Gene: WELLAND - Sequence: NG_Welland.1 Transcript: NM_000048.3 - Protein: FAKE_PROTEIN Date: February 12, 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 FIRST base of codon 4^{th} line: Amino acid numbering. Numbered on 1^{st} and increments of 10 Exon 1 | Start: 800 | End: 1020 | Length: 220 1-209 l-169 I-159 |-109

aaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaa													
aaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaa													
aaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaa													
Exon 2 Start: 1620 End: 1779 Length: 159													
gaaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaa													
gaaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaa													
gaaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaa													
gaaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaa													
gaaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaa													
-49 -39 -29 -19 -9 1 ACGTACGTACGTACGTACGTACGTACGTACGTACGTACGT													
11 21 31 41 51 61													
CGTACGTACGTACGTACGTACGTACGTACGTACGTACGTA													
C D E F G H I J K L M N O P Q R S T U V													
71 81 91 101													
CGTACGTACGTACGTACGTACGTACGTACGTACGTACGTA													
W X Y Z A B C D E F G H I													

aaag	aaa	aga:	aaag	gaaa	aaga	aaa	.gaa	aaį	gaaa	aag	aaa	agaa	aaa	gaaa	aaga	aaa	aga	aaa	ga
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aaagaaaagaa																			
Exon	. 3	S	tart	5: 2	2379	I	End	l: :	265	5	Le	ngtl	n: 1	276					
gaaa	.aga	.aaaį	gaaa	aaga	aaa	gaa	.aag	gaa	aaga	aaa	aga	aaag	gaa	aaga	aaaa	agaa	aaa	gaa	aa
gaaa	.aga	aaa	gaaa	aaga	aaaa	gaa	aag	gaa	aaga	aaa	aga	aaag	gaa	aaga	aaaa	agaa	aaa	gaa	aa
gaaa	.aga	.aaaį	gaaa	aaga	aaaa	gaa	aag	gaa	aaga	aaa	aga	aaag	gaa	aaga	aaaa	agaa	aaa	gaa	aa
gaaa	.aga	aaa	gaaa	aaga	aaaa	gaa	aag	gaa	aaga	aaa	aga	aaag	gaa	aaga	aaaa	agaa	aaa	gaa	aa
gaaa	.aga	.aaaį	gaaa	aaga	aaaa	gaa	aag	gaa	aaga	aaa	aga	aaag	gaa	aaga	aaaa	agaa	aaa	gaa	aa
ACGT J					12 ACGT 0 41	ACG P		CGT			GTA		\CG			CGT. Z		TAC	
ACGT D		17 TAC F	GTAC		CGT.	ACG J	TAC	CGT	ACG:	ГАС	GTA	201 CGT <i>I</i> P	ACG	TAC	GTAC	CGT. T	ACG' U	TAC	GΤ

		123:	L		12	41		ı	251			126	1		12	71	+1	
${\tt ACGTACGTACGTACGTACGTACGTACGTACGTACGTACGT$													ACGT					
Х	Y	Z	Α	В	C	D	Е	F	G	Н	Ι	J	K	L	М	*		
	_	_			18			_	_		_	-			19	1		
					10	_									10	_		
							١.	0.4									1.0	
	+11													+5				
ACGTACGTACGTACGTACGTACGTACGTACGTACGTACGT															CACGT			
	+71			+8	1		+	91		- 1	+10	1						
ACGT	'ACG'	TAC	GTA(CGT.	ACG'	TAC	GTA	CGT	ACG	TAC	:GTa	aaa	gaa	aaag	aaa	aga	aaag	gaaaa
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gaaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaa														gaaaa				
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gaaaagaaaagaaaagaaaagaaaagaaaagaaaagaaaa																		
•	•		•			•		•			•		•	•		•		
gaaaagaaaagaaaagaaaagaaaagaaaagaaaaga														gaaaa				
gaaa	າລຫລ	222	ກລລະ	າລຕ	aaa	ລσລ	222	σaa	່ລລດ	ลลล	aσ							
Sauce	-u5u	عسسع	وسمر	اكمت	uuu	سوم	Juu	صو	uug	uuc	مه							

GBK Parser: Version: 0.3, Version Date: 11/02/2015 Reader: Version: 0.3, Version Date: 11/02/2015 Writer: Version: 0.3, Version Date: 11/02/2015 Control: Version: 0.3, Version Date: 11/02/2015