Gene: ABL1 - Sequence:  $NG_012034.1$ Transcript:  $NM_007313.2$  - Protein:  $NP_009297.2$ Date: February 20, 2015

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

Exon 1 | Start: 5001 | End: 5575 | Length: 574 1-439 1 - 429I-419 1-409 1-399 1 - 389GGTTGGTGACTTCCACAGGAAAAGTTCTGGAGGAGTAGCCAAAGACCATCAGCGTTTCCT 1-379 1 - 3691 - 3591 - 3491-339 1 - 329TTATGTGTGAGAATTGAAATGACTAGCATTATTGACCCTTTTCAGCATCCCCTGTGAATA |-309 |-299 1-289 |-279 TTTCTGTTTAGGTTTTCTTCTTGAAAAGAAATTGTTATTCAGCCCGTTTAAAACAAATC 1 - 2591 - 2491 - 2391 - 2291 - 2191 - 209

|-199 |-189 |-179 |-169 |-159 |-149 |-150 |-149 |-150 |-149 |-150 |-149 |-150 |-140 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150 |-150

AAGAAACTTTTGGGTAACATTGCAATTACATGAAATTGATAACCGCGAAAATAATTGGAA

|-139 |-129 |-119 |-109 |-99 |-89 CTGTGATTGACTTCAATTGCTGACTTGTGGAGATGCAGCGAATGTGAAATCCCACGTATA

|-79 |-69 |-59 |-49 |-39 |-29 TGCCATTTCCCTCTACGCTCGCTGACCGTTCTGGAAGATCTTGAACCCTCTTCTGGAAAG

| 111 | 121 | 131 TGGGGGTCCACACTGCAATGTTTTTGTGGAACATG Exon 3 | Start: 145184 | End: 145357 | Length: 173

| 141 | 151 | 161 | 171 | 181 | 191

AAGCCCTTCAGCGGCCAGTAGCATCTGACTTTGAGCCTCAGGGTCTGAGTGAAGCCGCTC

E A L Q R P V A S D F E P Q G L S E A A

| 51 | | 61

| 261 | 271 | 281 | 291 | 301 TTGCACTGTATGATTTTGTGGCCAGTGGAGATAACACTCTAAGCATAACTAAAG | 91 | 101

Exon 4 | Start: 145921 | End: 146216 | Length: 295

|491 |501 |511 |521 |531 |541 TCAATGGCAGCTTCTTGGTGCGTGAGAGTGAGAGCAGTCCTGGCCAGAGGTCCATCTCGC I N G S F L V R E S E S S P G Q R S I S |171 | |181

|551 |561 |571 |581 |591 |601 TGAGATACGAAGGGAGGGTGTACCATTACAGGATCAACACTGCTTCTGATGGCAAG |191 |201 Exon 5 | Start: 153883 | End: 154155 | Length: 272 |611 621 |631 641 |651 |661 CTCTACGTCTCCCGAGAGCCGCTTCAACACCCTGGCCGAGTTGGTTCATCATCATTCA LYVSSESRFNTLAELVHHHS |211 |221 671 1681 |691 701 |711 ACGGTGGCCGACGGCTCATCACCACGCTCCATTATCCAGCCCCAAAGCGCAACAAGCCC T V A D G L I T T L H Y P A P K R N K P |231 |241 1731 1741 1751 1761 1771 1781  ${\tt ACTGTCTATGGTGTCCCCCAACTACGACAAGTGGGAGATGGAACGCACGGACATCACC}$ T V Y G V S P N Y D K W E M E R T D I T 251 |261 791 801 811 1821 831 1841 ATGAAGCACAAGCTGGGCGGGGCCAGTACGGGGAGGTGTACGAGGGCGTGTGGAAGAAA M K H K L G G G Q Y G E V Y E G V W K K 1271 1281 |851 861 1871 TACAGCCTGACGGTGGCCGTGAAGACCTTGAAG 1291 Exon 6 | Start: 163249 | End: 163333 | Length: 84 891 901 1911 1921 1931 GAGGACACCATGGAGGTGGAAGAGTTCTTGAAAGAAGCTGCAGTCATGAAAGAGATCAAA E D T M E V E E F L K E A A V M K E I K 301 1311 941 951 |961 CACCCTAACCTGGTGCAGCTCCTTG |321 Exon 7 | Start: 163980 | End: 164157 | Length: 177 971 |981 |991 1001 1011 1021 GGGTCTGCACCCGGGAGCCCCCGTTCTATATCATCACTGAGTTCATGACCTACGGGAACC

