



pSUPER RNAi System™

pSUPER : Manual

A Vector System for Expression of Short Interfering RNA

The following manual contains information about and
Instructions for the following pSUPER vectors:

<u>Vector Name</u>	<u>Catalog#</u>
pSUPER	VEC-PBS-0001 (linear) VEC-PBS-0002 (circular)
pSUPER.neo	VEC-PBS-0003 (linear) VEC-PBS-0004 (circular)
pSUPER.neo+gfp	VEC-PBS-0005 (linear) VEC-PBS-0006 (circular)
pSUPER.puro	VEC-PBS-0007 (linear) VEC-PBS-0008 (circular)
pSUPER.p53	VEC-P53-0001 (circular)

NEW : BgIII / XhoI Oligo Insert Design Option – See Inside...

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PRODUCT INFORMATION

Concentration:	0.5 mg/ml
Volume:	12 µl
Buffer:	10mM Tris-HCl pH 7.4, 1mM EDTA
Storage:	Store at 4°C
Shipping:	<i>pSUPER is shipped to customers at ambient temperature to reduce shipping and handling costs without affecting product quality and effectiveness.</i>

BACKGROUND: THE pSUPER RNAi SYSTEM

In several organisms, introduction of double-stranded RNA has proven to be a powerful tool to suppress gene expression through a process known as RNA interference (1). However, in most mammalian cells this provokes a strong cytotoxic response (2). This non-specific effect can be circumvented by use of synthetic short [21- to 22-nucleotide (nt)] interfering RNAs (siRNAs), which can mediate strong and specific suppression of gene expression (3). However, this reduction in gene expression is transient, which severely restricts its applications.

To overcome this limitation, the pSUPER RNAi system provides a mammalian expression vector that directs intracellular synthesis of siRNA-like transcripts. The vector uses the polymerase-III H1-RNA gene promoter, as it produces a small RNA transcript lacking a polyadenosine tail and has a well-defined start of transcription and a termination signal consisting of five thymidines in a row (T5). Most important, the cleavage of the transcript at the termination site is after the second uridine, yielding a transcript resembling the ends of synthetic siRNAs, which also contain two 3' overhanging T or U nucleotides (nt).

The pSUPER RNAi System has been used to cause efficient and specific down-regulation of gene expression (4, 5), resulting in functional inactivation of the targeted genes. Stable expression of siRNAs using this vector mediates persistent suppression of gene expression, allowing the analysis of loss-of-function phenotypes that develop over longer periods of time.

References:

1. P. A. Sharp, *Genes Dev.* 13, 139 (1999).
2. T. Hunter, T. Hunt, R. J. Jackson, H. D. Robertson, *J. Biol. Chem.* 250, 409 (1975).
3. S. M. Elbashir et al., *Nature* 411, 494 (2001).
4. T.R. Brummelkamp, R. Bernards, and R. Agami, *Science* 296, 550 (2002).
5. T.R. Brummelkamp, R. Bernards, and R. Agami, *Cancer Cell* Published online Aug. 22, 2002.

OLIGO INSERT DESIGN

To effect the silencing of a specific gene, the pSUPER vector is used in concert with a pair of custom oligonucleotides that contain, among other features, a unique 19-nt sequence derived from the mRNA transcript of the gene targeted for suppression (the "N-19 target sequence").

The N-19 target sequence corresponds to the sense strand of the pSUPER-generated siRNA, which in turn corresponds to a 19-nt sequence within the mRNA. In the mechanism of RNAi, the antisense strand of the siRNA duplex hybridizes to this region of the mRNA to mediate cleavage of the molecule.

These forward and reverse oligos are annealed and cloned by the user into the vector, between the unique BglIII and HindIII enzyme sites. This positions the forward oligo at the correct position downstream from the H1 promoter's TATA box to generate the desired siRNA duplex.

