

Gene: EGFR - Sequence: NG_007726.3 Date : January 8, 2015

Exon 1 | Start: 1 | End: 334 | Length: 333

tctctctcgcattctcctcctcctctgctcctcccgatccctcctccgccgcctggtccc

tctctcctcccgccctgcctccccgcgctcggcccgcgcgagctagacgtccgggcagcc

|-239 |-229 |-219 |-209 |-199 |-18-
CCCCGGCGCAGCGCGGCCGAGCAGCCTCCGCCCCCGCACGGTGTGAGCGCCGACGCG

|-179 |-169 |-159 |-149 |-139 |-12-
GCCGAGGCGGCGGAGTCCCGAGCTAGCCCCGCGGCCGCCGCCGCCCAGACCGGACGAC

|-119 |-109 |-99 |-89 |-79 |-69
AGGCCACCTCGTCGGCGTCCGCCCCGAGTCCCCGCCTCGCCGCCAACGCCACAACCACCGC

|-59 |-49 |-39 |-29 |-19 |-9
GCACGGCCCCCTGACTCCGTCCAGTATTGATCGGGAGAGCCGGAGCGAGCTCTTCGGGGA

 |1 |11 |21 |31 |41 |51
GCAGCGATGCGACCTCCGGGACGGCCGGGCAGCGCTCCTGGCGCTGCTGGCTGCGCTC
 M R P S G T A G A A L L A L L A A L
 |1 |11

GTCTTATCTAACTATGATGCAAATAAAACCGGACTGAAGGAGCTGCCCATGAGAAATTTA
V L S N Y D A N K T G L K E L P M R N L
|121 |131

|421
CAGGgtgagaggctgggatgcccaaggctgggggttcataaatgcagacagcagttccgat
Q E
|141

ggctcccagcgagcttgtcactcaattccacctcggagaaggctttttatttttaccagt

acac

Exon 4 | Start: 671 | End: 805 | Length: 134

actcttgttcgcaccatggcatctcttttagcagaacataaatgcaagagcacatgcatc

cttcatgggaatttaaaggagctggaaagagtgtcaccgcagttccattctccgcaga

431	441	451	461	471	481
AAATCCTGCATGGCGCCGTGCGGTTTCAGCAACAACCCTGCCCTGTGCAACGTGGAGAGCA					
I	L	H	G	A	V
	R	F	S	N	N
			P	A	L
				C	N
				V	E
				S	I
		151			161

491	501	511	521	531	541
TCCAGTGGCGGGACATAGTCAGCAGTGACTTTCTCAGCAACATGTCGATGGACTTCCAGA					

Q W R D I V S S D F L S N M S M D F Q N
|171 |181

|551
ACCACCTGGGCAGCTgtaagtgtcgcatcacactatctctgcctccagctcctatgggg
H L G S C

gacagctctacagcactggggcaggggagagaagccatgttttagtaagtcacattaatca

gaaacaaaaagtagt

Exon 5 | Start: 806 | End: 874 | Length: 68

agaataagttgaaaagattgtcttcatttattgaatgtgcttaactcaggcccggaag

ggcgtcatcagtttctcatctttcactgagatatgcatctattacttttacatttcagg

|561 |571 |581 |591 |601 |611
GCCAAAAGTGTGATCCAAGCTGTCCAATGGGAGCTGCTGGGGTGCAGGAGAGGAGAACT
Q K C D P S C P N G S C W G A G E E N C
|191 |201

|621
GCCAGAAACgtaagtcagtgaaacagcctcagacccatgtgtgaccgcccctctcttcctt
Q K L

cacttgcttaggtgattggatttgtttccctctgaagactccaaagagttactttatta

cagggtcag

Exon 6 | Start: 875 | End: 993 | Length: 118

gttgaagcaaatgtgtcttcactttttcatgaaaagtctgcaagtgctctgcgacatc

cctgggaaatgatcctaccctcactcttcagctcacagggaacctttgctctttttcagt

631	641	651	661	671	681														
TGACCAAAATCATCTGTGCCCAGCAGTGCTCCGGGCGCTGCCGTGGCAAGTCCCCCAGTG																			
T	K	I	I	C	A	Q	Q	C	S	G	R	C	R	G	K	S	P	S	D
211										221									

