

Gene: RUNX1 - Sequence: NG\_011402.2  
Transcript: NM\_001754.4 - Protein: NP\_001745.2  
Date : February 23, 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 MIDDLE base of codon  
4<sup>th</sup> line: Amino acid numbering. Numbered on 1<sup>st</sup> and increments of 10

Exon 1 | Start: 940414 | End: 940544 | Length: 130  
BE AWARE: This intron is shared with the following exon

```
. . . . .
atcagtagttccaaaaaccacaaataacaacaggagccgagttgtactaaatcagcaaag

. . . . .
accattgagatataataaagtgactgagtcacttttttctacatccccctcttgcaaag

. . . . .
tctactcggctattttcttgacagcctgggggagggcaggtggagggaaggaagggcat

. . . . .
tgctcagaaaagaaaaaaaaaattgacatcacttaagtcacatgattggcaagaaccaa

. . . . .
ttgagatgggctgtggaagggaacagttaaatttgtaatttgggttggtgtaaaactt

|-189    |-179    |-169    |-159    |-149    |-139
CTTTGGGCTCATAACAACACAGAACCACAAGTTGGGTAGCCTGGCAGTGTGAGAAGT

|-129    |-119    |-109    |-99     |-89     |-79
CTGAACCCAGCATAGTGGTCAGCAGGCAGGACGAATCACACTGAATGCAAACACAGGGT

|-69
TTCGACGCGTGgtgagcatcaccaacccacagccaaggcggcgctggctttttttttt

. . . . .
tttaatctttaacaatttgaatatttgtttttacaaaggtgcattttttaatagg
```

Exon 2 | Start: 940754 | End: 940870 | Length: 116  
BE AWARE: This intron is shared with the previous exon

. . . . .  
 gcttggggaggtcccagaggtatccagcagaggggagaagaaagagagatgtagggctaga  
 . . . . . | -59 | -49  
 ggggtgaggctgaaacagtgacctgtcttggttttcgctccgaagGTAAGAAATCATT  
 | -39 | -29 | -19 | -9 | 1 | 11  
 GAGTCCCCCGCCTTCAGAAGAGGGTGCATTTTCAGGAGGAAGCGATGGCTTCAGACAGCA  
 M A S D S I  
 | 1  
 | 21 | 31 | 41 | 51 . . .  
 TATTGAGTCATTTTCCTTCGTACCCACAGTGCCTTCATGAGAGgtgagtacatgctggtct  
 F E S F P S Y P Q C F M R E  
 | 11  
 . . . . .  
 tgtaatatctacttttgcctcagctttgcctgtaatgaaatggcagcttgtttcacctcgg  
 . . . . .  
 tgcagagatgcctcggcctgccagttccctgtcttgtttgtgagaggaattcaaactg  
 . . . . .  
 aggcataatgattacaagtctattggattacttactaatcagatggaagctcttcagaaat  
 . . . . .  
 gttttaataaatacttagttatgctgttgagtggtcagtcggtgcgtgagaactttgtc  
 . . . . .  
 aagtgcgagtaagttgtgctggtcttgtttaaaggcttagag

Exon 3 | Start: 1096749 | End: 1096787 | Length: 38

. . . . .  
 cattaaataaatcacattttaacacacacggagcttaaaagaggtgattctatgacaga  
 . . . . .  
 tattcagtagtagctagtgaaggagaacattgggtgaaggtacctaaccacgtgcata  
 . . . . .  
 aggaacagtttatttcttatttggtgatttactatttcaagtcatttttgtgtcaaagct  
 . . . . .  
 ataaaaatctccagtgctaaaaagtgtaagaagaatgagctagcaaaaaaaaaaagtc

. . . . .  
 gggggaaagttaatttttaaacttgcattacttttattttcctacttttctgttatttcag  
  
 |61 |71 |81 |91 . . . . .  
 AATGCATACTTGGAATGAATCCTTCTAGAGACGTCCACGgtatgttaaattccaagttat  
 C I L G M N P S R D V H D  
 |21 |31  
  
