

E-value scores of PFAM structural domains search

	HAP4L	bZIP
ScHAP4	1.6e-07	-
ScYAP7	1,00E-06	3.1e-08
ScYAP5	-	1.5e-06
CgAP7	-	5.6e-05
CgAP5	-	2.3e-07
KLTH	1.2e-05	1.6e-06
KLLA	1.5e-05	3.4e-07
LAKL	7,00E-07	2.9e-08
HAP43	7.3e-09	2.8e-05
HapX	1.3e-08	2.7e-05