691	701	711	721	731	741														
ACTGCTGCCACAACCAGTGTGCTGCAGGCTGCACAGGCCCCGGGAGAGCGACTGCCTGg																			
C	C	H	N	Q	C	A	A	G	C	T	G	P	R	E	S	D	C	L	
231										241									

taagatgccccctccagcagcctccctggagcaggctggggctgcacccgccccaccaca

ccaggacagaagacttcctgtgggggagctgtcaattagcatttgtcataacagacagg

Exon 7 | Start: 994 | End: 1135 | Length: 141

ggctgcaaagacagtaacttgggctttctgacgggagtcacaccgtgctgcgcttcctc

cgtgtgtggcgctgagtggtacttacctcacttggccagcgtgtcctctctcctccatagg

751	761	771	781	791	801
GTCTGCCG	AAATTCG	GAGACGA	AGCCACG	TGCAAGG	ACACCTG
V C R K	F R D E	A T C K	D T C P	P L M L	
251			261		

811	821	831	841	851	861
TACAACCC	CACGTA	CCAGATG	GATGTGA	ACCCGAG	GGCAAAT
Y N P T	T Y Q M	D V N P	E G K Y	S F G A	
271			281		

871	881
ACCTGCGT	GAAGAAG
T C V K	K C P R
291	

tccttgtcccgcctctgtctcctgctgagccctggagtatcccatcttggagagtctttgg

gtggatgtgtttgccttgcttg

Exon 8 | Start: 1136 | End: 1252 | Length: 116

ataaagccaaaggaggatggagcctttccatcacccctcaagaggacctggaccgcctgt

gtgaggcccgagcacctgggtgccaccgtcatcaccttcctttcatgctctcttccccagg

891	901	911	921	931	941														
GTAATTATGTGGTGACAGATCACGGCTCGTGCGTCCGAGCCTGTGGGGCCGACAGCTATG																			
N	Y	V	V	T	D	H	G	S	C	V	R	A	C	G	A	D	S	Y	E
			301																311

951	961	971	981	991	1001														
AGATGGAGGAAGACGGCGTCCGCAAGTGTAAGAAGTGCGAAGGGCCTTGCCGCAAAGgta																			
M	E	E	D	G	V	R	K	C	K	C	E	G	P	C	R	K	V		
			321																331

ggaagcccgcggtgtgcggacgaggcttggttctcggctgctgaggctgggctctcatgc

cacctccaaaggaacacatcttcctcttctcattaaaaaacaactatacatatcggt

Exon 9 | Start: 1253 | End: 1379 | Length: 126

atgtggcagtgggcggttccggtgaccggaattccttcctgcttcctctgcctgtggatc

cctagctattcttaatccaacaaatgtgaacggaatacacgtctctcttatctctgcagt

1011	1021	1031	1041	1051	1061																
TGTGTAACGGAATAGGTATTGGTGAATTTAAAGACTCACTCTCCATAAATGCTACGAATA																					
C	N	G	I	G	I	G	E	F	K	D	S	L	S	I	N	A	T	N	I		
										341											351

1071	1081	1091	1101	1111	1121																
TTAAACACTTCAAAAAGTGCACCTCCATCAGTGGCGATCTCCACATCCTGCCGGTGGCAT																					
K	H	F	K	N	C	T	S	I	S	G	D	L	H	I	L	P	V	A	F		
										361											371

1131
TTAGGGGgtgagtcacagggttcagttgcttggtataaagaaaaacaaaatctgccttttta
R G

actggtagagattggtgatcaataatcacctgttggttggttcagtgactccttcacac

atactcc

Exon 10 | Start: 1380 | End: 1453 | Length: 73

ctgccggtggcatttagggggtgagtcacaggttcagttgcttgataaaagaaaaacaaa

atctgcctttttaactggtagagattggtgatcaataatcacctggtgtttgtttcagt

1141	1151	1161	1171	1181	1111														
TGACTCCTTCACACATACTCCTCCTCGGATCCACAGGAACTGGATATTCTGAAAACCGT																			
D	S	F	T	H	T	P	P	L	D	P	Q	E	L	D	I	L	K	T	V
381										391									