The sequence of this forward oligo includes the unique N-19 target in both sense and antisense orientation, separated by a 9-nt spacer sequence. The 5' end corresponds to the BglIII site, while the 3' end contains the T5 sequence and any HindIII-corresponding nucleotides. *NOTE that while the 5' overhang of the oligo corresponds to the 3' BglIII overhang of the plasmid, the overhang sequence of the oligo actually corresponds to the BamHI, and thus destroys the BglIII site upon ligation to enable more efficient screening of positive clones.*

The resulting transcript of the recombinant vector is predicted to fold back on itself to form a 19–base pair stem-loop structure. Analysis indicates that the stem-loop precursor transcript is quickly cleaved in the cell to produce a functional siRNA (4). Figure 1 provides an overview of the insert design, and how the oligos are transcribed and process to functional siRNA.

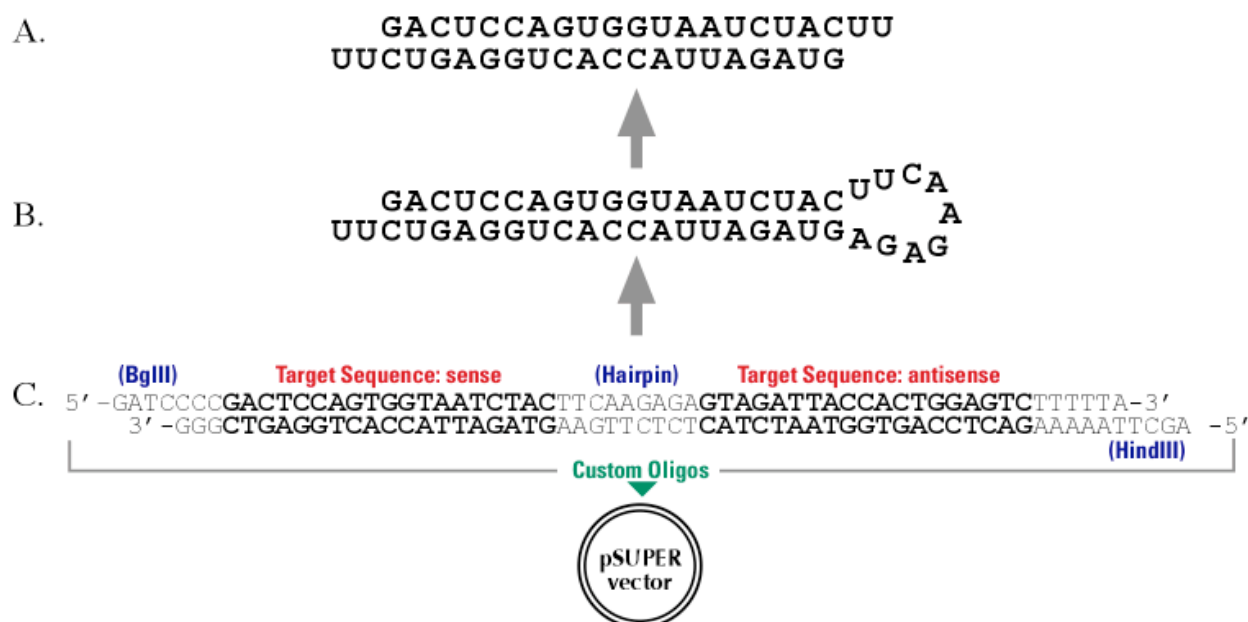


Fig. 1: Transcription of 60-nt oligo to hairpin RNA, processed to functional siRNA.

NEW: BglIII / XhoI INSERT OLIGO DESIGN

To facilitate easier linearization of the pSUPER vector, OligoEngine now offers the option to purchase oligo inserts with the following 5' / 3' ends:

- BglIII / HindIII (original format)
- BglIII / XhoI (new format)

When designing and/or purchasing oligos, the OligoEngine workstation gives users the option to select either configuration. See below for more information about using the workstation in conjunction with pSUPER, and refer to the Procedure section for instructions on using BglIII / XhoI oligos in pSUPER vectors.

OLIGOENGINE RNAi DESIGN TOOLS

It has been shown that a single nucleotide mismatch in the 19-nt targeting sequence abrogates the ability to suppress gene expression (4). Therefore, sequence design is critical.

