

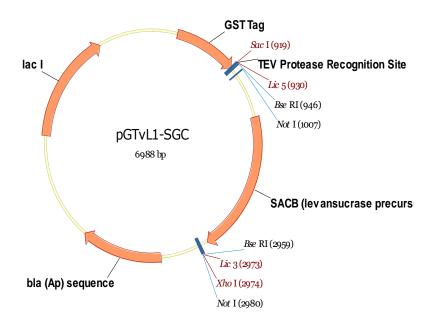
# Vector information sheet

Dated: 8<sup>th</sup> May 2013

Vector Name	pGTvL1-SGC
Source	Jonathan Elkins (SGC, Oxford)
Sequence accession/link	(SGC)

Description	pGEX expression vector with N-terminal GST tag and TEV				
	protease cleavage site. Includes sites for LIC cloning, and a				
	"stuffer" fragment that includes the SacB gene, allowing				
	negative selection on 5% sucrose.				

Antibiotic resistance	ampicillin
Promoter	Tac promoter (lac/IPTG inducible)
Cloning	LIC. (vector treated with BseRI, then with T4 DNA polymerase in presence of dGTP)
Initiation codon	(readthrough from GST gene).
N-terminal fusion – seq.	MSPILGYWKIKGLVQPTRLLLEYLEEKYEEHLYERDEGDKWRNKKFELGL EFPNLPYYIDGDVKLTQSMAIIRYIADKHNMLGGCPKERAEISMLEGAVL DIRYGVSRIAYSKDFETLKVDFLSKLPEMLKMFEDRLCHKTYLNGDHVTH PDFMLYDALDVVLYMDPMCLDAFPKLVCFKKRIEAIPQIDKYLKSSKYIA WPLQGWQATFGGGDHPPKSSSENLYFQ*S(M) (* - TEV cleavage site)
N-terminal fusion – MW	26752.6 including Methionine (26534.3 removed by TEV cleavage)
Termination codons	supplied in PCR primer
Protease cleavage	TEV protease
Additional features	
Preferred host	Many E. coli strains (not dependent on T7 RNA polymerase)
5' sequencing primer	pGEX-5': GGGCTGGCAAGCCACGTTTGGTG
3' sequencing primer	pGEX-3': CCGGGAGCTGCATGTGTCAGAGG



### Polylinker region:

## GST protein-····>

						M	S	P	I	L	G	Y	M	K	I	K	G	L	V	Q	
241	CACA	GGA.	AAC	AGTATTCATG			TCCCCTATAC			TAGGTTATTG			GAAAATTAAG			GGCCTTGTGC					
	· P	T	R	L	L	L	Ε	Y	L	Ε	E	K	Y	E	E	Н	L	Y	Ε	R	
301	AACC	CAC	TCG	ACTTCTTTTG			GAATATCTTG			AAGAAAAATA			TGAAGAGCAT			TTGTATGAGC					
	· D	E	G	D	K	W	R	N	K	K	F	Ε	L	G	L	E	F	P	N	L	
361	GCGA	TGA.	AGG	TGATAAATGG			CGAAACAAAA			AGTTTGAATT			GGGTTTGGAG			TTTCCCAATC					
	· P	Y	Y	I	D	G	D	V	K	L	T	Q	S	M	Α	I	I	R	Y	I	
421	TTCC	TTA	TTA	TAT	TGA	TGGT	GΑ	TGT	TAA	ΑT	TAAC	ACA	GTC	TAT	GGC	CATC	AT	ACG	TTA	TΑ	
	· A	D	K	Н	N	M	L	G	G	С	P	K	E	R	Α	E	I	S	M	L	
481	TAGC	TGA	CAA	GCA	CAA	CATG	TT	GGG	TGG	ΤТ	GTCC	AAA	AGA	GCG	TGC	AGAG	ΑT	TTC	AAT	GC	
	· E	G	Α	V	L	D	I	R	Y	G	V	S	R	I	Α	Y	S	K	D	F	
541	TTGA	AGG.	AGC	GGT	TTT	GGAT	ΑT	TAG	ATA	.CG	GTGT	TTC	GAG	AAT	TGC	TATA	AG	TAA	AGA	СТ	
	E	T	L	K	V	D	F	L	S	K	$_{ m L}$	P	E	M	L	K	Μ	F	Ε	D	
601	TTGA	AAC	TCT	CAAAGTTGAT		TGAT	TTTCTTAGCA		AGCTACCTGA		AATGCTGAAA			ATGTTCGAAG							
	· R	L	С	Н	K	T	Y	L	N	G	D	Н	V	T	Н	P	D	F	M	L	
661	ATCG	TTT.	ATG	TCA	TAA	AACA	TΑ	TTT	AAA	ΤG	GTGA	TCA	TGT	AAC	CCA	TCCT	GΑ	СТТ	CAT	GΤ	
	• У	D	Α	L	D	V	V	L	Y	Μ	D	P	M	С	L	D	Α	F	P	K	
721	TGTA	TGA	CGC	TCT	TGA	TGTT	GΤ	TTT	ATA	CA	TGGA	CCC	AAT	GTG	CCT	GGAT	GC	GTT	CCC.	AA	
	· L	V	С	F	K	K	R	I	Ε	Α	I	P	Q	I	D	K	Y	L	K	S	
781	AATT	AGT	TTG	TTT	TAA	AAAA	CG	TAT	TGA	AG	CTAT	CCC	ACA	AAT	TGA	TAAG	ΤA	СТТ	GAA.	AΤ	
	· S	K	Y	I	Α	M	Ρ	L	Q	G	W	Q	Α	Т	F	G	G	G	D	Η	
841	CCAG	AGCAAGTA TATAGCATGG		ATGG	CCTTTGCAGG				GCTG	AGC	CAC	CACGTTTGGT GGTG				CGA	CC				