 . . . . .  
 atttttaaataaagtacacaccttcatagatgtgtatatgattttatatctatgtatctc  
  
 . . . . .  
 tatgtataacttgtgatataatttaaccaaataattaaggctgtttctgctgattccaat  
  
 . . . . .  
 atacttaaaagcagttctccatattgtaaatgtataccatataaaatatttcaatgaaa  
  
 . . . . .  
 gcttaaatagataccctcagttctacagcgtttaccataggtgcattctgatgctattaaa  
  
 . . . . .  
 tgcctctatacaactctctaaaaacaaaacagagcttgg

Exon 4 | Start: 1102616 | End: 1102869 | Length: 253

. . . . .  
 tggaagatagtcctcactttttaatgcactaagcaatcgggtgctaggagcccatcctgg  
  
 . . . . .  
 gtcagaggccgatccgcagaaccagaacgttttcccctcctggactgtagtaacttagt  
  
 . . . . .  
 ctcccctctcccctaaccacccccgcccccccccccccccgagtaataaaggcccctg  
  
 . . . . .  
 aacgtgtatgttggctctccgggagctgcttgctgaagatccgcgcccctgtcgccgtct  
  
 . . . . .  
 ggtaggagctgtttgcagggtcctaactcaatcggccttggtgtgatgcgtatccccgtag  
  
 |101 |111 |121 |131 |141 |151  
 ATGCCAGCACGAGCCGCCGCTTCACGCCGCCTTCCACCGCGCTGAGCCCAGGCAAGATGA  
 A S T S R R F T P P S T A L S P G K M S



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. . . . .
aagtgtcttctctttataacttttctgttgtttgcttaagttatcattgctattcctct

. . . . .
gcaacctaaaaagaaatcattgaatatacatttaattttagaataatcactacacaaatg

. . . . .
ccctaaaagtgtatgtataacatccctgatgtctgcatttgctccttgactgggtgtag

      |361      |371      |381      |391      |401      |411
GTGGTGGCCCTAGGGGATGTTCCAGATGGCACTCTGGTCACTGTGATGGCTGGCAATGAT
V V A L G D V P D G T L V T V M A G N D
      |121                      |131

      |421      |431      |441      |451      |461      |471
GAAAACTACTCGGCTGAGCTGAGAAATGCTACCGCAGCCATGAAGAACCAGGTTGCAAGA
E N Y S A E L R N A T A A M K N Q V A R
      |141                      |151

      |481      |491      |501      . . . . .
TTTAATGACCTCAGGTTTGTCGGTCGAAGTGAAGAGgtacgttatctgtcaaaactatg
F N D L R F V G R S G R G
      |161

. . . . .
cttgaaacacgtttcatggcaacaacccacatttcaaaattctagtatttctgtgatg

. . . . .
gaatccctagaaactcggctctgtcttaacattcttattgctatgtcaagctcttaaaat

. . . . .
atttatttgaagttaatatatttttagcagaatctctatattattgctttatgttactgag

. . . . .
tggatttaagactgatttggaacatgcaacgtaggaattgtgaatttcaaagaaacgcct

. . . . .
gagatttttactaaaatgcaaaaaatataaaactgac

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Exon 6 | Start: 1130134 | End: 1130238 | Length: 104

```

. . . . .
atctgactctatggtttccggaaatcctgacttaacagaaaataaattctaattcttctt

```

. . . . .  
 gatcttgattgtttattgatgattacattaaagccatataactatgtgaagtgttacac  
 . . . . .  
 ttaggcaactatcataggtgtgtattccagaattgaagcaaagccaaaattccgggagtg  
 . . . . .  
 ttgtcaaaactggtaacttgtgctgaagggctggacagcataaattaatgattggttatt  
 . . . . .  
 caacagatatgttcaggccaccaacctcattctgttttgttctctatcgtgtccccacag  
 |511 |521 |531 |541 |551 |561  
 GGAAAAGCTTCACTCTGACCATCACTGTCTTCACAAACCCACGCAAGTCGCCACCTACC  
 K S F T L T I T V F T N P P Q V A T Y H  
 |171 |181  
 |571 |581 |591 |601 |611 . . .  
 ACAGAGCCATCAAAATCACAGTGGATGGGCCCCGAGAACCTCGAAgtaagtgcattccact  
 R A I K I T V D G P R E P R R  
 |191 |201  
 . . . . .  
 tggggctggtacaccctccaggctggtacaccctccaggctggtataactcagggaccatg  
 . . . . .  
 tctcagattccttgggttcaacctttccccagggggatgtcaggccccgtaaagccaa  
 . . . . .  
 aaagttgcattgtcaatactctcgctactcaaagcttgtttctgtgcactgagctctttt  
 . . . . .  
 tgaagcatccttttagagggcaaaaggaaattaggctgactgaatcatatgttagaacca  
 . . . . .  
 ttcagggcagcatgaaactaaggactgcttcctcaatatacagta

Exon 7 | Start: 1155111 | End: 1155302 | Length: 191

. . . . .  
 ttctgtatctcagtgagttgtgcccattggaagtgggtgtagagggttggggagggtgcc  
 . . . . .