OligoEngine provides a design tool for the pSUPER RNAi System – as well as for the SI² Silencing Duplex – that generates N-19 target sequences for any gene of interest. The tool can be accessed by clicking on the “Order Now” tab at the top of any page on our Web site, <http://www.oligoengine.com>. Or, you can click the “Download Workstation” link from the Web site home page to run this tool as a stand-alone application.

The RNAi Design Tool automates the target design process recommended based on the most recent published research on RNAi mechanisms, as well as our own proprietary design algorithms. It helps users choose and configure these oligos by analyzing a their gene sequence and applying various algorithms according to the chosen design method and user parameters.

Once the design is complete you can order your oligos right from the Design Tool. These are synthesized with BglIII (BamHI) and HindIII OR BglIII (BamHI) and XhoI ends, so no digestion is required prior to cloning.

For more information and instructions, visit www.oligoengine.com.

PROCEDURE

Outline

Here are the general steps for an experiment utilizing a pSUPER vector*:

1. Anneal the forward and reverse strands of the oligos that contain the siRNA-expressing sequence targeting your gene of interest.
2. Linearize the pSUPER vector with BglII and XhoI OR HindIII (**OMIT this step if you purchased a linear vector**)
3. Clone the annealed oligos into the vector
4. Transform the vector in bacteria
5. Transfect pSUPER vector into mammalian cells
6. Monitor EGFP fluorescence (for “+GFP” versions only)
7. Select with puromycin or neomycin to establish a stable cell line for siRNA expression (.neo or .puro versions)
8. Assay the effects on protein expression and/or mRNA levels

**Note that the pSUPER.p53 vector contains the p53-targeted insert already cloned into the pSUPER.basic vector.*

General Molecular Biology Techniques

For many of the steps described below you may use the method of choice for your lab or level of experience. For assistance with transformations, restriction enzyme digestion and analysis, DNA purification, sequence and biochemistry, please refer to *Molecular Cloning: A Laboratory Manual* (Sambrook *et al.*, 1989) or *Current Protocols in Molecular Biology* (Ausubel *et al.*, 1994). See the “Lab Notes” and “Lab Tips” for recommendations based on the experience of the vector’s inventors, and on feedback from our research customers

>> Step One: Anneal Oligos

Obtain two DNA oligonucleotides for hairpin RNA expression. Our experience demonstrates that gel purification of the oligos is not necessary for efficient ligation.

Dissolve the oligos in sterile, nuclease-free H₂O to a concentration of 3 mg/ml. If you need assistance to determine how much H₂O (or buffer, etc.) to add to your product on hand, see the Lab Tips section in this manual following the procedure.

Assemble the annealing reaction by mixing 1 µl of each oligo (forward + reverse) with 48 µl annealing buffer. You may choose to either a) use a similar buffer from another manufacturer, or b) create your own buffer stock from 100 mM NaCl and 50 mM HEPES pH 7.4.

Incubate the mixture at 90°C for 4 min, and then at 70°C for 10 minutes. Slowly cool the annealed oligos to 10°C (e.g., step-cool to 37 for 15-20 minutes, then to 10°C or room temperature before using or moving them to refrigerated storage). The annealed oligo inserts can be used immediately in a ligation reaction, or cooled further to 4°C. For longer storage, keep at -20°C until needed.

>> Step Two: Linearize the Vector (Omit if you have the “Linear” version of this vector)

Linearize 1 µl of the pSUPER vector with BglII and either HindIII OR XhoI restriction enzymes, depending on the configuration of your oligos as previously discussed. It is recommended to perform sequential reaction steps rather than simultaneous digestion, as follows: Digest with HindIII OR XhoI for 60 minutes, add BglII and continue reaction for 2 hours, then heat inactivate the reaction (raise the temperature to 65 or 80°C for 20 minutes).

Perform digestion according to the enzyme manufacturer’s instructions; most researchers follow the general rule that 10 units of restriction enzyme is sufficient to overcome variability in DNA source, quantity and purity. Generally, 1 µl of enzyme is added to 1 µg of purified DNA in a final volume of 50 µl of the appropriate buffer, followed by incubation for 1 hour at the recommended temperature.

