

CHROMOSOME_2
CHROMOSOME_1

P-value T term ID term domain and name

Each column represents one gene list of the multi-query. Each line stands for a functional term. Each table cell stands for one list/term association.

[#] shows number of annotated genes in this list/term.

Framed cells indicate highest statistical strength in this list/term.

Colour shows strength of statistics.

CHROMOSOME	P-value	T	term ID	term domain and name
1	6	3.13e-06	132	BP appendage development (1)
1	6	2.39e-06	126	BP appendage morphogenesis (2)
1	6	3.13e-06	132	BP limb development (2)
1	6	2.39e-06	126	BP limb morphogenesis (3)
5	2.20e-05	110	BP embryonic appendage morphogenesis (1)	
5	2.20e-05	110	BP embryonic limb morphogenesis (2)	
4	4.91e-06	37	BP embryonic digit morphogenesis (1)	
13	4.42e-08	903	BP macromolecular complex subunit organization (1)	
11	1.77e-14	149	BP protein-DNA complex subunit organization (2)	
12	1.76e-09	553	BP cellular macromolecular complex subunit organization (2)	
15	2	5.64e-08	1298	BP cellular component biogenesis (1)
11	2.20e-13	187	BP DNA conformation change (1)	
11	4.19e-14	161	BP DNA packaging (2)	
14	2	1.20e-07	1172	BP cellular component assembly (1)
13	1.68e-08	832	BP macromolecular complex assembly (2)	
13	2	2.32e-08	855	BP cellular component assembly at cellular level (2)
12	4.57e-10	491	BP cellular macromolecular complex assembly (3)	
11	1.12e-14	143	BP protein-DNA complex assembly (4)	
11	1.04e-07	641	BP chromosome organization (1)	
11	8.75e-09	503	BP chromatin organization (2)	
11	6.92e-15	137	BP nucleosome organization (3)	
11	1.13e-13	176	BP chromatin assembly or disassembly (3)	
11	4.18e-15	131	BP chromatin assembly (4)	
11	2.95e-15	127	BP nucleosome assembly (5)	

p-value for best gene list

click to visualise hierarchy of terms

term type or GO domain

term name, left alignment and (#) according to depth in local hierarchy

total # genes associated to functional term

term ID