1201
AAAGGAAATCACAGgtttgagctgaattatcacatgaatataaatgggaaatcagtgttt
K E I T G
401

tagagagagaacttttcgacatatttcctgttcccttggaataaaaacatttccttctgaa

attttaccgttaat

Exon 11 | Start: 1454 | End: 1544 | Length: 90

tccaatntttcccacttactgttcatataatacagagtccttgagagtctagagtaatgtc

tcatacaaaaaagaaactcctacgtgggtgtgtgtctgaagcttttcatctgccttacagg

1211	1221	1231	1241	1251	1261
GGTTTTTGCTGATT	CAGGCTTGGCCTG	AAAACAGGACG	GACCTCCATGC	CTTTGAGAACC	
F L L I Q A W P E N R T D L H A F E N L					
		411			421

1271	1281	1291
TAGAAATCATACG	CGGCAGGACCA	AGCAACAgtaag
E I I R G R T K Q H		ttgaccacagcca
	431	agcctggttag

attacatttgccttttttagttggaaattaggcttaacaggagagttgctaagatagggca

cagagctcctgcattctctcgccggcattccc

Exon 12 | Start: 1545 | End: 1744 | Length: 199

cctaccatcattggaaagcagttttagtcaatcaaagtggtctggagaaacaaagttt

tcagggatacattgtttttataatttttcaccacatgatttttcttctctccaatgtagt

1301	1311	1321	1331	1341	1351
GGTCAGT	TTTCTCT	TGCAGT	CGTCAGC	CTGAAC	ATAACAT
CCTTGGG	ATTACG	CTCCCT			
G	Q	F	S	L	A
V	V	S	L	N	I
T	S	L	G	L	R
S	L				
441			451		

1361	1371	1381	1391	1401	1411
CAAGG	AGATAA	GTGATG	GAGATG	TGATAA	TTTCAGG
AAACAAA	AAATTTG	TGCTATG	CAAAA		
K	E	I	S	D	G
D	V	I	I	S	G
N	K	N	L	C	Y
A	N				
461			471		

1421	1431	1441	1451	1461	1471
TACAATA	AACTG	GAATAA	CTGTTT	GGGAC	CTCCGGT
CAGAAA	ACCAAA	ATTATA	AGCAA		
T	I	N	W	K	K
L	F	G	T	S	G
Q	K	T	K	I	I
S	N				
481			491		

1481	1491
CAGAGGT	GAAAACAGCTGCA
gtaagtc	caccgctttctgttt
atgagtt	ggttct
R	G
E	N
S	C
K	

aatgggtcctttattttgtatttagaatattgaagggtattcccatttaaattacttttt

tcagttccttaagaagcaaa

Exon 13 | Start: 1745 | End: 1877 | Length: 132

catgtctgtgtcacccaaggtcatggagcacaggggcccctcccgggaaggtgccgtctcc

tccggcccctcgggtccctgctctgtcactgactgctgtgacccactctgtctccgcaga

1501	1511	1521	1531	1541	1551														
AGGCCACAGGCCAGGTCTGCCATGCCTTGTGCTCCCCGAGGGCTGCTGGGGCCCGGAGC																			
A	T	G	Q	V	C	H	A	L	C	S	P	E	G	C	W	G	P	E	P
501											511								

1561	1571	1581	1591	1601	1611														
CCAGGGACTGCGTCTCTTGCCGGAATGTCAGCCGAGGCAGGGAATGCGTGGACAAGTGCA																			
R	D	C	V	S	C	R	N	V	S	R	G	R	E	C	V	D	K	C	N
521											531								

1621	1631																		
ACCTTCTGGAGGGgtaggaggttatctttaaataccccttgcggttgatcaaaaataaggc																			
L	L	E	G																
541																			

tccaggttggtgttatagctttacaggcattctgtttgattttctcttccttttattctt

tgcccttggtttt

Exon 14 | Start: 1878 | End: 1968 | Length: 90

cctggagtcctcaactccttgaccattacctcaagttatttgaattttgaagaggtgatt

tgtgttcctgcaataatgtctcaggggtgggctgacgggtttcctcttcctcctctcagt

	1641		1651		1661		1671		1681		1196
TGAGCCAAGGGAGTTTGTGGAGAACTCTGAGTGCATACAGTGCCACCCAGAGTGCCTGCC											
E P R E F V E N S E C I Q C H P E C L P											
			551						561		