Following digestion, we recommend gel purifying the linearized vector on a 1% agarose gel to remove the fragment, and to help separate the prep from any undigested circular plasmid and to decrease the background in ligation and transformation.

Lab Note:

It is not necessary to CIP-treat the vector because the fully-digested plasmid will contain incompatible ends. However, some users who have failed to detect the presence of an insert have chosen to CIP-treat the vector – and phosphorylate the oligos – and have reported positive results. Again, however, this is only an optional step and is not necessary per se for effective ligation.

Prior to the cloning reaction in Step Three, normalize the concentration of your digested plasmid to between 0.2 and 0.5 mg/ml. If you need assistance in adjusting your concentration, see the Lab Tips section of this manual following the protocol.

>> Step Three: Ligation into pSUPER Vector

Assemble the cloning reaction by adding 2 µl of the annealed oligos to 1 µl of T4 DNA ligase buffer. Add 1 µl pSUPER vector, 5 µl nuclease-free H₂O, and 1 µl T4 DNA ligase.

Incubate overnight at room temperature. A negative control cloning reaction should be performed with the linearized vector alone and no insert.

After cloning and prior to transformation, plasmids should be treated with BglII to reduce the level of background in your transformation.

To perform this reaction:

1. Add 1.0 µl of BglII to your plasmid
2. Incubate for 30 minutes at 37°

Remember, the BglII site is destroyed upon successful cloning of the oligo pair, so those vectors cut by the enzyme will NOT contain the insert fragment.

>>Step Four: Transformation in Bacteria

Recombinant pSUPER vector should be transformed into competent cells of an appropriate host strain (e.g., DH5α) according to the supplier protocol or the transformation protocol routinely used in your laboratory. In order to monitor the efficiency of the transformation steps, as a negative control, cells should also be transformed either with a vector that has been ligated with a scrambled-base hairpin oligo, or with a circular vector containing no oligo insert.

Grow bacteria in amp-agarose plates overnight (16-24 hrs), then pick and grow colonies in an ampicilin broth for an additional cycle. Pick and miniprep several colonies (it can take many to locate a positive clone) according to supplier's instructions.

Check for the presence of positive clones (i.e., containing vector *with oligo insert*) by digesting with EcoRI and either HindIII or XhoI, depending on the configuration of your oligos as previously discussed. (Again, BglII can also be used since positive clones will not cut with BglII.) After digestion, determine your results as follows:

	Cut with EcoRI & HindIII	Cut with EcoRI & XhoI
Positive clone: vector with insert	281 bp	281 bp
Negative clone*: no insert	227 bp	248 bp

*e.g., supercoil that was nicked and not fully linearized with both enzymes) has a fragment of 227 or 248bp..

In addition, the presence of the correct insert within your recombinant pSUPER vector can be confirmed by sequencing prior to transfection in mammalian cells. For sequencing primer options, consult the appropriate pSUPER vector map included in this manual, or consult the sequence file, which is available for download from the pSUPER RNAi System section of the OligoEngine Web site (www.oligoengine.com).

>> Step Five: Transfection of Mammalian Cells

The pSUPER plasmid can now be transfected directly into your target cells. Methods for transfection include, but are not limited to, calcium phosphate (Chen and Okayama, 1987; Wigler et al., 1977), lipid-mediated (Felgner et al., 1989; Felgner and Ringold, 1989) and electroporation (Chu et al., 1987; Shigekawa and Dower, 1988). Researchers have also reported success utilizing Lipofectamine (Invitrogen) and the polyamine reagents LT1 and LT2 (from Mirus Corporation, www.genetransfer.com). Refer to these products' protocols for more specific transfection procedures. We recommend that you include the positive control vector and a mock transfection (negative control) to evaluate your results.

