

Gene: MECP2 - Sequence: NG\_007107.2  
 Transcript: NM\_004992.3 - Protein: NP\_004983.1  
 Date : February 26, 2015

1<sup>st</sup> line: Base numbering. Full stops for intronic +/- 5, 10, 15...  
 2<sup>nd</sup> line: Base sequence. lower case Introns, upper case Exons  
 3<sup>rd</sup> line: Amino acid sequence. Printed on FIRST base of codon  
 4<sup>th</sup> line: Amino acid numbering. Numbered on 1<sup>st</sup> and increments of 10

Exon 1 | Start: 44391 | End: 44518 | Length: 127

```

. . . . .
actaaaccagtcctccgcgcccaagccgcctcttttcccaaacgacggccgaaagcag

. . . . .
ccaatcaacagctggaggggtccgcccccttttccctggccgaaatggacaggaaatctc

. . . . .
gccaatgacggcatcgccgctgagacctccccctccccgtcctccccgtcccagccc

. . . . .
ggccatcacagccaatgacgggcgggctcgcagcggcgccgagggcgggcgcgggcgcg

. . . . .
caggtgcagcagcgcgggcgccgccaagaggcgggcgcgacgtcggccgtgcgggggt

      |-219      |-209      |-199      |-189      |-179      |-169
CCGGCGTCGGCGGCGCGCGCTCCCTCCTCTCGGAGAGAGGGCTGTGGTAAAGCCGTC

      |-159      |-149      |-139      |-129      |-119      |-109
CGGAAAATGGCCGCCGCCGCCGCCGCCGAGCGGAGGAGGAGGAGGAGCGAGGAG

      |-99 . . . . .
GAGAGACTgtgagtgggaccgccgtggccgcgggcggggacccttgccggggggcggggg

. . . . .
tcagggcggggacgtggcgcgggaggggcccgcgggggtcggacgacacggctggcggatg

. . . . .
gcgtccctcctctctaccctccccctcccgccgcgcggtggcgactctccctcggcc

. . . . .
cgtcaccctgtctcgcggtgacctcctcggcggcctccctggagccgccttcgcct

. . . . .

```

gacgcccctcttcctcccgccctcgacgcgcatcccgcccccgccccgcgggcgcccc

. .  
tgtcgccg

Exon 2 | Start: 49814 | End: 49937 | Length: 123

. . . . .  
tagaagaaatacttgccagaaatcgccactcatggatgctttttagtgctgaagtgtc

. . . . .  
ccctagaggtgacaaggcttgtagatgttgattctaacaagcatgaatctttccttta

. . . . .  
ttttagcactgtgtgttacgtgccagtaatttgcagcttatcctttgtttctagctaggt

. . . . .  
aagctgggaaatagcctagtactttgtctatgtgtttatcttcaaaatgtcccaaatagc

. . . . .  
cctgggaaaaggctgtgcagctcaatgggggctttcaactacaattttctttgtttta

          |-89      |-79      |-69      |-59      |-49      |-39  
GCTCCATAAAAATACAGACTCACCAGTTCCTGCTTTGATGTGACATGTGACTCCCCAGAA

          |-29      |-19      |-9      |1      |11      |21  
TACACCTTGCTTCTGTAGACCAGCTCCAACAGGATTCCATGGTAGCTGGGATGTTAGGGC  
                          M V A G M L G L  
                          |1

. . . . .  
TCAGgtaagtaaccttccttttttttttagtatatgtcctggtttggccatctgttt  
R

. . . . .  
tttttttttttaaaaaaaaaaaaaaaggaaaaggagaaaaaatatactactcttggac

. . . . .  
agtataaaagtaccccaaagactaaagacataactgtgccaactgtgccatataataaa

. . . . .  
aaaaagtcacttccctgagccctgaaaggtcagtgtgtgtagggttacttggtcgccaca

. . . . .  
gcgtgatctggggcgggcggtcagattagagccggaactgggtgatctgcaacttcagttc

.  
acct

Exon 3 | Start: 109571 | End: 109921 | Length: 350

. . . . .  
agccgcagtgtttccgctcagaggaaagggtctgtattctcctgcagtgcctaggagactt  
. . . . .  
gtgggtggccacagtgcaggtcaggcacaccggccagcaccacccacagcccaaattcct  
. . . . .  
aaagaaatatttgggtcccagcttggcccagtcctctgttgtcctggggaaggacatcaa  
. . . . .  
gatctgagtgtatgatggcctggggccttgcatgtggtgggggtccaagcctgcctctgc  
. . . . .  
tcacttgttctgcagactggcatgttctctgtgatacttacatacttgtttaacacttca

      |31      |41      |51      |61      |71      |81  
GGAAGAAAAAGTCAGAAGACCAGGACCTCCAGGGCCTCAAGGACAAACCCCTCAAGTTTAA  
E E K S E D Q D L Q G L K D K P L K F K  
      |11                  |21

      |91      |101      |111      |121      |131      |141  
AAAGGTGAAGAAAGATAAGAAAGAAGAGAAAGAGGGCAAGCATGAGCCCGTGCAGCCATC  
K V K K D K K E E K E G K H E P V Q P S  
      |31                  |41

      |151      |161      |171      |181      |191      |201  
AGCCCACTCTGCTGAGCCCGCAGAGGCAGGCAAAGCAGAGACATCAGAAGGGTCAGG  
A H H S A E P A E A G K A E T S E G S G  
      |51                  |61

