

truncated site

		10	20	30	40	50	60	70
1		ATGACGCGCCGAAAAACAGAGTATGCGCGTGCAATTCGGAGGCAGGTAAACTGCGCAAGGTCATGGTATGCAGTCCG						
1		M S A E K Q K Y G V H S E A G K L R K V M V C S P						
		85	95	105	115	125	135	145
76		GGTCTGGCACACAAACGTTTTCACCCCATCTAATTTGTGACGAACTGTTGTTTGTATGACGTCACTTTGGGTAGATCAG						
26		G L A H K R L T P S N C D E L L F D D V I W V D Q						
		160	170	180	190	200	210	220
151		GCCAAACCGGACCACTTTTGTATTTTCGTGACGAAATGCGGGAACGAGCGTGGACGTTCTGGAGATGCACAAITTA						
51		A K R D H F D F V T K M R E R G V D V L E M H N L						
		235	245	255	265	275	285	295
226		CTCACCGATATCGTTCAGCAACCGGAAGCGCTTAAATGGATTCTGGACCGTAAATCACAAGTGATACCGTCGGGA						
76		L T D I V Q Q P E A L K W I L D R K I T S D T V G						
		310	320	330	340	350	360	370
301		GTAGCGCTGACCAACGAAGTGCATCGTGCGTCTGAAGGTCTAGAGCCAGCTCATTGGCCAGAGTITTTTAAITGGC						
101		V G L T N E V R S W L E G L E P R H L A E F L I G						
		385	395	405	415	425	435	445
376		GGTGTGCTGTGTCAGGATCTGCCGGTTTCGGAAGGGGCGAGGTCATTAAATGTATAATAGTACCTGGGGCCAC						
126		G V A G Q D L P V S E G A E V I K M Y N K Y L G H						
		460	470	480	490	500	510	520
451		AGCTCGITTAICTTACCGCCACTCCCGAACACGCAATTCACTCGCGATACCACTGCTGGATTATGCGGGAGTG						
151		S S F I A L P P L P N T Q F T R D T T C W I Y G G V						
		535	545	555	565	575	585	595
526		ACGCTGAACCCCTATGTACTGGCCCGCACCGCGTCAGGAACCTTCTAACTACAGTATCTATAAATTTTCATAAG						
176		T L N P M Y W P A R R Q E T L L T T A I Y K F H K						
		610	620	630	640	650	660	670
601		GAGITTCACGGGTCCGGATTTTCAAGTGTGGTACGGGATCCTGACARAAGATCACGGCAATGCCACCCCTGGAAGGC						
201		E F T G A D F Q V W Y G D P D K D H G N A T L E G						
		685	695	705	715	725	735	745
676		GGTGTATGTCATGCCGGTGGGTAAAGGGAATTTGTGTTGATCGGAATGGGCGAAGCTACCAAGCGCGCATGCCATCGGC						
226		G D V M P V G K G I G V L I G M G E R T S R H A I G						
		760	770	780	790	800	810	820
751		CAGCTCGCCCAAAACCTTTTCGAAAAAGTTCGCCGCCGAGAAAATATCGTGGCGGGCTTACCGAAATCCCGTGCA						
251		Q L A Q N L L F E K G A A E K I I V A G L P K S R A						
		835	845	855	865	875	885	895
826		GCTATGCACCTTCGATACAGTITTTTCATTCGTGACCGCGAATCTGGTGACCGTITTTCCGGAAGTGGTTAAAGAG						
276		A M H L D T T V F S F C D R D L V T V F P E V V K E						
		910	920	930	940	950	960	970
901		ATTAAAGCGGTTTCATAATTACGCCAGACAGTTCTAARACCTTATGGTATGAATATCGCACCGCGAGGATGCTTCCCTT						
301		I K P F I I T P D S S K P Y G M N I A P Q D A S F						
		985	995	1005	1015	1025	1035	1045
976		TTAGAAGTCGTATCAGAGCACTGTTGGGCAAAAAGGATAAAATTCGCGGTGGTTGAAACCGGGGGAACAGCTTC						
326		L E V V S E E Q L L G K K D K L R V V E T G G N S F						
		1060	1070	1080	1090	1100	1110	1120
1051		GCGGCCGAACCGCGAGTGGGATGACGGCAATAACGTCGTAGCGCTGGAACCGGGAGTGTTATGGGTATGAT						
351		A A E R E Q W D D G N N V V A L E P G C V V I G Y D						
		1135	1145	1155	1165	1175	1185	1195
1126		CGGAATACCTACACGAACACTTTTACTCCGCAAGCGGGTATCGAAGTGATCAGATATCTGCCGCGAGCTGGGT						
376		R N T Y T N T L L R K A G I E V I T I S A G E L G						
		1210	1220	1230	1240	1250	1260	
1201		AGAGGGAGGGGCGGTGGGCATTGCATGACCTGTCCATTGTTTCGGCAGCCCTATCGATTAT						
401		R G R G G G G C H C M T C P I V R D P I D Y						