As an example, the procedure used by Drs. Agami and Brummelkamp and referenced in their original publication of the pSUPER system (4) notes that their target cells "...were transfected using an electroporation protocol, which results in more than 90% transfection efficiency." This protocol is in turn described in detail in a previous paper by Agami and Brummelkamp (Cell 102, 55 (2000)), as follows:

To obtain high transfection efficiency, 3 x 10⁵ MCF-7 cells were resuspended in 100 µl of electroporation buffer containing 2 mM HEPES (pH 7.2), 15 mM K₂HPO₄/KH₂PO₄, 250 mM manitol, and 1mM MgCl₂ at a final pH of 7.2. Either 1 or 2 µg of DNA was added and the cells and DNA were transferred to a 0.1 cm electroporation cuvette (BioRad) and electroporated with Gene Pulser II apparatus and Gene Pulser II RF module (BioRad) at 140 volts, 15 times 1.5 ms burst duration and 1.5 s intervals. Five minutes after electroporation, cells were seeded in either a 10 cm dish or an equivalent area. Cells were washed 16 hr after transfection and the experiment was performed either 24 or 48 hr later.

Since every cell line is different and may require a different method of transfection, some experimentation may be needed to determine the optimal conditions. We recommend that you follow exactly the protocol for your cell line (pay particular attention to medium requirements, when to pass the cells, and at what dilution to split the cells).

>> Step Six: Detection of EGFP Fluorescence

A fluorescent inverted microscope provides the easiest and most recommended method for detection of EGFP expression, and enables analysis of live cells.

If you do not have access to an inverted microscope, fluorescence can also be detected with either a photographic (35-mm) or digital camera (which frequently harbors a cooled charge-coupled device, a CCD), fitted with the appropriate filter. You can achieve excellent results using standard filter sets, such as FITC filters to detect EGFP, although optimized filter sets for detecting GFP are also available. Refer to the "Lab Tips" section of this manual following the protocol for this method of analyzing fixed cells.

>> Step Seven: Selection of Stable Transfectants (Omit if you do not have .neo or .puro vector)

The levels of siRNA expression and gene knockdown will typically vary widely among cells. In particular, transfection efficiency may be lower for primary cells; it is often difficult to obtain a stably expressing clone from normal (non-transformed) or primary cell lines using pSUPER vectors. If possible choose a transformed or immortal cell line instead.

Moreover, pSUPER-transfected cells that survive antibiotic selection may not have a significant reduction in expression of the target gene. Instead, they may have found a way to mitigate the effects of a reduction in the target gene expression by compensating in another fashion or by shutting down expression of the siRNA. Therefore, it can be useful to isolate clones that can be screened to identify the cells that cause the desired reduction in target gene expression.

Permanent cell lines may be created by growing the transfected cells under antibiotic selection. Begin by splitting the cells at 1:10-20 dilution.

Lab Note:

When selecting for positive clones, be sure to establish a kill curve for each lot of antibiotic to determine optimal effective dose. For puro selection, identify the lowest level of antibiotic that kills non-transfected cells within approximately 5 days by testing antibiotic concentrations from 1–10 µg/ml while keeping all other culture conditions equal. For neo/G418 selection, identify the lowest level that kills non-transfected cells within approximately 7 days by testing antibiotic concentrations from 25–4000 µg/ml

Note that the selection method described herein is optimized for 293 cells. For other cells, follow manufacturer's directions, or refer to the following procedure, which is provided as a reference method for stable cell selection (your standard laboratory procedure may work best for your specific conditions):

- a. Plate transfected cells at an optimized cell plating density: plating density can have a strong impact on antibiotic selection because cells at higher densities are less effectively killed off than cells at lower concentration. Also, cells that divide more rapidly typically have a lower optimal plating density than cells that double slowly.

It is important to include two non-transfected control cultures. One is subjected to antibiotic selection to control for cells that spontaneously become antibiotic resistant or are already antibiotic resistant; it will help determine the effectiveness of the transfection and selection. The second control is grown without antibiotic selection as a positive control for cell viability.

- b. After 24 hours, add culture medium containing the optimum antibiotic concentration:
 - **For .neo vectors:** Grow the cells in medium containing 500 µg/ml of neomycin or G418 (Invitrogen). Untransfected cells should die within 5-15 days.
 - **For .puro vectors:** Grow the cells in medium containing 1-10 µg/ml of puromycin. Untransfected cells should die within 5-10 days.
- c. Culture the cells in medium containing antibiotic until all of the cells in the non-transfected control culture are killed. At this point, the selection is complete and the cells can be grown without antibiotic selection.

