

XP_791566.1	MIGQAKISAFFSEKQVKRPLSSIENSQSPS-----PK-----KAKAVKDG-----	40
human_UNG2	MIGQKTLYSFFSESPARKRHAPSPEPAVQGTGVAGVPEESGDAAAIAPKKAPAGQEEPQT	60
XP_791566.1	-KSTGLSPEQKHRMEEKKKDAMEKLMKKTEISPINSIKNGPDGICASWAKALSSEFSKP	99
human_UNG2	PPSSPLSAEQLDRIQ-----RNKAAALLRLAARNVPVGFESWKKHLSGEFGKP	109
XP_791566.1	YFIKLQEFVAAERRSHITYPSSHQVFSWTQACDIKDVKVVLGQDPYHGPGQAHGLCFSV	159
human_UNG2	YFIKLMGFVAAERKHHTVYPSPHQVFTWTQACDIKDVKVVLGQDPYHGPNQAHGLCFSV	169
	***** **	
XP_791566.1	QKGVKPPPSLANMYKCLMKDIDGFEHPGHCHLIGWSKQGVLLLLNAVLTVRANNPNSHKDK	219
human_UNG2	QRPVPPPSLENITKELSTDIEDFVHPGHCDLSGWAKQGVLLLLNAVLTVRAHQANSHKER	229
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XP_791566.1	GWEKLTAVIAWLNSNLNGVVFMLWGSYAQKKGSTIINSKRHHVLKAVHPSPLSAHRGFFD	279
human_UNG2	GWEQFTDAVVSWLNLNSNLVFLWGSYAQKKGSATDRKRHHVLQTAHPSPLSVYRGFFG	289
	** ** *	
XP_791566.1	CGHFSRTNKLLOKAGKEPIDWTRLSDEN	307
human_UNG2	CRHFSKTNELLQKSGKKPIDWKEL----	313