	1701		1711		1721
TCAGGCCATGAACATCACCTGCACAGGACGGgtaagagccccttgctgctatccacgtcc					
Q A M N I T C T G R					
			571		

atttcatgggaagggccttcacagaagccgaacagtgatgatggcccagggcatcctgtg

tgggcaggacggccatcagagccacttccca

Exon 15 | Start: 1969 | End: 2126 | Length: 157

tttctatcatttggctttccccactcacacacactaaatattttaagtaaaaagttactt

ccattttgaaagagaaaagaaagagacatgcatgaacatttttctccaccttggtgcagg

1731	1741	1751	1761	1771	1118
GGACCAGACAACTGTATCCAGTGTGCCCCTACATTGACGGCCCCCACTGCGTCAAGACC					
G	P	D	N	C	I
			Q	C	A
			H	Y	I
			D	G	P
			H	C	V
			K	T	
		581		591	

1791	1801	1811	1821	1831	1114
TGCCCGGCAGGAGTCATGGGAGAAAACAACCCCTGGTCTGGAAGTACGCAGACGCCGGC					
C	P	A	G	V	M
			G	E	N
			N	T	L
			V	W	K
			Y	A	D
			A	G	
		601		611	

1851	1861	1871
CATGTGTGCCACCTGTGCCATCCAACTGCACCTACGGgtgagtggaaagtgaaggagaa		
H	V	C
	H	L
	C	H
	P	N
	C	T
	Y	G
	621	

cagaacatttcctctcttgcaaattcagagatcaaaaatgtctccaagttttccggcaa

caaattgccgaggtttgtatttgagtcagttacttaag

Exon 16 | Start: 2127 | End: 2165 | Length: 38

ttttaattatttaagagtagtttagcatatattgctttatgatttaattaaaaatctcca

aaatatatgccaaagaagtagaatgagaaaaatgtatatatttctctttcacttcctacaga

|2041 |2051 |2061
 GAGGCTGCTGCAGGAGAGGGAGgtgagtgccagtcctgggtgggctcaggagccctcgca
 R L L Q E R E
 |681

ccccgacaggaacaagggccagccccgagaacgggccattagcagttgtgtatgttagat

acataattgtattatgatgcag

Exon 18 | Start: 2308 | End: 2430 | Length: 122

tcctggcacccaagcccatgccgtggctgctgggtccccctgctgggccatgtctggcact

gctttccagcatggtgagggctgaggtgacccttgctctctgtgttcttgccccccagc

	2071	2081	2091	2101	2111	2121
	CTTGTGGAGCCTCTTACACCCAGTGGAGAAGCTCCCAACCAAGCTCTCTTGAGGATCTTG					
	L	V	E	P	L	T
	P	S	G	E	A	P
	N	Q	A	L	L	R
	I	L				
	691			701		
	2131	2141	2151	2161	2171	2181
	AAGGAAACTGAATTCAAAAAGATCAAAGTGCTGGGCTCCGGTTCGGTTCGGCACGGTGTAT					
	K	E	T	E	F	K
	K	I	K	V	L	G
	S	G	A	F	G	T
	V	Y				
	711			721		

AAGtaaggTccctggcacaggcctctgggctgggccgcagggcctctcatggtctggtg
K

gggagcccagagtccttgcaagctgtatatttccatcatctactttactctttgtttcac

tga

Exon 19 | Start: 2431 | End: 2529 | Length: 98

agccccagtgTccctcaccttcggggTgcatcgctggtaacatccaccagatcactggg

cagcatgtggcaccatctcacaattgccagttaacgtcttccttctctctgtcatagg

2191	2201	2211	2221	2231	2242
GGACTCTGGATCCCAGAAGGTGAGAAAAGTTAAAATTCCCGTCGCTATCAAGGAATTAAGA					
G L W I P E G E K V K I P V A I K E L R					
731			741		
2251	2261	2271	2281		
GAAGCAACATCTCCGAAAGCCAACAAGGAAATCCTCGATgtgagtttctgctttgctgtg					
E A T S P K A N K E I L D					
751			761		