RNAi ANALYSIS : Measurement of siRNA-induced Silencing

The level of suppression of your target gene can be measured by using different techniques:

- To determine the amount of protein expressed by the gene, a Western Blot analysis can be performed.
- For a measurement of the mRNA transcript of your target gene, Northern analysis and quantitative RT-PCR are the most widely used techniques.

Please refer to supplier protocols or standard lab methods handbooks for more information on the appropriate protocol for each technique.

Lab Tips

Diluting Oligonucleotides

To dilute your lyophilized oligos to a specified concentration, use the following equation to determine how much H₂O (or buffer, etc.) to add to your product on hand:

$$\text{ml H}_2\text{O required for concentration of Xmg/ml} = (\mu\text{g oligos} \times 10^{-3}) / \text{X}$$

Thus, if you have 200 μg of oligo*, add ~0.067 ml of H₂O to achieve a concentration of 3mg/ml [(200 $\times 10^{-3}$) / 3 = ~0.067]. When diluting your oligos, you may first wish to create a “master stock” of 10 mg/ml, which you can store and dilute further (e.g., to 3mg/ml) as needed.

You can access OligoEngine’s “Concentration calculator” online at www.oligoengine.com/calculator.html.

**If you purchased your oligos through OligoEngine, you can refer to their accompanying Data Sheets for the specific quantity, in μg , of each oligo as delivered. If you do not have this data but instead know the quantity of an oligo in pmoles, you can calculate μg weight by multiplying pmoles $\times 10^{-6} \times$ molecular weight (MW) of the oligo. Likewise, you can use a UV spectrophotometer at 260 nm to determine the optical density (“OD”) of your oligos and calculate the quantity of each in μg (1 A₂₆₀ OD = 33 $\mu\text{g/ml}$ ssDNA), then adjust your concentration accordingly.*

Normalization of Plasmid Concentration

If you are working with the circular version of the pSUPER vector and have performed the BglIII / HindIII digestion yourself, you may wish to normalize the concentration of vector after the purification process in Step 2:

- Using a UV spectrophotometer at 260 nm to determine the OD of your vector and calculate the quantity of each in μg , based on the equation 1 A₂₆₀ OD = 50 $\mu\text{g/ml}$ dsDNA.
- Using on this measurement, adjust your concentration of vector to between 0.2 and 0.5 mg/ml before proceeding to ligation (ethanol precipitation and re-dilution may be required).

Step-Cooling Suggestions for Annealing Oligos

On rare occasions, researchers have not been able to identify a positive clone after transformation. Based on extensive evaluation, we believe this is likely due to occasional problems with self-annealing oligos during the initial annealing step.

If experience such difficulties, you may wish to alter the anneal procedure to include more steps in the cooling process. Since the melting temperature of the hairpin structure will be less than that of two full-length oligos, this should help to ensure proper annealing if problems arise.

The following procedure has proven to reduce the occurrence of self-hybridizing oligonucleotides:

- Raise the initial annealing reaction temperature to 94°C for 4 min., then cool to 80°C for 4 min.
- Continue by cooling to 75°C for 4 min., then down to 70°C, before proceeding with any further cooling.

If this doesn't yield any better results, you may want to add even more steps: 94°C for 4 min, then 85°C for 4 min, then 82°C, 80°C, 78°C, 75°C, etc. (If you have access to a PCR block, you may want to use that to perform your annealing reaction for easy and automatic step-cooling.)

Standard Microscopy Procedure for Fluorescence Detection

The protocol provided below is one possible microscopy procedure for detection of the EGFP expression by pSUPER vectors. Other equally suitable and more detailed microscopy procedures may be found elsewhere (e.g., Ausubel et al., 1995 et seq.)