Exon 8 | Start: 1190250 | End: 1190411 | Length: 161

```
. . . . .
gacattaatacttctaccttgaaagtgtaatccattcctctcatcattgttagcatgatt

. . . . .
aaataaaacctttggaaggaatagtatatcaggtgaaaatctccaagaatcagtctctttt

. . . . .
gggggaaaataatccaacagaggcagatacttggacttgagtaggcttattaaaccctgg

. . . . .
tacataggccacatacatgtatgtgacatatttgaacaagggccactcatttcttattaa

. . . . .
aagacattttttaaatcccaccccactttacatataattgacctttctgattctcttcag

      |811      |821      |831      |841      |851      |861
ATACAAGGCAGATCCAACCATCCCCACCGTGGTCCTACGATCAGTCCTACCAATACCTGG
  T  R  Q  I  Q  P  S  P  P  W  S  Y  D  Q  S  Y  Q  Y  L  G
      |271                      |281

      |871      |881      |891      |901      |911      |921
GATCCATTGCCTCTCCTTCTGTGCACCCAGCAACGCCCATTTACCTGGACGTGCCAGCG
  S  I  A  S  P  S  V  H  P  A  T  P  I  S  P  G  R  A  S  G
      |291                      |301

      |931      |941      |951      |961      .      .      .
GCATGACAACCCTCTCTGCAGAACTTTCCAGTCGACTCTCAAgtaagccacttgaaaaca
  M  T  T  L  S  A  E  L  S  S  R  L  S  T
      |311                      |321

. . . . .
cattctttgcagctgagctggggtggaaggtccaggagactagaggtgcatgaaggagtt

. . . . .
ggcagaacattattgagtaaaacatggtttaaagggtgacctatctctactctgagtccca

. . . . .
tgcccatgcactgattttcttggatacataagaggcagagattgtggatgtgaaggcaga

. . . . .
agtggcttggaagaagtaggttccttagacatctaaagttgttcacatttggcaagaaagg
```



. . . . .  
atggtgactgaagggttaagaacaatatataaaaagccaaagt

Exon 9 | Start: 1197102 | End: 1201911 | Length: 4809

. . . . .  
gcatagcatcatgggtagtagtcccatcctcttgggaggtgacatgctgggtgatcctcgtc  
. . . . .  
atctcaggaggtggcatcctgggtgggtccctgtccccctgggtatagcatcctgggtaat  
. . . . .  
cctcgtcctcttgggagtagcatcccggtgggtccccgtcctccccagcagtagcatcct  
. . . . .  
gggtggcttcccatcctcctaggcggtatcatcctgggtagccccctggggcagagggaa  
. . . . .  
gagctgtggcctccgcaacctcctactcacttccgctccgttctcttggccgcctgcag

971	981	991	1001	1011	1021
CGGCACCCGACCTGACAGCGTT	CAGCGACCCGCGCCAGTT	CCCCGCGCTGCCCTCCATCT			
A P D L T A F S D P R Q F P A L P S I S					
	331				341

1031	1041	1051	1061	1071	1081
CCGACCCCGCATGCACTATCCAGGCGCCTT	CACCTACTCCCGACGCGGT	CACCTCGG			
D P R M H Y P G A F T Y S P T P V T S G					
	351				361

1091	1101	1111	1121	1131	1141
GCATCGGCATCGGCATGTCGGCCATGGGCTCGGCCACGCGTACCACACCTACCTGCCGC					
I G I G M S A M G S A T R Y H T Y L P P					
	371				381

1151	1161	1171	1181	1191	1201
CGCCCTACCCCGGCTCGTCGCAAGCGCAGGGAGGCCGTTCCAAGCCAGCTCGCCCTCCT					
P Y P G S S Q A Q G G P F Q A S S P S Y					
	391				401

1211	1221	1231	1241	1251	1261
ACCACCTGTACTACGGCGCCTCGGCCGGCTCCTACCAGTTCTCCATGGTGGGCGGCGAGC					
H L Y Y G A S A G S Y Q F S M V G G E R					
	411				421



