

	1	10	20	30	40	50
NP_462349.1
NP_462349.1
NP_250790.1
NP_00348919.1
NP_00350795.1
NP_008406338.1
NP_003494706.1
NP_003494908.1
NP_215668.1
NP_00363767.1

[illegible]

	120	130	140
NP_062949.1	ALP	SVQKAGKSPFEYCRYVYLSEYFGFIHYH	
NP_01278659.1	RUVSEFNK.		
NP_20790.1	EAFSTAAALRGDEKMLTMR.		
NP_00394819.1	SALDGFALRGDEKMGD.		
NP_003950795.1	ETLAE..PA..		
NP_008406338.1	QVAAAAHPAARG..		
NP_003949076.1	AVLDSTYGEDR.		
NP_003949208.1	ACRAEYVATDSQAQTHP.		
NP_215668.1	HVPDD.		
NP_010636767.1	EALRAAAYTR..		