901 ATCCTCCAAA ATCGAGCTCA GAGAACCTGT ACTTCCAATC CATAAGCTAG CTTCTCCTCC 5' LIC sequence BseRI

<	····· (SacB spacer)
•••••	•>

2941 TCGAGGAGTT TAC**TAGTAAG TAAAGGTGGA TA**CTCGAGCG GCCGCATCGT GACTGACTGA

\*\*BseRI\*\* 3' LIC sequence XhoI\*\* NotI

Primers for LIC cloning:

Upstream: add TACTTCCAATCCATG to the 5' end (ATG in-frame with the desired coding sequence).

Downstream: add TATCCACCTTTACTG to 5' end of downstream primer; add termination codon, if necessary.

#### pGTvL1-SGC sequence:

 ${\tt acgttatcgactgcaccggtgcaccaatgcttctggcgtcaggcagccatcggaagctgtggtatggctg}$ tgcaggtcgtaaatcactgcataattcgtgtcgctcaaggcgcactcccgttctggataatgttttttg cgccgacatcataacggttctggcaaatattctgaaatgagctgttgacaattaatcatcggctcgtataatgtgtggaattgtgagcggataacaatttcacacaggaaacagtattcatgtcccctatactaggttattggaaaattaagggccttgtgcaacccactcgacttcttttggaatatcttgaagaaaaatatgaag agcatttgtatgagcgcgatgaaggtgataaatggcgaaacaaaaagtttgaattgggttttggagtttc $\tt ccaatcttccttattattattgatggtgatgttaaattaacacagtctatggccatcatacgttatatag$ ctgacaagcacaacatgttgggtggttgtccaaaagagcgtgcagagatttcaatgcttgaaggagcgg  $\verb|ttttggatattagatacggtgtttcgagaattgcatatagtaaagactttgaaactctcaaagttgatt|\\$  $\verb|ttcttagcaagctacctgaaatgctgaaaatgttcgaagatcgtttatgtcataaaacatatttaaatg|$  $\tt gtgatcatgtaacccatcctgacttcatgttgtatgacgctcttgatgttgttttataccatggacccaa$  ${\tt tgtgcctggatgcgttcccaaaattagtttgttttaaaaaaacgtattgaagctatcccacaaattgata}$  ${\tt agtacttgaaatccagcaagtatatagcatggcctttgcagggctggcaagccacgtttggtggtggcg}$  $\verb|atccata| act tcgt a tag cata cat tatac gaa get tat gcggccgcgacgtccacat at acct gccgt tatacgcd gas get to the control of th$  $\verb|cactattatttagtgaaatgagatattatgatattttctgaattgtgattaaaaaggcaactttatgcc|$  $\verb|cgagagtcta| at a gaat | \verb|gagatcgaaaagta| a at \verb|cgcgcgggtttgtta| ctgataaagcaggcaagacc| a gaat | \verb|gagagtcta| a gaat | cgagagatcta| a gaat | cgagagatc$ taaaatgtgtaaagggcaaagtgtatactttggcgtcaccccttacatattttaggtctttttttattgtgcgtaactaacttgccatcttcaaacaggaggctggaagaagcagaccgctaacacagtacataaaa aaggagacatgaacgatgaacatcaaaaagtttgcaaaacaagcaacagtattaacctttactaccgca ggcatttcccatattacacgccatgatatgctgcaaatccctgaacagcaaaaaaatgaaaaatatcaa qttcctqaattcqattcqtccacaattaaaaatatctcttctqcaaaaqqcctqqacqtttqqqacaqc tggccattacaaaacgctgacggcactgtcgcaaactatcacggctaccacatcgtctttgcattagcc qqaqatcctaaaaatqcqqatqacacatcqatttacatqttctatcaaaaaqtcqqcqaaacttctatt qacaqctqqaaaaacqctqqccqcqtctttaaaqacaqcqacaaattcqatqcaaatqattctatccta aaagaccaaacacaagaatggtcaggttcagccacatttacatctgacggaaaaatccgtttattctac actgatttctccggtaaacattacggcaaacaacactgacaactgcacaagttaacgtatcagcatca cactacgtagaagataaaggccacaaatacttagtatttgaagcaaacactggaactgaagatggctac caaggcgaagaatctttatttaacaaagcatactatggcaaaagcacatcattcttccgtcaagaaagt  $\verb|caa| a a a cttctg | caa a a g | cgata a a a a a cgca cgg | ctgag | ctag | ctgag |$  $\verb|cta| a a c gat gat ta ca cact gaa a a a a g t gat gaa a c c g c t gat t g cat c t a a c a c a g a t g a a cac g a t g a cac g a$  $\verb|attgaacgcgcgaacgtctttaaaatgaacggcaaatggtacctgttcactgactcccgcggatcaaaa| \\$  $\verb|atgacg| attgacggcattacgtctaacgatatttacatgcttggttatgtttctaattctttaactggc| |$  $\verb|ccatacaagccgctgaacaaaactggccttgtgttaaaaatggatcttgatcctaacgatgtaaccttt|\\$ agaggattctacgcagacaaacaatcaacgtttgcgccaagcttcctgctgaacatcaaaggcaagaaa

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