*611	*621	*631	*641	*651	*661
ATGGAAGAAATATGATTCCCTTTTCTTTAAGTTTATTTAACTTTTCTTTTGGACTTTT					
*671	*681	*691	*701	*711	*721
GGGTAGTTGTTTTTTTGTGTTTGTGTTTTTTGAGAAACAGCTACAGCTTTGGGTC					
*731	*741	*751	*761	*771	*781
ATTTTAACTACTGTATTCCCAAGGAATCCCAGATATTTATGTATCTTGATGTTTCA					
*791	*801	*811	*821	*831	*841
ACATTTATGTGTTGATAATTTTAAATTATTTAAATGTACTTATATTAAGAAAAATATCA					
*851	*861	*871	*881	*891	*901
AGTACTACATTTTCTTTGTGTTCTGATAGTAGCCAAAGTTAAATGTATCACATTGAAGAA					
*911	*921	*931	*941	*951	*961
GGCTAGAAAAAAGAATGAGTAATGTGATCGCTTGGTTATCCAGAAGTATTGTTTACATT					
*971	*981	*991	*1001	*1011	*1021
AAACTCCCTTTTCATGTTAATCAAACAAGTGAGTAGCTCACGCAGCAACGTTTTTAATAGG					
*1031	*1041	*1051	*1061	*1071	*1081
ATTTTGTAGACTGAGGGTCACTCCAAGGATCAGAAGTATGGAATTTTCTGCCAGGCTCA					
*1091	*1101	*1111	*1121	*1131	*1141
ACAAGGGTCTCATATCTAACTTCCTCCTTAAACAGAGAAGGTCAATCTAGTTCCAGAGG					
*1151	*1161	*1171	*1181	*1191	*1201
GTTGAGGCAGGTGCCAATAATTACATCTTTGGAGAGGATTTGATTTCTGCCAGGGATTT					
*1211	*1221	*1231	*1241	*1251	*1261
GCTCACCCCAAGGTCATCTGATAATTTACAGATGCTGTGTAACAGAACACAGCCAAAGT					
*1271	*1281	*1291	*1301	*1311	*1321
AAACTGTGTAGGGGAGCCACATTTACATAGGAACCAATCAATGAATTTAGGGGTTACGA					
*1331	*1341	*1351	*1361	*1371	*1381
TTATAGCAATTTAAGGGCCCACCAGAAGCAGGCCTCGAGGAGTCAATTTGCCTCTGTGTG					
*1391	*1401	*1411	*1421	*1431	*1441
CCTCAGTGGAGACAAGTGGGAAAACATGGTCCCACCTGTGCGAGACCCCTGTCCTGTGC					
*1451	*1461	*1471	*1481	*1491	*1501
TGCTCACTCAACAACATCTTTGTGTTGCTTTCACCAGGCTGAGACCCTACCCTATGGGGT					
*1511	*1521	*1531	*1541	*1551	*1561

