

1	MKLITSAICL	LSLAILTQAM	PRARRDSCDS	SSGTVCGSNG	RTYHNECDVE
51	DDSSRSGVSI	SVNHKGSK	TTFQVPRAPS	RPAPRAPYAP	RVPVAPRAPY
101	APPVPAAPRA	PYAPRVPPVP	RAPLPPARAP	VVPPRAPLPP	ARAPVVPPRA
151	PLPRAPGFAP	AAPVPGVDAD	VDVTVTEDYL	DNSADGGNSD	SGSSEEEYEYE
201	STGGDTTDGT	DGTDGIDGVT	NGLNVVFPVA	PANAPVPDVD	VTTEYENAA
251	DGEYEYGSDE	DSSEEEYGTS	IVVGGTDGTD	GNTSEASTG	GVVVPGTSAG
301	IPGFPGVSSP	SRPRAPTFGS	PALPRVPFPA	PRAPVVAPPL	IPRAPAPPTA
351	PGAPYAPSAP	LPPQTPYAPP	APRVPPQVPQT	PYTPQAPRAP	QTPYAPQAPR
401	APQTPYAPQA	PSAPQTPYAP	QAPSAPQTPY	APQAPSAPQT	PYAPQAPYAT
451	GS				