

1

2

3

4

Name	E Score	410	420	430	Filter Options
SYK_GLOVI	1e-19	N P Y P Y R Y E	. R T H M A . . G D L Q A K		E Score: e-5 <input type="text"/>
666876	2e-19	T Q I C K I K S			Redundancy Cutoff: 100 <input type="text"/>
67920132	2e-19	N G E E V E V D			Superkingdom: All
23130228	3e-19	A D L A S G E E			Archaea
57159018	3e-19	M I D K V	. . . Y C . . A D V T P E		Bacteria
1N9W	4e-19	R V L V R D L K			Eukaryota
46199389	5e-19	R V L V R D L K			
SYK_SYNY3	5e-19	R D L S N G E E			Kingdom: All
SYK_SYNEL	1e-18	A H L A A G E A			Fungi
SYK_STRMU	1e-18	D P F G K R F E	. R T A T S . . . G Q L K E K Y A D K T K E E L H		Metazoa
50256771	1e-18	E E V I D M P A			Viridiplantae
57227974	1e-18	E E V I D M P A			
68179432	3e-18	A A A L E G C E			Phylum: All
SYK_PROMA	4e-18	P N G Q D R E I			Actinobacteria
55738646	5e-18	D P F G K R F E	. R T A T S . . . G Q L K E K Y A D K T K E E L H		Aquificae
SYK_STRR6	5e-18	K Y A N L D K E			Arthropoda
55820759	5e-18	D P F G K R F E	. R T A T S . . . G Q L K E K Y A D K T K E E L H		Ascomycota
SYK_STRPN	6e-18	K Y A N L D K E			Bacteroidetes
15900610	6e-18	K Y A N L D K E			Chlamydiae
62526807	6e-18	D P F G K R F E	. R T A T S . . . G Q L K E K Y A D K T K E E L H		Apply Filter
SYK1_SALTI	6e-18	E E L E A L N I			
SYK_ENTFA	8e-18	Y D N H T K E E			
56707357	8e-18	E L E E L D N K			