ATATGGGCTTTTACCTGTGCACCACTGTGACAGGAAAGATTTCATGTCTACTACTGTCCGTG  
 |\*1571 |\*1581 |\*1591 |\*1601 |\*1611 |\*1621  
 GCTACAATTCAAAGGTATCCAATGTCGCTGTAAATTTTATGGCACTATTTTATTGGAGG  
 |\*1631 |\*1641 |\*1651 |\*1661 |\*1671 |\*1681  
 ATTTGGTCAGAATGCAGTTGTTGTACAACTCATAAATACTAACTGCTGATTTTGACACAT  
 |\*1691 |\*1701 |\*1711 |\*1721 |\*1731 |\*1741  
 GTGTGCTCCAAATGATCTGGTGGTTATTTAACGTACCTCTTAAATTCGTTGAAACGATT  
 |\*1751 |\*1761 |\*1771 |\*1781 |\*1791 |\*1801  
 TCAGGTCAACTCTGAAGAGTATTTGAAAGCAGGACTTCAGAACAGTGTGATTTTATT  
 |\*1811 |\*1821 |\*1831 |\*1841 |\*1851 |\*1861  
 TTATAAATTTAAGCATTCAAATTAGGCAAATCTTTGGCTGCAGGCAGCAAAAACAGTGG  
 |\*1871 |\*1881 |\*1891 |\*1901 |\*1911 |\*1921  
 ACTTATTTAAAACAACTTGTTTTTGAGTTTTCTTATATATATATTGATTATTTGTTTTAC  
 |\*1931 |\*1941 |\*1951 |\*1961 |\*1971 |\*1981  
 ACACATGCAGTAGCACTTTGGTAAGAGTTAAAGAGTAAAGCAGCTTATGTTGTCAGTCTG  
 |\*1991 |\*2001 |\*2011 |\*2021 |\*2031 |\*2041  
 TTCTTATCTAGAGAAGAGCTATAGCAGATCTCGGACAACTCAGAATATATTCACTTTCA  
 |\*2051 |\*2061 |\*2071 |\*2081 |\*2091 |\*2101  
 TTTTGGACAGGATTCCCTCCACAACTCAGTTTCATATATTATTCCGTATTACATTTTGC  
 |\*2111 |\*2121 |\*2131 |\*2141 |\*2151 |\*2161  
 AGCTAAATTACCATAAAATGTCAGCAAATGTAAAAATTTAATTTCTGAAAAGCACCATTA  
 |\*2171 |\*2181 |\*2191 |\*2201 |\*2211 |\*2221  
 GCCCATTTCGCCCAAATTAACGTAATGTTTTTTTTCAGCACATGTTACCATGTCTGAC  
 |\*2231 |\*2241 |\*2251 |\*2261 |\*2271 |\*2281  
 CTGCAAAAATGCTGGAGAAAAATGAAGGAAAAAATTATGTTTTTCAGTTTAATTCTGTTA  
 |\*2291 |\*2301 |\*2311 |\*2321 |\*2331 |\*2341  
 ACTGAAGATATTCCAACCTCAAACCCAGCCTCATGCTCTGATTAGATAATCTTTTACATTG  
 |\*2351 |\*2361 |\*2371 |\*2381 |\*2391 |\*2401  
 AACCTTTACTCTCAAAGCCATGTGTGGAGGGGCTTGTCCTATTGTAGGCTCACTGGAT  
 |\*2411 |\*2421 |\*2431 |\*2441 |\*2451 |\*2461  
 TGGTCATTTAGAGTTTCACAGACTCTTACCAGCATATATAGTATTTAATTGTTTTCAAAA

*2471	*2481	*2491	*2501	*2511	*2521
AAATCAAAGTGTAGTTGTTTTGGCGATAGGTCTCACGCAACACATTTTTGTATGTGTGTG					
*2531	*2541	*2551	*2561	*2571	*2581
TGTGTGCGTGTGTGTGTGTGTGTGAAAAATTGCATTCATTGACTTCAGGTAGATTAAG					
*2591	*2601	*2611	*2621	*2631	*2641
GTATCTTTTTATTTCATTGCCCTCAGGAAAGTTAAGGTATCAATGAGACCCTTAAGCCAAT					
*2651	*2661	*2671	*2681	*2691	*2701
CATGTAATAACTGCATGTGTCTGGTCCAGGAGAAGTATTGAATAAGCCATTTCTACTGCT					
*2711	*2721	*2731	*2741	*2751	*2761
TACTCATGTCCCTATTTATGATTTCAACATGGATACATATTCAGTTCTTTCTTTTTCTC					
*2771	*2781	*2791	*2801	*2811	*2821
ACTATCTGAAAATACATTTCCCTCCCTCTCTCCCCCAATATCTCCCTTTTTTTCTCTC					
*2831	*2841	*2851	*2861	*2871	*2881
TTCTCTATCTTCCAAACCCACTTTCTCCCTCCTCTTTCTGTGTCTCTTAAGCAG					
*2891	*2901	*2911	*2921	*2931	*2941
ATAGCACATACCCCCACCCAGTACCAAATTCAGAACACAAGAAGGTCCAGTTCTTCCCC					
*2951	*2961	*2971	*2981	*2991	*3001
CTTCACATAAAGGAACATGGTTTGTGAGCCTTTCTCCTGTTTATGGGTTTCTTCCAGCAG					
*3011	*3021	*3031	*3041	*3051	*3061
AACAGAGACATTGCCAACCATATTGGATCTGCTTGCTGTCCAAACCAGGAAACTTTCTCTG					
*3071	*3081	*3091	*3101	*3111	*3121
GGCAAATCACAATCAGTGAGTAAATAGACAGCCTTTCTGCTGCCTTGGGTTTCTGTGCAG					
*3131	*3141	*3151	*3161	*3171	*3181
ATAAACAGAAATGCTCTGATTAGAAAGGAAATGAATGGTTCCACTCAAATGTCCTGCAAT					
*3191	*3201	*3211	*3221	*3231	*3241
TTAGGATTGCAGATTTCTGCCTTGAAATACCTGTTTCTTTGGGACATTCCGTCTGATGA					
*3251	*3261	*3271	*3281	*3291	*3301
TTTTTATTTTTGTTGGTTTTTATTTTTGGGGGAATGACATGTTTGGGTCTTTTATACAT					
*3311	*3321	*3331	*3341	*3351	*3361
GAAAATTTGTTTGACAATAATCTCACAAAACATATTTTACATCTGAACAAAATGCCTTTT					