Materials required

- 70% Ethanol
- Dulbecco's Phosphate buffered saline (DPBS; pH 7.4)
- DPBS/4% paraformaldehyde (pH 7.4–7.6)
Add 4 g of paraformaldehyde to 80 ml of DPBS. Heat to dissolve. Once the solution has cooled, readjust the pH if necessary, then dilute to a final volume of 100 ml. Store at –20°C.
- Rubber cement, molten agarose, or commercial mounting medium (e.g., ProLong® Antifade Kit, Molecular Probes)

Procedure

In a tissue culture hood:

- a. Sterilize a glass coverslip with 70% ethanol.
- b. Place the coverslip in a sterile tissue culture dish.
- c. Plate and transfect cells in the tissue-culture dish containing the coverslip.
Note: Some cell types may not adhere to the glass coverslip. In these cases, you may need to pre-treat the glass coverslip with a substrate that promotes cell adhesion (e.g., lamin, or poly-D-lysine, or both).
- d. At the end of the culture period, remove the tissue culture media and wash once with DPBS.
- e. Fixing cells:
 - i. After cells have been washed with DPBS, add freshly made DPBS/4% paraformaldehyde directly to the coverslip.
 - ii. Incubate cells in solution at room temperature for 30 min.
 - iii. Wash cells twice with DPBS. Allow cells to soak in DPBS for 10 min during each wash.
- f. Mounting the coverslip onto a glass microscope slide:
 - i. Carefully remove the coverslip from the plate with forceps. Pay close attention to which side of the coverslip contains the cells.
 - ii. Place a tiny drop of commercial mounting solution (e.g., ProLong® Antifade Kit, Molecular Probes) on the slide, and allow the coverslip to slowly contact the solution and to lie down on the slide, cell side down.
 - iii. Carefully aspirate the excess solution around the edge of the coverslip using a Pasteur pipette connected to a vacuum pump.
 - iv. If desired, seal the coverslip to the microscope slide using molten agarose, rubber cement, or black nail polish.
 - v. Allow to dry. The drying time may vary depending on the mounting solution used.
 - vi. Examine slides by fluorescence microscopy. Once fixed, cells can be stored in the dark at 4°C.

Troubleshooting

Difficulty in achieving detectable levels of gene suppression may be due to problems that can occur at each step of this protocol. Begin by reviewing supplier guides and protocols (e.g., for transformation, transfection, etc.) to eliminate the possibility of error leading beyond the scope of this product/protocol. Once some of the more basic problems have been ruled out, consider the following:

- ***Synthesized oligo may contain incorrect sequence.*** Occasional errors in the process of DNA synthesis can cause an incorrect nucleotide to be added within an oligo. Check your data sheet to confirm that the sequence of your synthesized oligo matches what is required for proper ligation and effective target knockdown.
- ***Oligo pairs may not have annealed properly.*** Self-ligation has been known to occur in rare cases, but can be avoided by following the steps outlined in the “Lab Tip” in Step Four of the procedure.
- ***Transfection may be unsuccessful/too low.*** To confirm successful transfection of the pSUPER vector, use a positive control plasmid of equivalent size, such as a GFP vector.
- ***Mutation may have occurred.*** The process of apurination or other factors may lead to a mutation in a nucleotide of an insert oligo. A single mutation is enough to significantly affect gene knockdown. Sequence your plasmid construct using the primers described in the appropriate vector map or sequence file to confirm or rule out such problems.
- ***Target sequence may not be appropriate for silencing.*** This is one of the most common problems for lack of any observable suppression, and to date one of the least understood. Review the target selection strategy described earlier in this protocol, and choose another target sequence from different region of the gene.

Other problems may be more specific to the cell type, target gene, system or species with which you are working. You may wish to consult PubMed for recent articles in the field of RNAi that may provide some insight, or contact OligoEngine technical support (customerservice@oligoengine.com) to discuss your experiment in greater detail.



