

Additional file 2: AA2 Two Hybrid Screen Results using drosophila cDNA library

Gene Name (Best Match)	Start	Stop	Frame	Orientation	Global PBS
Drosophila melanogaster - BtbVII ; GID: 24656485	1086	1911	IF	Sense	E
Drosophila melanogaster - Dlc90F ; GID: 24647918	51	ND	IF	Sense	D
Drosophila melanogaster - Hsc70-4 ; FBgn0001219	1293	1898	IF	Sense	D
Drosophila melanogaster - Rab6 ; FBgn0015797	3	834	IF	Sense	A
Drosophila melanogaster - Rab6 ; FBgn0015797	18	ND	IF	Sense	A
Drosophila melanogaster - Rab6 ; FBgn0015797	39	1015	IF	Sense	A
Drosophila melanogaster - Rab6 ; FBgn0015797	39	ND	IF	Sense	A
Drosophila melanogaster - mor ; FBgn0002783	909	2387	IF	Sense	C
Drosophila melanogaster - mor ; FBgn0002783	909	ND	IF	Sense	C

ND : no data

PBS, Predicted Biological Score