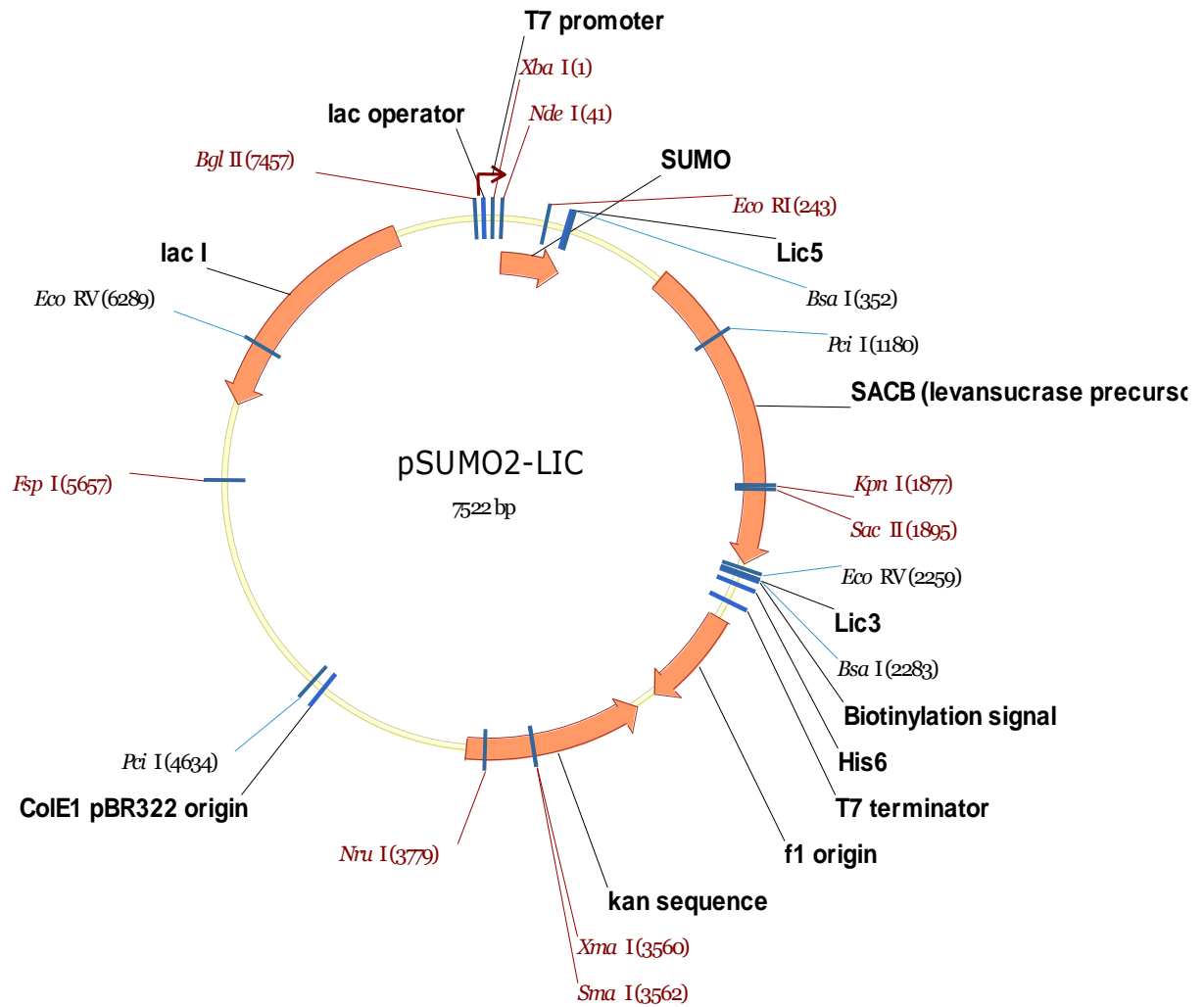


Vector information sheet

Dated: 16th September 2013

Vector Name	pSUMO2-LIC
Source	Pavel Savitsky
Sequence accession/link	Genbank XXXXXXXX
Description	pET expression vector with N-terminal SUMO (96-aa sequence derived from SUMO1 protein) tags followed by a SUMO protease cleavage site and C-terminal biotinylation and I His ₆ tags. SUMO tag has been shown to improve solubility of the fusion partners. The vector includes sites for LIC cloning, and a "stuffer" fragment that includes the SacB gene, allowing negative selection on 5% sucrose.
Antibiotic resistance	Kanamycin, 50 µg/ml
Promoter	T7 - lacO
Cloning	LIC. (vector treated with BsaI, then with T4 DNA polymerase in presence of dTTP)
Initiation codon	Supplied in PCR primer
N-terminal fusion – seq.	MCSSGSGSGSDQEAKPSTEDLGDKKEGEYIKLVIGQDSSEI HFKVKMTTHLKKLKESYQQRQGVPMNSLRFLFEGQRIADNHT PKELGMEEEDVIEVYQEQTGG* (* - SUMO cleavage site)
N-terminal fusion – MW	11755.1 Da
C-terminal fusion – seq	SGGGLNDIFEAQKIEWHEHHHHHH
C-terminal fusion – MW	2853 Da
Termination codons	No stop codon
Protease cleavage	SUMO
Additional features	.
Preferred host	DE3 hosts: BL21, Rosetta, etc. MUST express T7 RNA polymerase.
5' sequencing primer	pLIC-for: TGTGAGCGGATAACAATTCC
3' sequencing primer	pLIC-rev: AGCAGCCAACTCAGCTTCC