*3371	*3381	*3391	*3401	*3411	*3421
TGTTTACCGTAGCGTATACATTTGTTTGGGATTTTGTGTGTTTGTGGGAATTTTGT					
*3431	*3441	*3451	*3461	*3471	*3481
TTTAGCCAGGTCAGTATTGATGAGGCTGATCATTTGGCTCTTTTTTCCTCCAGAAGAG					
*3491	*3501	*3511	*3521	*3531	*3541
TTGCATCAACAAAGTTAATTGTATTATGTATGTAAATAGATTTTAAGCTTCATTATAAA					
*3551	*3561	*3571	*3581	*3591	*3601
ATATTGTTAATGCCTATAACTTTTTTCAATTTTTTGTGTGTGTTTCTAAGGACTTTTT					
*3611	*3621	*3631	*3641	*3651	*3661
CTTAGGTTTGCTAAATACTGTAGGGAAAAAATGCTTCTTCTACTTTGTTTATTTTAGA					
*3671	*3681	*3691	*3701	*3711	*3721
CTTTAAATGAGCTACTTCTTATCACTTTTGTAAACAGCTAATAGCATGGTTCCAATTT					
*3731	*3741	*3751	*3761	*3771	*3781
TTTTTAAGTTCACTTTTTTTGTCTAGGGGAAATGAATGTGCAAAAAAAGAAAAAGAACT					
*3791	*3801	*3811	*3821	*3831	*3841
GTTGGTTATTTGTGTTATTCTGGATGTATAAAAAATCAATGGAAAAAATAAACTTTCAAA					
*3851	*3861	*3871	*3881	*3891	*3901
TTGAAATGACGGTATAACACATCTACTGAAAAAGCAACGGGAAATGTGGTCCTATTTAAG					
*3911	*3921	*3931	*3941	*3951	*3961
CCAGCCCCACCTAGGGTCTATTTGTGTGGCAGTTATTGGGTTTGGTCACAAAACATCCT					
*3971	*3981	*3991	*4001	*4011	*4021
GAAAATTTCGTGCGTGGGCTTCTTTCTCCCTGGTACAAACGTATGGAATGCTTCTTAAAGG					
*4031	*4041	*4051	*4061	*4071	*4081
GGAACTGTCAAGCTGGTGTCTTCAGCCAGATGACATGAGAGAATATCCAGAACCCTCTC					
*4091	*4101	*4111	*4121	*4131	*4141
TCCAAGGTGTTTCTAGATAGCACAGGAGAGCAGGCACTGCACTGTCCACAGTCCACGGTA					
*4151	*4161	*4171	*4181	*4191	*4201
CACAGTCGGGTGGGCCGCTCCCTCTCCTGGGAGCATTTCGTGCGTCCCGAGCCTGAGCAG					
*4211	*4221	*4231	*4241	*4251	*4261
GGCAGCTGGACTGCTGCTGTTTCCAGGAGCCACCAGAGCCTTCTCTCTTTGTACCACAGTT					
*4271	*4281	*4291	*4301	*4311	*4321

TCTTCTGTAAATCCAGTGTTACAATCAGTGTGAATGGCAAATAAACAGTTTGACAAGTAC

      |\*4331 . . . . .  
ATACACCATAtccaattggttttgccttctctgagatggagggttgggaaactgagtaa  
      . . . . .  
ggcccactgcacacactaggaagatacctccagtgaacagcttcaggctgaagctatgtc  
      . . . . .  
ctcacccccggtgctcctagcacactctggcttctgaaggcatgagggcctggaaaaggc  
      . . . . .  
gcaggctcccagggttgcttctttgcatatgaaagagcactgaagcatgttttaatgctt  
      . . . . .  
gttttaaaaaataataataggccggcgcggtggctcacgcctgtaatcccagcacgttgg  
      . . .  
gaggccgagg

LRG Parser: Version: 1.1, Version Date: 11/02/2015  
Reader: Version: 1, Version Date: 11/02/2015  
Writer: Version: 1, Version Date: 11/02/2015  
Control: Version: 1, Version Date: 11/02/2015