      |211      |221      |231      |241      |251      |261  
CTCCGCCCCGGCTGTGCCGAAGCTTCTGCCTCCCCAAACAGCGGCGTCCATCATCCG  
S A P A V P E A S A S P K Q R R S I I R  
      |71                  |81

      |271      |281      |291      |301      |311      |321  
TGACCGGGGACCCATGTATGATGACCCACCCCTGCCTGAAGGCTGGACACGGAAGCTTAA  
D R G P M Y D D P T L P E G W T R K L K  
      |91                  |101

      |331      |341      |351      |361      |371      .  
GCAAAGGAAATCTGGCCGCTCTGCTGGGAAGTATGATGTGTATTTGATCAAgtagtaag  
Q R K S G R S A G K Y D V Y L I N  
      |111                  |121

. . . . .  
agcaactcctatctctacagggcagggagggcagggacaaggatccctcatggagcagga  
. . . . .  
aaatgtatgtgccaggggtggggtcggggggaacataaacaatgaacactgagaccaggt  
. . . . .  
gtgcttgaaatgaccgtgtacagaggtcgctgccctgagtgggaagttctcaaggtagca  
. . . . .  
ggccctctatcctctccacacctaagtctttatctggggatggaatagctgcggaagca  
. . . . .  
gaggaacttgcagagctaggggttcagaggggtgaagaagcatgtttcagt

Exon 4 | Start: 110678 | End: 120315 | Length: 9637

. . . . .  
tgttctagatggtgactcaggcccaggcaccaaccagcagaatgggcctcagcctgacaa  
. . . . .  
cccttctgtaccaggcctgactctttggttgctgaactttggagaggcctgggggggtca  
. . . . .  
gcggcaggcagacgagtgagtggctttggtgacaggtcctcaggggcagccaggcagtggt  
. . . . .  
gactctcgttcaatagtaacgtttgtcagagcgttggtcaccaccatccgctctgcctat  
. . . . .  
ctctgacattgctatggagagcctctaattgttccttggtgtctttctgtttgtccccaca

|381 |391 |401 |411 |421 |431  
TCCCAGGGAAAAGCCTTTCGCTCTAAAGTGGAGTTGATTGCGTACTTCGAAAAGGTAGG  
P Q G K A F R S K V E L I A Y F E K V G  
|131 |141

|441 |451 |461 |471 |481 |491  
CGACACATCCCTGGACCCTAATGATTTTGA CTTCACGGTAACTGGGAGAGGGAGCCCCCTC  
D T S L D P N D F D F T V T G R G S P S  
|151 |161

|501 |511 |521 |531 |541 |551  
CCGGCGAGAGCAGAAAACCACCTAAGAAGCCCCAAATCTCCCAAAGCTCCAGGAACTGGCAG  
R R E Q K P P K K P K S P K A P G T G R  
|171 |181

|561 |571 |581 |591 |601 |611  
AGGCCGGGGACGCCCCAAAGGGAGCGGCACCACGAGACCCAAGGCGGCCACGTCAGAGGG  
G R G R P K G S G T T R P K A A T S E G  
|191 |201

|621 |631 |641 |651 |661 |671  
TGTGCAGGTGAAAAGGGTCCTGGAGAAAAGTCCTGGGAAGCTCCTTGTCAAGATGCCTTT  
V Q V K R V L E K S P G K L L V K M P F  
|211 |221

|681 |691 |701 |711 |721 |731  
TCAAACCTTCGCCAGGGGGCAAGGCTGAGGGGGGTGGGGCCACCACATCCACCCAGGTCAT  
Q T S P G G K A E G G G A T T S T Q V M  
|231 |241

741	751	761	771	781	791
GGTGATCAAACGCCCCGGCAGGAAGCGAAAAGCTGAGGCCGACCCTCAGGCCATTCCCAA					
V	I	K	R	P	G
			R	K	R
			K	A	E
			A	D	P
			Q	A	I
			P	K	
		251		261	
801	811	821	831	841	851
GAAACGGGGCCGAAAGCCGGGGAGTGTGGTGGCAGCCGCTGCCGCCGAGGCCAAAAAGAA					
K	R	G	R	K	P
			G	S	V
			V	A	A
			A	A	A
			A	E	A
			K	K	K
		271		281	
861	871	881	891	901	911
AGCCGTGAAGGAGTCTTCTATCCGATCTGTGCAGGAGACCGTACTCCCCATCAAGAAGCG					
A	V	K	E	S	S
			I	R	S
			V	Q	E
			T	V	L
			P	I	K
			K	K	R
		291		301	
921	931	941	951	961	971
CAAGACCCGGGAGACGGTCAGCATCGAGGTCAAGGAAGTGGTGAAGCCCTGCTGGTGTC					
K	T	R	E	T	V
			S	I	E
			V	K	E
			V	V	K
			P	L	L
			V	S	
		311		321	
981	991	1001	1011	1021	1031
CACCTCGGTGAGAAGAGCGGGAAAGGACTGAAGACCTGTAAGAGCCCTGGGCGGAAAAAG					
T	L	G	E	K	S
			G	K	G
			L	K	T
			C	K	S
			P	G	R
			K	S	
		331		341	
1041	1051	1061	1071	1081	1091
CAAGGAGAGCAGCCCCAAGGGGCGCAGCAGCAGCGCCTCCTACCCCCAAGAAGGAGCA					
K	E	S	S	P	K
			G	R	S
			S	S	S
			A	S	S
			P	P	K
			K	E	H
		351		361	
1101	1111	1121	1131	1141	1151
CCACCACCATCACCACCACTCAGAGTCCCCAAAGGCCCCCGTGCCACTGCTCCCACCCCT					
H	H	H	H	H	H
			S	E	S
			P	K	A
			P	V	P
			L	L	P
			P	P	L
		371		381	
1161	1171	1181	1191	1201	1211
GCCCCACCTCCACCTGAGCCCGAGAGCTCCGAGGACCCACCAGCCCCCTGAGCCCCA					
P	P	P	P	P	E
			P	E	S
			S	S	E
			D	P	T
			S	P	P
			E	P	Q
		391		401	
1221	1231	1241	1251	1261	1271
GGAATTGAGCAGCAGCGTCTGCAAAGAGGAGAAGATGCCAGAGGAGGCTCACTGGAGAG					
D	L	S	S	S	V
			C	K	E
			E	K	M
			P	R	G
			G	G	S
			L	E	S
		411		421	