341

G V C T R E P P F Y I I T E F M T Y G N |331

Exon 8 | Start: 165988 | End: 166172 | Length: 184

CTGGG

Exon 9 | Start: 169535 | End: 169687 | Length: 152

|1451 |1461 |1471

## ${\tt GCCCAGAGAAGGTCTATGAACTCATGCGAGCAT}$

1491

Exon 10 | Start: 171188 | End: 171277 | Length: 89

|1541 |1551 |1561 CAATGTTCCAGGAATCCAGTATCTCAGACG |521

Exon 11 | Start: 171620 | End: 171784 | Length: 164

| 1631 | 1641 | 1651 | 1661 | 1671 | 1681 | CCCCAGAGCTGCCCACCAAGACGAGGACCTCCAGGAGAGCTGCAGAGCACAGAGACACCA | A P E L P T K T R T S R R A A E H R D T | 1551 | 1561

|1691 |1701 |1711 |1721 |1731 CTGACGTGCCTGAGATGCCTCACTCCAAGGGCCAGGGAGAGAGCG |571

Exon 12 | Start: 175089 | End: 178795 | Length: 3706

| 1741 | 1751 | 1761 | 1771 | 1781 | 1791
ATCCTCTGGACCATGAGCCTGCCGTGTCTCCATTGCTCCCTCGAAAAGAGCGAGGTCCCC
D P L D H E P A V S P L L P R K E R G P | 581 | 591

| 1861 | 1871 | 1881 | 1891 | 1901 | 1911

F	S	A L  621	1	K	K	K	K	K	Т	A	P  6		Р	Р	K	R	S	S
CC'		1921 CGGGAG	ΔТС	-	931 GGC(			194		'AGA	19		ሮሮር	. –	961 GAA		•	1971 CGAG
	F		М		G						G  6	A				Е	G	R
AC		1981 AGCAAC	GGG	•	991 CTG(			200		TTG	20		GCT	•	021 CCA			2031 TCCC
D	I	S N  661	G		L			Т		L		T	A		P		K	S
C A		2041	· A A T	•	051		•	206			120		aaa		081		•	2091
P		CCCAGC P S 1681	N	G	A	G	V		N	G		L	R	E E	S	G	G	S
		2101		12	111		ı	212	:1		21			12	141		1:	2151
GC'	TTC	CGGTCT	CCC	CAC	CTG	TGG	AAG	AAG	TCC	AGC	CACG	CTG	ACC	AGC	AGC	CGC	CTA	GCCA
G	F	R S  701	P	Н	L	W	K	K	S	S	T  7		T	S	S	R	L	A
		2161		•	171			218			21			•	201		•	2211
		GAGGAG																
T	G	E E  721	Е	G	G	G	S	S	S	K	к  7	F 31	L	R	S	С	S	A
aa'		2221 GTTCCC		•	231		•	224			122		ата	•	261		•	2271
S		JIICCC V P	H	G G		AAG K	GAC D			W W	ragg R		GIC V		L		CGG R.	D
D	C	741	11	ď	А	I	ט	1	ü	W	17		V	1	ь	Г	11	D
		2281		12	291		ı	230	1		123	11		12	321		1:	2331
TG	CAG	TCCACG	GGA	AGA	CAG	TTT	GAC	TCG	TCC	ACA			GGG	CAC	AAA	AGT	GAG.	AAGC
L	Q	S T  761	G	R	Q	F	D	S	S	Т	F  7		G	Н	K	S	E	K
		2341		12	351		1	236	1		123	71		12	381		1:	2391
CG	GCT(	CTGCCT	'CGG	AAG	AGG(	GCA	GGG	GAG	AAC	AGG	TCT	GAC	CAG	GTG	ACC	CGA	GGC.	ACAG
P	A	L P  781	R	K	R	Α	G	E	N	R	S  7		Q	V	T	R	G	T
		2401		12	411		I	242	1		124	31		12	441		1:	2451

TCAGCGCCTTGATCAAGAAGAAGAAGAAGACAGCCCCAACCCCTCCCAAACGCAGCT

 ${\tt TAACGCCTCCCCCAGGCTGGTGAAAAAGAATGAGGAAGCTGCTGATGAGGTCTTCAAAG}$ 

- V T P P P R L V K K N E E A A D E V F K |801 | |811
- | 2461 | 2471 | 2481 | 2491 | 2501 | 2511 | ACATCATGGAGTCCAGCCCGGGCTCCAGCCCGCCCAACCTGACTCCAAAACCCCTCCGGC D I M E S S P G S S P P N L T P K P L R | 821 | 831
- | 2581 | 2591 | 2601 | 2611 | 2621 | 2631 GTGCCTTAGGGACCCCTGCTGCAGCTGAGCCAGTGACCCCCACCAGCAAAGCAGGCTCAG S A L G T P A A A E P V T P T S K A G S | 861 | 871
- | 2701 | 2711 | 2721 | 2731 | 2741 | 2751