tgggggtccatggctctgaacctcaggccaccttttctcatgtctggcagctgctctgc

tctagaccctgctcatctccacatcctaaatgttcactt

Exon 20 | Start: 2530 | End: 2715 | Length: 185

tgagtacgtattttgaaactcaagatcgcatcatgcgtcttcacctggaaggggtccat

gtgcccctccttctggccaccatgcgaagccacactgacgtgcctctccctccctccagg

2291	2301	2311	2321	2331	2321
GAAGCCTACGTGATGGCCAGCGTGGACAACCCCCACGTGTGCCGCTGCTGGGCATCTGC					
E A Y V M A S V D N P H V C R L L G I C					
		771			787

2351	2361	2371	2381	2391	2421
CTCACCTCCACCGTGCAGCTCATCACGCAGCTCATGCCCTTCGGCTGCCTCCTGGACTAT					
L T S T V Q L I T Q L M P F G C L L D Y					
		791			808

2411	2421	2431	2441	2451	2421
GTCCGGGAACACAAAGACAATATTGGCTCCCAGTACCTGCTCAACTGGTGTGTGCAGATC					
V R E H K D N I G S Q Y L L N W C V Q I					
		811			828

GCAAAGgtaatcagggaaggagatacggggaggggagataaggagccaggatcctcaca

A K

tgcggtctgcgctcctgggatagcaagagtttgccatggggatatgtgtgtgcgtgcatg

cagcac

Exon 21 | Start: 2716 | End: 2871 | Length: 155

agccataagtcctcgacgtggagaggctcagagcctggcatgaacatgaccctgaattcg

gatgcagagcttcttcccatgatgatctgtccctcacagcagggtcttctctgtttcagg

2471	2481	2491	2501	2511	2521
GGCATGAACTACTTGGAGGACCGTCGCTTGGTGCACCGCGACCTGGCAGCCAGGAACGTA					
G M N Y L E D R R L V H R D L A A R N V					
	831			841	

2531	2541	2551	2561	2571	2581
CTGGTGA AAAACACCGCAGCATGTCAAGATCACAGATTTTGGGCTGGCCAAACTGCTGGGT					
L V K T P Q H V K I T D F G L A K L L G					
	851			861	

2591	2601	2611	2621
GCGGAAGAGAAAGAATACCATGCAGAAGGAGGCAAAgtaaggaggtggcttttaggtcagc			
A E E K E Y H A E G G K			

|871

cagcattttcctgacaccagggaccaggctgccttcccactagctgtattgtttaacaca

tgcaggggaggatgctctccagacattctgggtgag

Exon 22 | Start: 2872 | End: 2947 | Length: 75

cctggggtgatctggctcgtctgtgtgtgtcactcgtaattaggtccagagtgagttaac

ttttccaacagagggaactaatagttgtctcactgcctcatctctcaccatcccaagg

2631	2641	2651	2661	2671	2681														
GTGCCTATCAAGTGGATGGCATTGGAATCAATTTACACAGAATCTATACCCACCAGAGT																			
V	P	I	K	W	M	A	L	E	S	I	L	H	R	I	Y	T	H	Q	S
											881								891

2691	2701																		
GATGTCTGGAGCTACG	gtgagtcataatcctgatgctaatagtttgtactgaggccaag																		
D	V	W	S	Y	G														
											901								

ctggcttttattgttagttaatttacattatcctctgacatgcaagtattttctttcg

agataatgactaatga

Exon 23 | Start: 2948 | End: 3094 | Length: 146

tacagaaatgtaggtttctaaacatcaagaaacagtaaccagtaatgatgactaaagcaa

gggattgtgattgttcattcatgatccactgccttcttttcttgcttcacctctcagg

2711	2721	2731	2741	2751	2167																
GGGTGACCGTTTGGGAGTTGATGACCTTTGGATCCAAGCCATATGACGGAATCCCTGCCA																					
V	T	V	W	E	L	M	T	F	G	S	K	P	Y	D	G	I	P	A	S		
										911											912

