMATION_OF_TUBULIN_FOLDING_INTERMEDIATES_BY_CCT_TRIC
	8		1				7		REACTOME_COOPERATION_OF_PDCL_PHLP1_AND_TRIC_CCT_IN_G_PROTEIN_BETA_FOLDING
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		a		crispr_highPaths		inpath_highPaths		dd	
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