1281	1291	1301	1311	1321	1331
CGACGGCTGCCCCAAGGAGCCAGCTAAGACTCAGCCCGCGGTTGCCACCGCCGCCACGGC					
D G C P K E P A K T Q P A V A T A A T A					
	431			441	

  

1341	1351	1361	1371	1381	1391
CGCAGAAAAGTACAAACACCGAGGGGAGGGAGAGCGCAAAGACATTGTTTCATCCTCCAT					
A E K Y K H R G E G E R K D I V S S S M					
	451			461	

  

1401	1411	1421	1431	1441	1451
GCCAAGGCCAAACAGAGAGGAGCCTGTGGACAGCCGGACGCCCGTGACCGAGAGAGTTAG					
P R P N R E E P V D S R T P V T E R V S					
	471			481	

  

1461	*11	*21	*31	*41	*51
CTGACTTTACACGGAGCGGATTGCAAAGCAAACCAACAAGAATAAAGGCAGCTGTTGTCT					
*					

  

*61	*71	*81	*91	*101	*111
CTTCTCCTTATGGGTAGGGCTCTGACAAAGCTTCCCGATTAACTGAAATAAAAAATATTT					

  

*121	*131	*141	*151	*161	*171
TTTTTCTTTTCAGTAACTTAGAGTTTCGTGGCTTCAGGGTGGGAGTAGTTGGAGCATTG					

  

*181	*191	*201	*211	*221	*231
GGGATGTTTTTCTTACCGACAAGCACAGTCAGGTTGAAGACCTAACCAGGGCCAGAAAGTA					

  

*241	*251	*261	*271	*281	*291
GCTTTGCACTTTTCTAAACTAGGCTCCTTCAACAAGGCTTGCTGCAGATACTACTGACCA					

  

*301	*311	*321	*331	*341	*351
GACAAGCTGTTGACCAGGCACCTCCCCTCCCGCCCAAACCTTTCCCCCATGTGGTCGTTA					

  

*361	*371	*381	*391	*401	*411
GAGACAGAGCGACAGAGCAGTTGAGAGGACACTCCCGTTTTTCGGTGCCATCAGTGCCCCG					

  

*421	*431	*441	*451	*461	*471
TCTACAGCTCCCCAGCTCCCCCAGCTCCCCCACTCCCAACCACGTTGGGACAGGGAGG					

  

*481	*491	*501	*511	*521	*531
TGTGAGGCAGGAGAGACAGTTGGATTCTTTAGAGAAGATGGATATGACCAGTGGCTATGG					

  

*541	*551	*561	*571	*581	*591
CCTGTGCGATCCCACCCGTGGTGGCTCAAGTCTGGCCCCACACCAGCCCCAATCCAAAAC					



*601	*611	*621	*631	*641	*651
TGGCAAGGACGCTTCACAGGACAGGAAAGTGGCACCTGTCTGCTCCAGCTCTGGCATGGC					
*661	*671	*681	*691	*701	*711
TAGGAGGGGGGAGTCCCTTGAACACTGGGTGTAGACTGGCCTGAACCACAGGAGAGGAT					
*721	*731	*741	*751	*761	*771
GGCCCAGGGTGAGGTGGCATGGTCCATTCTCAAGGGACGTCCTCCAACGGGTGGCGCTAG					
*781	*791	*801	*811	*821	*831
AGGCCATGGAGGCAGTAGGACAAGGTGCAGGCAGGCTGGCCTGGGGTCAGGCCGGGCAGA					
*841	*851	*861	*871	*881	*891
GCACAGCGGGGTGAGAGGGATTCCCTAATCACTCAGAGCAGTCTGTGACTTAGTGGACAGG					
*901	*911	*921	*931	*941	*951
GGAGGGGGCAAAGGGGAGGAGAAGAAAATGTTCTTCCAGTTACTTTCCAATTCTCCTTT					
*961	*971	*981	*991	*1001	*1011
AGGGACAGCTTAGAATTATTTGCACTATTGAGTCTTCATGTTCCCACTTCAAAACAAACA					
*1021	*1031	*1041	*1051	*1061	*1071
GATGCTCTGAGAGCAAACCTGGCTTGAATTGGTGACATTTAGTCCCTCAAGCCACCAGATG					
*1081	*1091	*1101	*1111	*1121	*1131
TGACAGTGTGAGAACTACCTGGATTTGTATATATACCTGCGCTTGTTTTAAAGTGGGCT					
*1141	*1151	*1161	*1171	*1181	*1191
CAGCACATAGGGTTCCACGAAGCTCCGAACTCTAAGTGTTGCTGCAATTTTATAAGG					
*1201	*1211	*1221	*1231	*1241	*1251
ACTTCCTGATTGTTTTCTTCTCCCTTCCATTTCTGCCTTTTGTTTCATTTTCATCCTTT					
*1261	*1271	*1281	*1291	*1301	*1311
CACTTCTTTCCCTTCCGTCCTCCTCCTCCTAGTTCATCCCTTCTCTTCCAGGCAGC					
*1321	*1331	*1341	*1351	*1361	*1371
CGCGGTGCCCAACCACACTTGTCGGCTCCAGTCCCCAGAACTCTGCCTGCCCTTTGTCCT					
*1381	*1391	*1401	*1411	*1421	*1431
CCTGCTGCCAGTACCAGCCCCACCCTGTTTTGAGCCCTGAGGAGGCCTTGGGCTCTGCTG					
*1441	*1451	*1461	*1471	*1481	*1491
AGTCCGACCTGGCCTGTCTGTGAAGAGCAAGAGAGCAGCAAGGTCTTGCTCTCCTAGGTA					
*1501	*1511	*1521	*1531	*1541	*1551