2771	2781	2791	2801	2811	2128																
GCGAGATCTCCTCCATCCTGGAGAAAGGAGAACGCCTCCCTCAGCCACCCATATGTACCA																					
E	I	S	S	I	L	E	K	G	E	R	L	P	Q	P	P	I	C	T	I		
										931											914

2831	2841							
TCGATGTCTACATGATCATGGTCAAGTgtgagtgactggtgggtctgtccacactgccta								
D	V	Y	M	I	M	V	K	C

gctgagccttggtggctgctcttagccaaacagctgaggcctttgcatccctggagaaat

gtcatcacattacttaaggcaggcaca

Exon 24 | Start: 3095 | End: 3192 | Length: 97

ttctttcatcacttatttgactggaagtgtcgcatcaccaatgccttctttaagcaatgc

catctttatcatttcttccagtgttctaattgcactgttttttctcattccttccccagg

2851	2861	2871	2881	2891	2901
GCTGGATGATAGACGCAGATAGTCGCCCAAAGTTCCGTGAGTTGATCATCGAATTCTCCA					
W M I D A D S R P K F R E L I I E F S K					
951			961		

2911	2921	2931	2941
AAATGGCCCGAGACCCCGAGCGCTACCTTGTTCATTGAGgtacaaattgcagtctgtgctt			
M A R D P Q R Y L V I Q			
971		981	

ccattgggaagagtcctctaatagagcatctcatgtcactgtgttctgtcacatgccagc

ctggcctccctgtgtcccagatcgcatattattaaaccct

Exon 25 | Start: 3193 | End: 3360 | Length: 167

ttataacaaaattggcaaacacacaggcacctgctggcaatagaccctgctcctatagc

caagaagtggaatagcatctctacgggccattctaatagcctcaaaatctctgcaccagg

2951	2961	2971	2981	2991	3001
GGGGATGAAAGAATGCATTTGCCAAGTCCTACAGACTCCAACCTTCTACCGTGCCCTGATG					
G D E R M H L P S P T D S N F Y R A L M					
		991			1001

3011	3021	3031	3041	3051	3061
GATGAAGAAGACATGGACGACGTGGTGGATGCCGACGAGTACCTCATCCCACAGCAGGGC					
D E E D M D D V V D A D E Y L I P Q Q G					
		1011			1021

3071	3081	3091	3101	3111
TTCTTCAGCAGCCCCCTCCACGTCACGGACTCCCTCCTGAGCTCTCTGgtatgaaatctc				
F F S S P S T S R T P L L S S L				
		1031		

tgtctctctctctctcaagctgtgtctactcatttgaacaaattgaattttaggghaaa

ataaccatctagtgaactcacatggatatgaagtcaattttaaccaa

Exon 26 | Start: 3361 | End: 3408 | Length: 47

agacaaaaattaaacaccttcacaatataccctccatgaggcacaccacctgcattcagg

aaaagtggatgagatgtggtacaagcattccatgggcaacttctctgtttcttttcaga

3121	3131	3141	3151	3161											
AGTGCAACCAGCAACAATTCCACCGTGGCTTGCATTGATAGAAATGGG	gtatgtatgaac														
S	A	T	S	N	N	S	T	V	A	C	I	D	R	N	G
1041															1051

accttataagccagaatttacagctctccactatggctctattttacatggaaaatgcct

taacctaaataattttaaccagataatcttgagttttcttcctgtgt

Exon 27 | Start: 3409 | End: 3517 | Length: 108

caggcctgccccaacctactaatcagaaccagcatctcaaggagatctcgggtgatttttg

caaacactgaagttggggcagccctgaccggagtaaccttccctcatttcctcctgcagc

3171	3181	3191	3201	3211	3312																
CTGCAAAGCTGTCCCATCAAGGAAGACAGCTTCTTGCAGCGATACAGCTCAGACCCACA																					
L	Q	S	C	P	I	K	E	D	S	F	L	Q	R	Y	S	S	D	P	T		
										1061											1071
3231	3241	3251	3261	3271																	
GGCGCCTTGACTGAGGACAGCATAGACGACACCTTCCTCCCAGTGCCTGgtgagtggtt																					
G	A	L	T	E	D	S	I	D	D	T	F	L	P	V	P	E					
										1081											1091

