



gi		50659080	ref	NP_001076.2	TQNIFFMSKVTNPQA	423
gi		114654577	ref	XP_510145.2	TQNIFFMSKVTNPQA	448
gi		31340900	ref	NP_777193.2	TQSIIFLGKVTNPSEA	411
gi		126165236	ref	NP_001075213.1	TQSIIFLGKVTNPSEA	411
gi		84000377	ref	NP_001033293.1	TQSIIFLGKVTNPSEA	411
gi		119914040	ref	XP_001252712.1	-----	299
gi		119903758	ref	XP_001254254.1	-----	276
gi		126165264	ref	NP_001075181.1	TQSIIFLGKVTNPQA	418
gi		75832097	ref	NP_001012283.2	TQSIIFLGKVTNPQA	417
gi		194676990	ref	XP_001252647.2	DPSFLWLS-----	405
gi		194677056	ref	XP_001787316.1	TQSIIFLGKVTNPQA	382
gi		73964432	ref	XP_537546.2	TEIILFWGKVTNPQA	421
gi		130503301	ref	NP_033278.2	TEIAPFLAKIANPK--	418
gi		13928716	ref	NP_113719.1	TEIAPFLAKIFNPK--	408
					.....460.....	