GCCCCCTCTTCCCTGGTAAGAAAAAGCAAAAGGCATTTCCCACCCTGAACAACGAGCCTT

|\*1561      |\*1571      |\*1581      |\*1591      |\*1601      |\*1611  
TTCACCCTTCTACTCTAGAGAAGTGGACTGGAGGAGCTGGGCCCATTGTTAGTTGAGG

|\*1621      |\*1631      |\*1641      |\*1651      |\*1661      |\*1671  
AAAGCACAGAGGCCTCCTGTGGCCTGCCAGTCATCGAGTGGCCCAACAGGGGCTCCATGC

|\*1681      |\*1691      |\*1701      |\*1711      |\*1721      |\*1731  
CAGCCGACCTTGACCTCACTCAGAAGTCCAGAGTCTAGCGTAGTGCAGCAGGGCAGTAGC

|\*1741      |\*1751      |\*1761      |\*1771      |\*1781      |\*1791  
GGTACCAATGCAGAACTCCCAAGACCCGAGCTGGGACCAGTACCTGGGTCCCCAGCCCTT

|\*1801      |\*1811      |\*1821      |\*1831      |\*1841      |\*1851  
CCTCTGCTCCCCCTTTCCCTCGGAGTTCTTCTTGAATGGCAATGTTTTGCTTTTGCTCG

|\*1861      |\*1871      |\*1881      |\*1891      |\*1901      |\*1911  
ATGCAGACAGGGGGCCAGAACACCACACATTTCACTGTCTGTCTGGTCCATAGCTGTGGT

|\*1921      |\*1931      |\*1941      |\*1951      |\*1961      |\*1971  
GTAGGGGCTTAGAGGCATGGGCTTGCTGTGGGTTTTTAATTGATCAGTTTTCATGTGGGA

|\*1981      |\*1991      |\*2001      |\*2011      |\*2021      |\*2031  
TCCCATCTTTTTAACCTCTGTTTCAGGAAGTCCTTATCTAGCTGCATATCTTCATCATATT

|\*2041      |\*2051      |\*2061      |\*2071      |\*2081      |\*2091  
GGTATATCCTTTTCTGTGTTTACAGAGATGTCTCTTATATCTAAATCTGTCCAACCTGAGA

|\*2101      |\*2111      |\*2121      |\*2131      |\*2141      |\*2151  
AGTACCTTATCAAAGTAGCAAATGAGACAGCAGTCTTATGCTTCCAGAAACACCCACAGG

|\*2161      |\*2171      |\*2181      |\*2191      |\*2201      |\*2211  
CATGTCCCATGTGAGCTGCTGCCATGAACTGTCAAGTGTGTGTTGTCTTGTGTATTTAG

|\*2221      |\*2231      |\*2241      |\*2251      |\*2261      |\*2271  
TTATTGTCCCTGGCTTCCTTACTATGGTGTAAATCATGAAGGAGTGAAACATCATAGAAAC

|\*2281      |\*2291      |\*2301      |\*2311      |\*2321      |\*2331  
TGTCTAGCACTTCCTTGCCAGTCTTTAGTGATCAGGAACCATAGTTGACAGTTCCAATCA

|\*2341      |\*2351      |\*2361      |\*2371      |\*2381      |\*2391  
GTAGCTTAAGAAAAAACCGTGTGTTGTCTCTTCTGGAATGGTTAGAAGTGAGGGAGTTTGC

|\*2401      |\*2411      |\*2421      |\*2431      |\*2441      |\*2451  
CCCGTTCTGTTTGTAGAGTCTCATAGTTGGACTTTCTAGCATATATGTGTCCATTTCTT

