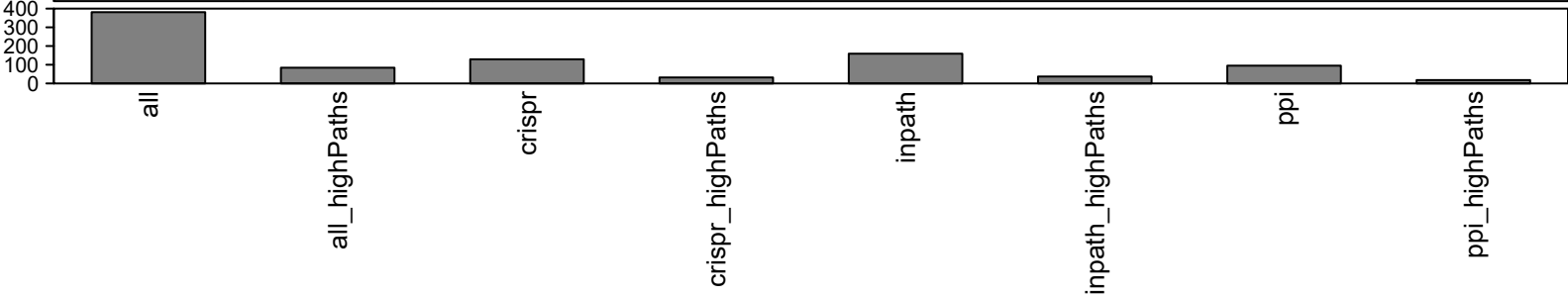


-log10(adj.p)
0 5 10
thresholded to number of paths ipap

51	13	10	4	28	4	13	6	Myc_Targets_V1_MSigDB-Hallmark-2020
31	8	10	3	11	4	10	2	E2F_Targets_MSigDB-Hallmark-2020
33	18	14	3	17	7	2	8	EIF2_Signaling
11	11	6	2	3	8	2	1	Hereditary_Breast_Cancer_Signaling
2	6	2	3		3			Nucleotide_Excision_Repair_Pathway
6	7	4	4	2	3			NER_(Nucleotide_Excision_Repair;_Enhanced_Pathway)
30	11	10	2	13	7	7	2	MicroRNA_Biogenesis_Signaling_Pathway
12	8	4		8	7		1	Thyroid_hormone_signaling_pathway_hsa04919
10	7	4		6	6		1	Prostate_Cancer_Signaling
13	7	2		7	7	4		WNT/β-catenin_Signaling
18	3	5		13	3			BAG2_Signaling_Pathway
22	4	4	0	15	3	3	1	Spliceosome_hsa03040
12	3	2	1	8	1	2	1	Myc_Targets_V2_MSigDB-Hallmark-2020
13	5	1		12	3		2	ErbB_signaling_pathway_hsa04012
25	4	14	2	10	1	1	1	NOD-like_receptor_signaling_pathway_hsa04621
6	1	6	1	0				Kinetochores_Metaphase_Signaling_Pathway
12	1	1	1	4		7		mRNA_surveillance_pathway_hsa03015



Group Sizes