

Sequence ID	Start	Alignment	End	Organism								
	1	50	100	150	200	250	300	350	400	440		
NP_848862.1 (+)	1										440	Mus musculus
XP_031758963.1 (+)	1										440	Xenopus tropicalis
XP_012998768.1 (+)	4	L-gulonolactone oxidase FAD lactone ox									491	Cavia porcellus
XP_002931724.3 (+)	66	LOW QUALITY PROTEIN: L-gul... FAD lactone ox FAD binding 4	LOW QUAL.	LOW QUALITY PROTEIN: L-gulonolactone oxidase-like	FAD lacton...	FAD lactone ox					483	Xenopus tropicalis
NP_919417.1 (+)	66	probable ...probable D-lac.. GlcD FAD bindi..	proba.. GlcD FAD binding 4	prob... GlcD FAD_b..	probable.. GlcD FAD-oxi...	probabl.. GlcD FAD-o..	probable D-l.. GlcD FAD-oxidasa...	probable D-la.. GlcD FAD-oxidase...	probab.. GlcD FAD...		483	Homo sapiens
XP_003472138.1 (+)	66										483	Cavia porcellus
EAW95665.1 (+)	66										494	Homo sapiens
NP_705690.2 (+)	66										506	Homo sapiens
XP_024305946.1 (+)	66										523	Homo sapiens
XP_002936654.1 (+)	95										519	Xenopus tropicalis
XP_004917918.1 (+)	127										551	Xenopus tropicalis
6LPN_A (+)	45										469	Homo sapiens
XP_013010325.1 (+)	66										512	Cavia porcellus
AAH36604.2 (+)	91										515	Homo sapiens
NP_689996.4 (+)	91										515	Homo sapiens
XP_003474407.1 (+)	108										532	Cavia porcellus
XP_024305943.1 (+)	66										546	Homo sapiens
XP_011510038.1 (+)	91										529	Homo sapiens
XP_013010326.1 (+)	66										440	Cavia porcellus
XP_011510037.1 (+)	91										541	Homo sapiens
AAM50322.1 (+)	66										485	Homo sapiens
XP_024308870.1 (+)	25										479	Homo sapiens
XP_011510036.1 (+)	91										555	Homo sapiens
XP_024305947.1 (+)	121										503	Homo sapiens
KAE8609327.1 (+)	1										368	Xenopus tropicalis
KAE8609324.1 (+)	66										406	Xenopus tropicalis
XP_024305945.1 (+)	121										526	Homo sapiens
KAE8609325.1 (+)	66										391	Xenopus tropicalis
NP_001274178.1 (+)	1										381	Homo sapiens
BAG52828.1 (+)	1										381	Homo sapiens
XP_024305944.1 (+)	121										543	Homo sapiens
4BCA_A (+)	200										654	Cavia porcellus
EAX11057.1 (+)	78										532	Homo sapiens
XP_005007711.2 (+)	200										654	Cavia porcellus
XP_011510343.1 (+)	110										564	Homo sapiens
4BC7_A (+)	200										654	Cavia porcellus
NP_003650.1 (+)	200										654	Homo sapiens
EAX11058.1 (+)	226										680	Homo sapiens
XP_024305942.1 (+)	121										566	Homo sapiens
XP_012997896.1 (+)	172										629	Cavia porcellus
KAE8604466.1 (+)	95										444	Xenopus tropicalis
XP_016860317.1 (+)	91										437	Homo sapiens
XP_031758405.1 (+)	1										363	Xenopus tropicalis
NP_001120301.1 (+)	174										622	Xenopus tropicalis
KAE8609326.1 (+)	66										360	Xenopus tropicalis
KAE8580862.1 (+)	211										659	Xenopus tropicalis
KAE8609328.1 (+)	66										334	Xenopus tropicalis
XP_011510039.1 (+)	91										463	Homo sapiens
KAE8609330.1 (+)	66										264	Xenopus tropicalis
KAE8604471.1 (+)	95										274	Xenopus tropicalis
XP_011510047.1 (+)	91										398	Homo sapiens
KAE8609329.1 (+)	1										317	Xenopus tropicalis
XP_011510052.1 (+)	91										398	Homo sapiens
XP_011510054.1 (+)	91										283	Homo sapiens
XP_011510051.1 (+)	91										283	Homo sapiens
XP_016860319.1 (+)	91										381	Homo sapiens
XP_016860321.1 (+)	91										291	Homo sapiens
XP_011510049.1 (+)	91										283	Homo sapiens
XP_011510043.1 (+)	91										283	Homo sapiens
XP_011510062.1 (+)	91										283	Homo sapiens
XP_011510048.1 (+)	91										350	Homo sapiens
BAG65519.1 (+)	91										283	Homo sapiens
XP_011510055.1 (+)	91										283	Homo sapiens
XP_011510060.1 (+)	91										283	Homo sapiens
BAG52404.1 (+)	91										310	Homo sapiens
XP_011510059.1 (+)	91										283	Homo sapiens
XP_011510053.1 (+)	91										283	Homo sapiens
EAW71291.1 (+)	91										310	Homo sapiens
XP_024308871.1 (+)	91										283	Homo sapiens
XP_016860316.1 (+)	91										283	Homo sapiens
XP_011510045.1 (+)	91										283	Homo sapiens
XP_011510040.1 (+)	91										283	Homo sapiens
XP_011510058.1 (+)	91										364	Homo sapiens
NP_001339753.1 (+)	4										328	Homo sapiens
XP_011510345.1 (+)	11										409	Homo sapiens
BAH14611.1 (+)	70										294	Homo sapiens
AAH71598.1 (+)	91										277	Homo sapiens
KAE8609331.1 (+)	2										263	Xenopus tropicalis
KAE8604472.1 (+)	20										307	Xenopus tropicalis
AAY14986.1 (+)	200										331	Homo sapiens
BAE45721.1 (+)	26										396	Homo sapiens
BAH14790.1 (+)	74										444	Homo sapiens
BAH13803.1 (+)	74										444	Homo sapiens
NP_055577.1 (+)	115										485	Homo sapiens
XP_003464117.1 (+)	115										497	Cavia porcellus
XP_031755958.1 (+)	115										498	Xenopus tropicalis
NP_001016800.1 (+)	115										498	Xenopus tropicalis
EAW956												