*2461	*2471	*2481	*2491	*2501	*2511
ATGCTGTAAAAGCAAGTCCTGCAACCAAACCTCCCATCAGCCCAATCCCTGATCCCTGATC					
*2521	*2531	*2541	*2551	*2561	*2571
CCTTCCACCTGCTCTGCTGATGACCCCCCAGCTTCACTTCTGACTCTTCCCCAGGAAGG					
*2581	*2591	*2601	*2611	*2621	*2631
GAAGGGGGGTCAGAAGAGAGGGTGAGTCCTCCAGAACTCTTCTCCAAGGACAGAAGGCT					
*2641	*2651	*2661	*2671	*2681	*2691
CCTGCCCCCATAGTGGCCTCGAACTCCTGGCACTACCAAAGGACACTTATCCACGAGAGC					
*2701	*2711	*2721	*2731	*2741	*2751
GCAGCATCCGACCAGGTTGTCACTGAGAAGATGTTTATTTTGGTCAGTTGGGTTTTTATG					
*2761	*2771	*2781	*2791	*2801	*2811
TATTATACTTAGTCAAATGTAATGTGGCTTCTGGAATCATTGTCCAGAGCTGCTTCCCCG					
*2821	*2831	*2841	*2851	*2861	*2871
TCACCTGGGCGTCATCTGGTCCTGGTAAGAGGAGTGCGTGGCCCACCAGGCCCCCCTGTC					
*2881	*2891	*2901	*2911	*2921	*2931
ACCCATGACAGTTCATTTCAGGGCCGATGGGGCAGTCGTGGTTGGGAACACAGCATTTCAA					
*2941	*2951	*2961	*2971	*2981	*2991
GCGTCACTTTATTTTCATTTCGGGCCCCACCTGCAGCTCCCTCAAAGAGGCAGTTGCCAGC					
*3001	*3011	*3021	*3031	*3041	*3051
CTCTTTCCCTTCCAGTTTATTCCAGAGCTGCCAGTGGGGCCTGAGGCTCCTTAGGGTTTT					
*3061	*3071	*3081	*3091	*3101	*3111
CTCTCTATTTCCCCCTTCTTCCCTCATTCCCTCGTCTTTCCCAAAGGCATCACGAGTCAG					
*3121	*3131	*3141	*3151	*3161	*3171
TCGCCTTTCAGCAGGCAGCCTTGGCGGTTTATCGCCCTGGCAGGCAGGGGCCCTGCAGCT					
*3181	*3191	*3201	*3211	*3221	*3231
CTCATGCTGCCCCCTGCCTTGGGGTCAGGTTGACAGGAGGTTGGAGGAAAGCCTTAAGCT					
*3241	*3251	*3261	*3271	*3281	*3291
GCAGGATTCTCACCAGCTGTGTCCGGCCAGTTTTGGGGTGTGACCTCAATTTCAATTTT					
*3301	*3311	*3321	*3331	*3341	*3351
GTCTGTACTTGAACATTATGAAGATGGGGGCCTTTTCAGTGAATTTGTGAACAGCAGAA					

*3361	*3371	*3381	*3391	*3401	*3411
TTGACCGACAGCTTTCCAGTACCCATGGGGCTAGGTCATTAAGGCCACATCCACAGTCTC					
*3421	*3431	*3441	*3451	*3461	*3471
CCCCACCCTTGTTCCAGTTGTTAGTTACTACCTCCTCTCCTGACAATACTGTATGTCGTC					
*3481	*3491	*3501	*3511	*3521	*3531
GAGCTCCCCCAGGTCTACCCCTCCCGGCCCTGCCTGCTGGTGGGCTTGTCATAGCCAGT					
*3541	*3551	*3561	*3571	*3581	*3591
GGGATTGCCGGTCTTGACAGCTCAGTGAGCTGGAGATACTTGGTCACAGCCAGGCGCTAG					
*3601	*3611	*3621	*3631	*3641	*3651
CACAGCTCCCTTCTGTTGATGCTGTATTCCCATATCAAAAGACACAGGGGACACCCAGAA					
*3661	*3671	*3681	*3691	*3701	*3711
ACGCCACATCCCCCAATCCATCAGTGCCAACTAGCCAACGGCCCCAGCTTCTCAGCTCG					
*3721	*3731	*3741	*3751	*3761	*3771
CTGGATGGCGGAAGCTGCTACTCGTGAGCGCCAGTGCGGGTGCAGACAATCTTCTGTTGG					
*3781	*3791	*3801	*3811	*3821	*3831
GTGGCATCATTCCAGGCCGAAGCATGAACAGTGCACCTGGGACAGGGAGCAGCCCCAAA					
*3841	*3851	*3861	*3871	*3881	*3891
TTGTACCTGCTTCTCTGCCCAGCTTTTCATTGCTGTGACAGTGATGGCGAAAGAGGGTA					
*3901	*3911	*3921	*3931	*3941	*3951
ATAACCAGACACAACTGCCAAGTTGGGTGGAGAAAGAGTTTCTTTAGCTGACAGAATC					
*3961	*3971	*3981	*3991	*4001	*4011
TCTGAATTTTAAATCACTTAGTAAGCGGCTCAAGCCCAGGAGGGAGCAGAGGGATACGAG					
*4021	*4031	*4041	*4051	*4061	*4071
CGGAGTCCCCTGCGCGGGACCATCTGGAATTGGTTTAGCCCAAGTGGAGCCTGACAGCCA					
*4081	*4091	*4101	*4111	*4121	*4131
GAACTCTGTGTCCCCCGTCTAACCACAGCTCCTTTTCCAGAGCATTCCAGTCAGGCTCTC					
*4141	*4151	*4161	*4171	*4181	*4191
TGGGCTGACTGGGCCAGGGGAGGTTACAGGTACCAGTTCTTTAAGAAGATCTTTGGGCAT					
*4201	*4211	*4221	*4231	*4241	*4251
ATACATTTTTCAGCTGTGTCATTGCCCCAAATGGATTCTGTTTCAAGTTCACACCTGCA					
*4261	*4271	*4281	*4291	*4301	*4311

