

Additional file 1: AA2 Two Hybrid Screen Results using human cDNA library

Gene Name (Best Match)	Start	Stop	Frame	Orientation	Global PBS
hRAB6A; NM_198896.1	3	537	IF	Sense	D
hRAB6B; NM_016577.2	39	616	IF	Sense	D
hSNX5; NM_152227.1	-109	743	IF	Sense	E

PBS, Predicted Biological Score