gtctggaacagtcctgctcctcaacctcctcgaccactcagcagcagccagtctccag

tgtccaagccaggtgctccctccagcatctccagagggggaacagtgg

Exon 28 | Start: 3518 | End: 5600 | Length: 2082

ctcctgctccctgtcataagtctccttggtgaggacattcacagggttcagaaccaggg

atcctgcatgggatgggtgctttgctgattacttcacctctgatttctttccactttcaga

3281	3291	3301	3311	3321	3133
AATACATAAACAGTCCGTTCCCAAAAGGCCGCTGGCTCTGTGCAGAATCCTGTCTATC					
Y	I	N	Q	S	V
			P	K	R
			P	A	G
			S	V	Q
			N	P	V
			Y	H	
		1101			1111
3341	3351	3361	3371	3381	3193
ACAATCAGCCTCTGAACCCCGCGCCAGCAGAGACCCACACTACCAGGACCCACAGCA					
N	Q	P	L	N	P
		A	P	S	R
		D	P	H	Y
		Q	D	P	H
		S	T		
		1121			1131
3401	3411	3421	3431	3441	3154
CTGCAGTGGGCAACCCGAGTATCTCAACACTGTCCAGCCACCTGTGTCAACAGCACAT					
A	V	G	N	P	E
		Y	L	N	T
		V	Q	P	T
		C	V	N	S
		T	F		
		1141			1151
3461	3471	3481	3491	3501	3115
TCGACAGCCCTGCCCACTGGGCCAGAAAGGCAGCCACCAAATTAGCCTGGACAACCCTG					
D	S	P	A	H	W
		A	Q	K	G
		S	H	Q	I
		S	L	D	N
		P	D		
		1161			1171
3521	3531	3541	3551	3561	3175
ACTACCAGCAGGACTTCTTTCCCAAGGAAGCCAAGCCAAATGGCATCTTTAAGGGCTCCA					
Y	Q	Q	D	F	F
		P	K	E	A
		K	P	N	G
		I	F	K	G
		S	T		
		1181			1191
3581	3591	3601	3611	3621	3136
CAGCTGAAAATGCAGAATACCTAAGGGTCGCGCCACAAAGCAGTGAATTTATTGGAGCAT					
A	E	N	A	E	Y
		L	R	V	A
		P	Q	S	S
		E	F	I	G
		A			
		1201			
3641	3651	3661	3671	3681	3196
GACCACGGAGGATAGTATGAGCCCTAAAAATCCAGACTCTTTCGATACCCAGGACCAAGC					
3701	3711	3721	3731	3741	3157
CACAGCAGGTCCTCCATCCCAACAGCCATGCCCGCATTAGCTCTTAGACCCACAGACTGG					
3761	3771	3781	3791	3801	3118
TTTTGCAACGTTTACACCGACTAGCCAGGAAGTACTTCCACCTCGGGCACATTTTGGGAA					