GATTCTAGGACCTGTGTCCTAGACTTCAGGGAGTCAGCTGTTTCTAGAGTTCCTACCATG

|\*4321      |\*4331      |\*4341      |\*4351      |\*4361      |\*4371  
GAGTGGGTCTGGAGGACCTGCCCCGGTGGGGGGGCAGAGCCCTGCTCCCTCCGGGTCTTCC

|\*4381      |\*4391      |\*4401      |\*4411      |\*4421      |\*4431  
TACTCTTCTCTCTGCTCTGACGGGATTTGTTGATTCTCTCCATTTTGGTGTCTTTCTCTT

|\*4441      |\*4451      |\*4461      |\*4471      |\*4481      |\*4491  
TTAGATATTGTATCAATCTTTAGAAAAGGCATAGTCTACTTGTATAAATCGTTAGGATA

|\*4501      |\*4511      |\*4521      |\*4531      |\*4541      |\*4551  
CTGCCTCCCCCAGGGTCTAAAATTACATATTAGAGGGGAAAAGCTGAACACTGAAGTCAG

|\*4561      |\*4571      |\*4581      |\*4591      |\*4601      |\*4611  
TTCTCAACAATTTAGAAGGAAAACCTAGAAAACATTTGGCAGAAAATTACATTTTCGATGT

|\*4621      |\*4631      |\*4641      |\*4651      |\*4661      |\*4671  
TTTTGAATGAATACGAGCAAGCTTTTACAACAGTGCTGATCTAAAAATACTTAGCACTTG

|\*4681      |\*4691      |\*4701      |\*4711      |\*4721      |\*4731  
GCCTGAGATGCCTGGTGAGCATTACAGGCAAGGGGAATCTGGAGGTAGCCGACCTGAGGA

|\*4741      |\*4751      |\*4761      |\*4771      |\*4781      |\*4791  
CATGGCTTCTGAACCTGTCTTTTGGGAGTGGTATGGAAGGTGGAGCGTTCACCAGTGACC

|\*4801      |\*4811      |\*4821      |\*4831      |\*4841      |\*4851  
TGGAAGGCCCAGCACCACCCTCCTTCCCACTCTTCTCATCTTGACAGAGCCTGCCCCAGC

|\*4861      |\*4871      |\*4881      |\*4891      |\*4901      |\*4911  
GCTGACGTGTCAGGAAAACACCCAGGGAAGTGAAGGCACCTTCTGCCTGAGGGGCAGCC

|\*4921      |\*4931      |\*4941      |\*4951      |\*4961      |\*4971  
TGCCTTGCCCACTCCTGCTCTGCTCGCCTCGGATCAGCTGAGCCTTCTGAGCTGGCCTCT

|\*4981      |\*4991      |\*5001      |\*5011      |\*5021      |\*5031  
CACTGCCTCCCCAAGGCCCTGCTGCCCCTGTCAGGAGGCAGAAGGAAGCAGGTGTGAG

|\*5041      |\*5051      |\*5061      |\*5071      |\*5081      |\*5091  
GGCAGTGAAGGAGGGAGCACAACCCCAAGCTCCCGCTCCGGGCTCCGACTTGTGCACAG

|\*5101      |\*5111      |\*5121      |\*5131      |\*5141      |\*5151  
GCAGAGCCCAGACCCTGGAGGAAATCCTACCTTTGAATTCAAGAACATTTGGGGAATTTG

|\*5161      |\*5171      |\*5181      |\*5191      |\*5201      |\*5211  
GAAATCTCTTTGCCCCAAACCCCAATTCTGTCTACCTTTAATCAGGTCCTGCTCAGCA