Cloning region in the vector:

```

XbaI                                     NdeI
~~~~~                                ~~~~~
1  CTAGAAATAA TTTTGTTTAA CTTTAAAGAAG GAGATATACA TATGTGCAGC AGCGGCAGCG
   · S G S D Q E A K P S T E D L G D K K E G ·
61 GCAGCGGCTC TGACCAGGAG GCAAAACCTT CAACTGAGGA CTTGGGGGAT AAGAAGGAAG
   · E Y I K L K V I G Q D S S E I H F K V K ·
121 GTGAATATAT TAAACTCAA GTCATTGGAC AGGATAGCAG TGAGATTCAC TTCAAAGTGA
   · M T T H L K K L K E S Y C Q R Q G V P M ·
181 AAATGACAAC ACATCTCAAG AACTCAAAG AATCATACTG TCAAAGACAG GGTGTTCCAA

EcoRI
~~~~~
241 TGAATTCAC T CAGGTTTCTC TTTGAGGGTC AGAGAATTGC TGATAATCAT ACTCCAAAAG

                                     Lic5      BsaI
                                     ~~~~~      ~~~~~
301 AACTGGGAAT GGAGGAAGAA GATGTGATTG AAGTTTACCA GGAGCAAACG GGAGGTGAGA

BsaI
~~~
361 CCGACGTCCA CATATACCTG CCGTTCCTA TTATTTAGTG AAATGAGATA TTATGATATT
  
```

SacB

```

                                EcoRV                                BsaI
                                ~~~~~~                                ~~~~~~
2221 AACAGTTAAC AAATAAAAC GCAAAAGAAA ATGCCGATAT CCTATTGGCA TTGACGGTCT

BsaI
~
  S G G G L N D I F E A Q K I E W H E H H .
2281 CAGCGGTGGC GGTCTGAACG ACATCTTCGA GGCTCAGAAA ATCGAATGGC ACGAACACCA

. H H H H *
2341 CCACCACCAC CACTGAGATC CGGCTGCTAA CAAAGCCCGA AAGGAAGCTG AGTTGGCTGC

```

Lic cloning scheme:

Treat vector with dTTP:

```

V Y Q E Q T G      G      BsaI
G TTT ACCAGGAGCAAACGG      GAGGT GAGACC GACGTCCACATataacctgccgttcactattattt
CAAAATGGTCCTCGTTTGCCCTCC      ACTCTGG CTGCAGGTGTAtatggacggcaagtataataaa

```

(SacB fragment)

```

                                BsaI      S G G G L N D I F E A Q
GATATCCTAT TGGCATTGAC GGTCTCA GCGGTGGCGGTCTGAACGACATCTTCGAGGCTCAG
CTATAGGATA ACCGTAAC TG CAGAGTCGC CACCGCCAGACT GCTGTAGAAGCTCCGAGTC

K I E W H E H H H H H H
AAAATCGAATGGCACGAACA CCACCACCAC CACCACTGAGATCCGGCTGC
TTTAGCTTACCGTGCTTGT GGTGGTGGTG GTGGT GACTTAGGCCGACG

```

Treat PCR product with dATP:

```

ACCAGGAGCAAACGGGAGGT      PCR product      AGCGGTGGCGGTCTG
TGGTCCTCGTTTGCCCTCC      TCGCCACCGCCAGAC

```

Primers LIC extensions:

```

Forward      ACCAGGAGCAAACGGGAGGT
Reverse      CAGACCGCCACCGCT

```

Primers for LIC cloning (treat PCR fragment with T4 DNA polymerase in presence of dATP)

Upstream: add ACCAGGAGCAAACGGGAGGT to the 5' end of upstream primer

Downstream: add CAGACCGCCACCGCT to 5' end of downstream primer

>pSUM02-LIC

```

ctagaaataatTTTTgtttaactttaagaaggagatatatacatatgtgcagcagcggcagcggcagcggct
ctgaccaggaggcaaaaccttcaactgaggacttgggggataagaaggaaggtgaatatattaaactca
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```

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