|3821 |3831 |3841 |3851 |3861 |3178
GTTGCATTCCCTTTGTCTTCAAACGTGAAGCATTACAGAAACGCATCCAGCAAGAATAT

|3881 |3891 |3901 |3911 |3921 |3139
TGTCCCTTTGAGCAGAAATTTATCTTTCAAAGAGGTATATTGAAAAAAAAAAAAAGTAT

|3941 |3951 |3961 |3971 |3981 |3199
ATGTGAGGATTTTTATTGATTGGGATCTTGGAGTTTTTCATTGTCGCTATTGATTTTA

|4001 |4011 |4021 |4031 |4041 |4150
CTTCAATGGGCTCTTCCAACAAGGAAGAAGCTTGCTGGTAGCACTTGCTACCCCTGAGTTC

|4061 |4071 |4081 |4091 |4101 |4111
ATCCAGGCCCAACTGTGAGCAAGGAGCACAAGCCACAAGTCTTCCAGAGGATGCTTGATT

|4121 |4131 |4141 |4151 |4161 |4171
CCAGTGTTCTGCTTCAAGGCTTCCACTGCAAAACACTAAAGATCCAAGAAGGCCTTCAT

|4181 |4191 |4201 |4211 |4221 |4132
GGCCCCAGCAGGCCGATCGGTACTGTATCAAGTCATGGCAGGTACAGTAGGATAAGCCA

|4241 |4251 |4261 |4271 |4281 |4192
CTCTGTCCCTTCTGGGCAAAGAAGAAACGGAGGGGATGGAATTCTTCCTTAGACTTACT

|4301 |4311 |4321 |4331 |4341 |4153
TTTGTAATAATGTCCCCACGGTACTTACTCCCCACTGATGGACCAGTGGTTTCCAGTCAT

|4361 |4371 |4381 |4391 |4401 |4114

GAGCGTTAGACTGACTTGTTTGTCTTCCATTCCATTGTTTTGAAACTCAGTATGCTGCCC

|4421 |4431 |4441 |4451 |4461 |4174
CTGTCTTGCTGTCATGAAATCAGCAAGAGAGGATGACACATCAAATAATAACTCGGATTC

|4481 |4491 |4501 |4511 |4521 |4135
CAGCCCACATTGGATTCATCAGCATTTGGACCAATAGCCCACAGCTGAGAATGTGGAATA

|4541 |4551 |4561 |4571 |4581 |4195
CCTAAGGATAGCACCGCTTTTGTTCGCAAAAACGTATCTCCTAATTTGAGGCTCAGAT

|4601 |4611 |4621 |4631 |4641 |4156
GAAATGCATCAGGTCCTTTGGGGCATAGATCAGAAGACTACAAAAATGAAGCTGCTCTGA

|4661 |4671 |4681 |4691 |4701 |4117
AATCTCCTTTAGCCATCACCCCAACCCCCAAAATTAGTTTGTGTTACTTATGGAAGATA

|4721 |4731 |4741 |4751 |4761 |4177
GTTTTCTCCTTTTACTTCACTTCAAAAGCTTTTACTCAAAGAGTATATGTTCCCTCCAG

|4781 |4791 |4801 |4811 |4821 |4138
GTCAGCTGCCCCAAACCCCTCCTTACGCTTGTGTCACAAAAAGTGTCTCTGCCTTGA

|4841 |4851 |4861 |4871 |4881 |4198
GTCATCTATTCAAGCACTTACAGCTCTGGCCACAACAGGGCATTTTACAGGTGCGAATGA

|4901 |4911 |4921 |4931 |4941 |4159
CAGTAGCATTATGAGTAGTGTGGAATTCAGGTAGTAAATATGAAACTAGGGTTTGAAATT

|4961 |4971 |4981 |4991 |5001 |5110
GATAATGCTTTCACAACATTTGCAGATGTTTTAGAAGGAAAAAAGTTCCTTCCTAAAATA

|5021 |5031 |5041 |5051 |5061 |5170
ATTTCTCTACAATTGGAAGATTGGAAGATTCAGCTAGTTAGGAGCCCACCTTTTTTCCTA

|5081 |5091 |5101 |5111 |5121 |5131
ATCTGTGTGTGCCCTGTAACTGACTGGTTAACAGCAGTCCTTTGTAAACAGTGTTTTAA

|5141 |5151 |5161 |5171 |5181 |5191
ACTCTCCTAGTCAATATCCACCCCATCCAATTTATCAAGGAAGAAATGGTTCAGAAAAATA

|5201 |5211 |5221 |5231 |5241 |5152
TTTTTCAGCCTACAGTTATGTTTCAGTCACACACATACAAAATGTTTCCTTTTGCTTTTAA

|5261 |5271 |5281 |5291 |5301 |5113
AGTAATTTTGGACTCCAGATCAGTCAGAGCCCCTACAGCATTGTTAAGAAAGTATTTGA

|5321 |5331 |5341 |5351
TTTTTGTCTCAATGAAAATAAACTATATTCAATTTCCACTCTAttatgctctcaaatacc

cctaagcatctatactagcctgggtatgggtatgaaagatacaaagataaataaaacatag

tccctgattctaagaaattcacaatttagcaaaggaaatggac