*5221	*5231	*5241	*5251	*5261	*5271
GTGAGAGCAGATGAGGTGAAAAGGCCAAGAGGTTTGGCTCCTGCCCCTGATAGCCCTC					
*5281	*5291	*5301	*5311	*5321	*5331
TCCCCGAGTGTTTGTGTGTCAAGTGGCAAAGCTGTTCTTCCTGGTGACCCTGATTATAT					
*5341	*5351	*5361	*5371	*5381	*5391
CCAGTAACACATAGACTGTGCGCATAGGCCTGCTTTGTCTCCTCTATCCTGGGCTTTTGT					
*5401	*5411	*5421	*5431	*5441	*5451
TTTGCTTTTGTAGTTTGTGCTTTTGTGCTTTTCTGTCCCTTTTATTTAACGCACCGACTAGAC					
*5461	*5471	*5481	*5491	*5501	*5511
ACACAAAGCAGTTGAATTTTATATATATATCTGTATATTGCACAATTATAAACTCATTT					
*5521	*5531	*5541	*5551	*5561	*5571
TGCTTGTGGCTCCACACACACAAAAAAGACCTGTTAAAATTATACCTGTGCTTAATTA					
*5581	*5591	*5601	*5611	*5621	*5631
CAATATTTCTGATAACCATAGCATAGGACAAGGAAAATAAAAAAAGAAAAAAGAAAA					
*5641	*5651	*5661	*5671	*5681	*5691
AAAAACGACAAATCTGTCTGCTGGTCACTTCTCTGTCCAAGCAGATTCTGCGTCTTTTC					
*5701	*5711	*5721	*5731	*5741	*5751
CTCGCTTCTTTCAAGGGCTTTCTGTGCCAGGTGAAGGAGGCTCCAGGCAGCACCCAGGT					
*5761	*5771	*5781	*5791	*5801	*5811
TTTGCACTCTTGTTTCTCCCGTGCTTGTGAAAGAGGTCCCAAGGTTCTGGGTGCAGGAGC					
*5821	*5831	*5841	*5851	*5861	*5871
GCTCCCTTGACCTGCTGAAGTCCGGAACGTAGTCGGCACAGCCTGGTCGCCTTCCACCTC					
*5881	*5891	*5901	*5911	*5921	*5931
TGGGAGCTGGAGTCCACTGGGGTGGCCTGACTCCCCAGTCCCCTTCCCGTGACCTGGTC					
*5941	*5951	*5961	*5971	*5981	*5991
AGGGTGAGCCCATGTGGAGTCAGCCTCGCAGGCCTCCCTGCCAGTAGGGTCCGAGTGTGT					
*6001	*6011	*6021	*6031	*6041	*6051
TTCATCCTTCCCACTCTGTGAGCCTGGGGGCTGGAGCGGAGACGGGAGGCCTGGCCTGT					
*6061	*6071	*6081	*6091	*6101	*6111
CTCGGAACCTGTGAGCTGCACCAGGTAGAACGCCAGGGACCCAGAATCATGTGCGTCAG					

*6121	*6131	*6141	*6151	*6161	*6171
TCCAAGGGGTCCCCTCCAGGAGTAGTGAAGACTCCAGAAATGTCCCTTTCTTCTCCCCCA					
*6181	*6191	*6201	*6211	*6221	*6231
TCCTACGAGTAATTGCATTTGCTTTTGTAATTCTTAATGAGCAATATCTGCTAGAGAGTT					
*6241	*6251	*6261	*6271	*6281	*6291
TAGCTGTAAACAGTTCTTTTGATCATCTTTTTTTAATAATTAGAAACACCAAAAAAATCC					
*6301	*6311	*6321	*6331	*6341	*6351
AGAAACTTGTCTTCCAAAGCAGAGAGCATTATAATCACCAGGGCCAAAAGCTTCCCTCC					
*6361	*6371	*6381	*6391	*6401	*6411
CTGCTGTCAATTGCTTCTTCTGAGGCCTGAATCCAAAAGAAAAACAGCCATAGGCCCTTC					
*6421	*6431	*6441	*6451	*6461	*6471
AGTGGCCGGGCTACCCGTGAGCCCTTCGGAGGACCAGGGCTGGGGCAGCCTCTGGGCCCA					
*6481	*6491	*6501	*6511	*6521	*6531
CATCCGGGGCCAGCTCCGGCGTGTGTTTCAGTGTTAGCAGTGGGTCATGATGCTCTTTCCC					
*6541	*6551	*6561	*6571	*6581	*6591
ACCCAGCCTGGGATAGGGGCAGAGGAGGCGAGGAGGCCGTTGCCGCTGATGTTTGCCGT					
*6601	*6611	*6621	*6631	*6641	*6651
GAACAGTGGGTGTCTGCGTGCGTCCACGTGCGTGTTTTCTGACTGACATGAAATCGACG					
*6661	*6671	*6681	*6691	*6701	*6711
CCCGAGTTAGCCTCACCCGGTGACCTCTAGCCCTGCCCGGATGGAGCGGGGCCACCCGG					
*6721	*6731	*6741	*6751	*6761	*6771
TTCAGTGTTTCTGGGGAGCTGGACAGTGGAGTGCAAAAGGCTTGCAGAACTGAAGCCTG					
*6781	*6791	*6801	*6811	*6821	*6831
CTCCTTCCCTTGCTACCACGGCCTCCTTTCCGTTTGATTTGTCAGTCTTCAATCAATAA					
*6841	*6851	*6861	*6871	*6881	*6891
CAGCCGCTCCAGAGTCAGTAGTCAATGAATATATGACCAAAATATCACCAGGACTGTTACT					
*6901	*6911	*6921	*6931	*6941	*6951
CAATGTGTGCCGAGCCCTTGCCCATGCTGGGCTCCCGTGATCTGGACACTGTAACGTGT					
*6961	*6971	*6981	*6991	*7001	*7011
GCTGTGTTTGCTCCCTTCCCTTCCTTCTTTGCCCTTTACTTGTCTTTCTGGGGTTTTT					
*7021	*7031	*7041	*7051	*7061	*7071