  ACTCCTCTGAGTCGCCAGGGAGGGACAAGGGGAAATTGTCCAGGCTCAAACCTGCCCCGC

  H S S E S P G R D K G K L S R L K P A P | 901 | 911
- | 2821 | 2831 | 2841 | 2851 | 2861 | 2871
  AGGAGGCGGCGGGAGGCAGTCCTGGGCGCAAAGACAAAAGCCACGAGTCTGGTTGATG
  Q E A A G E A V L G A K T K A T S L V D
  | 941 | 951
- | 2881 | 2891 | 2901 | 2911 | 2921 | 2931 | 2961 | 2961 | 2961 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 | 2971 |

|981 |991

																		3051
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$																		
P	V	P S  1001		L	P	S	Α	S	S	A		A 011	G	D	Q	P	S	S
		3061		130	71		[3	308:	1		309	91		31	L01		13	3111
CC	GCC1	TTCATO	CCT	CTC	ATA	ΓCA <i>I</i>	ACCO	CGA	GTG	TCT	CTT	CGG	AAA	ACCO	CGC	CAG	CCT	CCAG
Т	A	F I  1021	_	L	Ι	S	Т	R	V	S	L  10	R 031	K	Т	R	Q	P	P
		3121		31	l31		[3	314:	1		31	51		31	l61		[3	3171
AG	CGG/	ATCGCC	AGC	GGCC	GCC/	ATC!	ACC/	AAG	GGC	GTG	GTC	CTG	GAC	AGC/	ACCO	GAG	GCG	CTGT
E	R	I A  1041		G	Α	Ι	Т	K	G	V	V  10	L 051	D	S	Т	E	Α	L
		3181		31	L91		[3	320:	1		32:	11		32	221		[3	3231
GC	CTC	GCCATC	TCT	AGG <i>I</i>	AACT	CCC(	GAG	CAG	ATG	GCC	AGC	CAC	AGC(	GCAC	GTG	CTG	GAG	GCCG
С	L	A I	S	R	N	S	E	Q	M	Α	S	Н	S	Α	V	L	E	Α
		1061									10	071						
		3241		132	251		[3	326:	1		32	71		32	281		[3	3291
GC	AAA <i>I</i>	ACCTC	TAC	ACGT	TC:	rgc(	GTG/	\GC'	ΓΑΤ	GTG	GAT	TCC	ATC	CAGO	CAA	ATG	AGG	AACA
G	K	N L	Y	T	F	C	V	S	Y	V	D	S	Ι	Q	Q	М	R	N
		1081									10	091						
		3301		33	311		[3	332:	1		33:	31		33	341		[3	3351
AG	TTT	GCCTTC	CGA	GAGO	GCC/	ATC <i>I</i>	AAC <i>I</i>	AAA	CTG	GAG	AAT	AAT	CTC	CGGC	GAG	CTT	CAG	ATCT
K	F	A F	R	E	Α	I	N	K	L	E	N	N	L	R	E	L	Q	I
		1101									1:	111						
		3361		33	371		[3	338:	1		339	91		34	101		:	3411
GC	CCG	GCGACA	GCA(	GGC <i>I</i>	AGTO	GTC	CCAC	GCG(	GCC	ACT	CAG	GAC:	ГТС	AGC/	AAG	CTC	CTC	AGTT
С	P	A T	Α	G	S	G	P	Α	Α	T	Q	D	F	S	K	L	L	S
		1121									1:	131						
		3421		34	131		[3	344:	1		34	51		+	<b>-11</b>			+21
CG	$\tt CGGTGAAGGAAATCAGTGACATAGTGCAGAGGTAGCAGCAGTCAGGGGTCAGGTGTCAGG$																	
S	V	K E	I	S	D	Ι	V	Q	R	*								
		1141									1:	151						
		+31		+	-41			+5:	1		+(	61		+	<b>⊦</b> 71			+81
CC	CGT	CGGAGC																ACAA