CTGTTTGGGTTTGGTTTGGTTTTTATTTCTCCTTTTGTGTTCCAAACATGAGGTTCTCTC

|\*7081      |\*7091      |\*7101      |\*7111      |\*7121      |\*7131  
TACTGGTCCTCTTAACTGTGGTGTGAGGCTTATATTTGTGTAATTTTGGTGGGTGAAA

|\*7141      |\*7151      |\*7161      |\*7171      |\*7181      |\*7191  
GGAATTTTGCTAAGTAAATCTCTTCTGTGTTTGAAGTGAAGTCTGTATTGTAAGTATGTT

|\*7201      |\*7211      |\*7221      |\*7231      |\*7241      |\*7251  
TAAAGTAATTGTTCCAGAGACAAATATTTCTAGACACTTTTCTTTACAAACAAAAGCAT

|\*7261      |\*7271      |\*7281      |\*7291      |\*7301      |\*7311  
TCGGAGGGAGGGGGATGGTGAAGTGAAGAGGGGAGAGCTGAACAGATGACCCCTGCC

|\*7321      |\*7331      |\*7341      |\*7351      |\*7361      |\*7371  
CAGATCAGCCAGAAGCCACCCAAAGCAGTGGAGCCCAGGAGTCCCACTCCAAGCCAGCAA

|\*7381      |\*7391      |\*7401      |\*7411      |\*7421      |\*7431  
GCCGAATAGCTGATGTGTTGCCACTTTCCAAGTCACTGCAAAACCAGGTTTGTTCGCC

|\*7441      |\*7451      |\*7461      |\*7471      |\*7481      |\*7491  
CAGTGGATTCTTGTGTTTGCTTCCCCTCCCCCGAGATTATTACCACCATCCCGTGCTTTT

|\*7501      |\*7511      |\*7521      |\*7531      |\*7541      |\*7551  
AAGGAAAGGCAAGATTGATGTTTCCTTGAGGGGAGCCAGGAGGGGATGTGTGTGTGCAGA

|\*7561      |\*7571      |\*7581      |\*7591      |\*7601      |\*7611  
GCTGAAGAGCTGGGGAGAATGGGGCTGGGCCCACCCAAGCAGGAGGCTGGGACGCTCTGC

|\*7621      |\*7631      |\*7641      |\*7651      |\*7661      |\*7671  
TGTGGGCACAGGTCAGGCTAATGTTGGCAGATGCAGCTCTTCTGACAGGCCAGGTGGT

|\*7681      |\*7691      |\*7701      |\*7711      |\*7721      |\*7731  
GGGCATTCTCTCTCCAAGGTGTGCCCCGTGGGCATTACTGTTTAAGACACTTCCGTCACA

|\*7741      |\*7751      |\*7761      |\*7771      |\*7781      |\*7791  
TCCCACCCCATCCTCCAGGGCTCAACACTGTGACATCTCTATTCCCACCCCTCCCCTTCC

|\*7801      |\*7811      |\*7821      |\*7831      |\*7841      |\*7851  
CAGGGCAATAAAATGACCATGGAGGGGGCTTGCACTCTCTTGGCTGTCAACCGATCGCCA

|\*7861      |\*7871      |\*7881      |\*7891      |\*7901      |\*7911  
GCAAAACTTAGATGTGAGAAAACCCCTTCCCATTCCATGGCGAAAACATCTCCTTAGAAA

|\*7921      |\*7931      |\*7941      |\*7951      |\*7961      |\*7971  
AGCCATTACCCCTATTAGGCATGGTTTTGGGCTCCCAAAACACCTGACAGCCCTCCCTC



*7981	*7991	*8001	*8011	*8021	*8031
CTCTGAGAGGCGGAGAGTGCTGACTGTAGTGACCATTCATGCCGGGTGCAGCATCTGGA					
*8041	*8051	*8061	*8071	*8081	*8091
AGAGCTAGGCAGGGTGTCTGCCCCCTCCTGAGTTGAAGTCATGCTCCCCTGTGCCAGCCC					
*8101	*8111	*8121	*8131	*8141	*8151
AGAGGCCGAGAGCTATGGACAGCATTGCCAGTAACACAGGCCACCCTGTGCAGAAGGGAG					
*8161	*8171	*8181	*8191	*8201	*8211
CTGGCTCCAGCCTGGAAACCTGTCTGAGGTTGGGAGAGGTGCACTTGGGGCACAGGGAGA					
*8221	*8231	*8241	*8251	*8261	*8271
GGCCGGGACACACTTAGCTGGAGATGTCTCTAAAAGCCCTGTATCGTATTCACCTTCAGT					
*8281	*8291	*8301	*8311	*8321	*8331
TTTTGTGTTTTGGGACAATTACTTTAGAAAATAAGTAGGTCGTTTTAAAAACAAAATTA					
*8341	*8351	*8361	*8371	*8381	*8391
TTGATTGCTTTTTTGTAGTGTTCAAAAAAGGTTCTTTGTGTATAGCCAAATGACTGAA					
*8401	*8411	*8421	*8431	*8441	*8451
AGCACTGATATATTTAAAAACAAAAGGCAATTTATTAAGGAAATTTGTACCATTTTCAGTA					
*8461	*8471	*8481	*8491	*8501	*8511
AACCTGTCTGAATGTACCTGTATACGTTTCAAAAACACCCCCCCCCCACTGAATCCCTGT					
*8521	*8531	*8541	*8551	.	.
AACCTATTTATTATATAAAGAGTTTGCCTTATAAATTTacataaaaatgtccgtttgtgt					
.	.	.	.	.	.
cttttggtgtaaaaatcaagtgattttttcataagggtcttttactattggaaaagatgg					
.	.	.	.	.	.
gcagcacgcagttttattttatttttgtaagttttttaatacatgtgaaagcaaagaata					
.	.	.	.	.	.
ctcagcatgcctttctaagtgacgcgtttgcaccttttggtgggaagtactgtatcctgt					
.	.	.	.	.	.
gctgttagcattctcgataaatctctctgtgaaagtgactcaaggtctgggctttcatta					
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Control: Version: 1, Version Date: 11/02/2015