+121

|+131

|+141

|+111

+91

|+101

GGGACT	AGTGAGTCA	GCACCTTGGC	CCAGGAGCTC	rgcgccaggc <i>i</i>	AGAGCTGAGGO	CCCT
ашаала				+181		
GTGGAG				CTGCCCTCCC		
CCGCTC				+241 GTTCCTGCTC		
CCATCC				+301  GCCCCAGACT		
donido						
A ССТСС				+361  TTTTTCTCTC		
AGGIGG	GAACGGCIGA	AIGIGGACIG.	ICITITICAT	11111101010	JIGGAGCCCCI	.0010
22222				+421		
CCCCGG	CTGGGCCTCC	CTTCTTCCAC.	ITCTCCAAGA <i>I</i>	ATGGAAGCCTO	JAACTGAGGCC	TTGT
				+481		
GTGTCA	GGCCCTCTG	CCTGCACTCC	CTGGCCTTGC	CCGTCGTGTGC	CTGAAGACATO	TTTC
	+511	l+521	l+531	l+541	+551	+561
AAGAAC	CGCATTTCG	GGAAGGGCAT	GCACGGGCAT	GCACACGGCT	GGTCACTCTGC	CCCTC
	+571	l+581	l+591	l+601	+611	+621
TGCTGC	TGCCCGGGGT	rggggtgcac:	rcgccatttc(	CTCACGTGCAC	GGACAGCTCTT	GATT
	l+631	l+641	l+651	l+661	l+671	+681
TGGGTG	GAAAACAGG	GTGCTAAAGC	CAACCAGCCT	TTGGGTCCTG	GGCAGGTGGG <i>A</i>	AGCTG
	+691	+701	+711	+721	+731	+741
AAAAGG	ATCGAGGCAT	rggggcatgt(	CCTTTCCATC	rgtccacatco	CCCAGAGCCCA	AGCTC
	<del>+</del> 751	l+761	+771	+781	<del>+</del> 791	+801
TTGCTC				AGAAAGCTTG <i>I</i>		
	l+811	l+821	l+831	l+841	l+851	l+861
AGGTCA				GAATGGAGGC <i>I</i>		
	l±971	l±001	l±801	+901	I±011	1±021
CAGTGG				GCCCCAGACT(		
	1.021	1.041	LiOE1	+961	1.071	1.004
TGCAAG	•	-	•	T+961 CTGGGTGCCCT	-	•

|+1021

|+1031

|+1041

+1011

 $\tt CCTCCCCACTCCTCTAAGACAAAGTAGATTCTTACAAGGCCCTTTCCTTTGGAACAAGA$ 

+991

+1001

- |+1051 |+1061 |+1071 |+1081 |+1091 |+1101 |
  CAGCCTTCACTTTTCTGAGTTCTTGAAGCATTTCAAAGCCCTTGCCTCTGTGTAGCCGCCC
- |+1111 |+1121 |+1131 |+1141 |+1151 |+1161 TGAGAGAGAATAGAGCTGCCACTGGGCACCTGCGCACAGGTGGGAGGAAAGGGCCTGGCC
- |+1171 |+1181 |+1191 |+1201 |+1211 |+1221
  AGTCCTGGTCCTGGCTGCACTCTTGAACTGGGCGAATGTCTTATTTAATTACCGTGAGTG
- |+1231 |+1241 |+1251 |+1261 |+1271 |+1281 ACATAGCCTCATGTTCTGTGGGGGTCATCAGGGAGGGTTAGGAAAACCACAAACGGAGCC
- |+1291 |+1301 |+1311 |+1321 |+1331 |+1341 |+1341 |+1360 |
- |+1351 |+1361 |+1371 |+1381 |+1391 |+1401 GGCCGGAGCCCAGATACGGGGGCTGTGACTCTGGGCAGGGACCCGGGGTCTCCTGGACCT
- |+1411 |+1421 |+1431 |+1441 |+1451 |+1461 TGACAGAGCAGCTAACTCCGAGAGCAGTGGCCAGGTGGCCGCCCCTGAGGCTTCACGCCG
- |+1471 |+1481 |+1491 |+1501 |+1511 |+1521 GGAGAAGCCACCTTCCCACCCCTTCATACCGCCTCGTGCCAGCAGCCTCGCACAGGCCCT
- |+1531 |+1541 |+1551 |+1561 |+1571 |+1581 AGCTTTACGCTCATCACCTAAACTTGTACTTTATTTTTCTGATAGAAATGGTTTCCTCTG
- |+1591 |+1601 |+1611 |+1621 |+1631 |+1641 |
  GATCGTTTTATGCGGTTCTTACAGCACATCACCTCTTTGCCCCCGACGGCTGTGACGCAG
- |+1651 |+1661 |+1671 |+1681 |+1691 |+1701 |+1681 |+1691 |+1701 |
- |+1771 |+1781 |+1791 |+1801 |+1811 |+1821 |
  CACTATATTTTACACGTATCTCTTGGTATGCATCTTTTATAGACGCTCTTTTCTAAGTGG
- |+1831 |+1841 |+1851 |+1861 |+1871 |+1881 |+1871 |+1881 |

## |+1951 |+1961 |+1971 |+1981 |+1991 TCTGTCCTCTGTAGTATTTTTTAAATAAATCAGTGTTTACATTAGAA

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