pSUPER RNAi System™

VECTOR: pSUPER.basic
CATALOG#: VEC-PBS-0001/0002

Length: 3176 bp

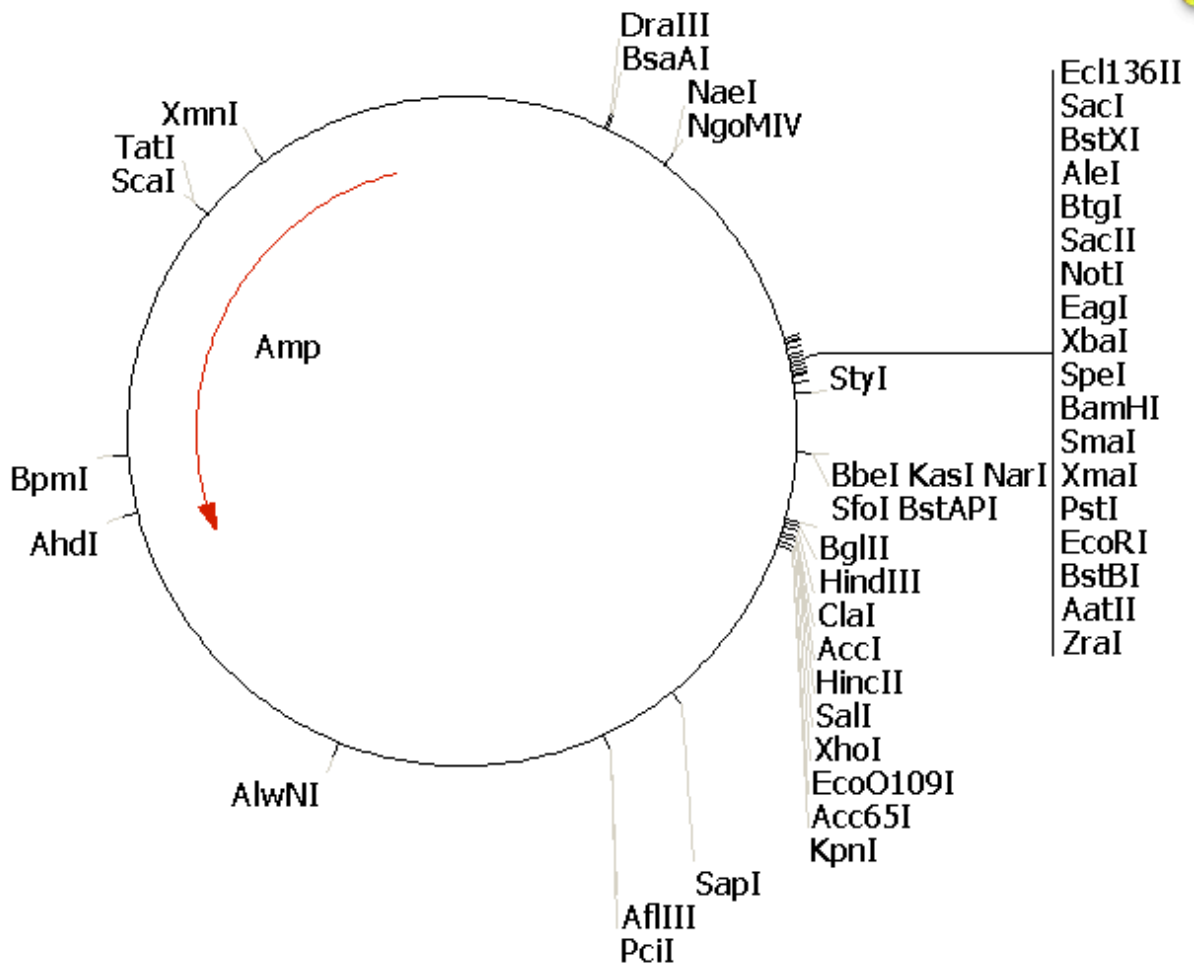
Key Sites

BglII: 928
HindIII: 934
EcoRI: 707
Sall: 949
XhoI: 955

Vector Features

f1(+) origin: 441-135
H1 promoter: 708 - 934
pUC origin: 1373-2040
Ampicillin resistance ORF: 3048-2191

T7 primer (AATACGACTCACTATAG): 627-643
M13 (-20) primer (GTAAAACGACGGCCAGT): 600-616
M13 reverse primer (AACAGCTATGACCATG): 1038-1023





pSUPER RNAi System™

VECTOR: pSUPER.neo
CATALOG#: VEC-PBS-0003/0004

Length: 4699 bp

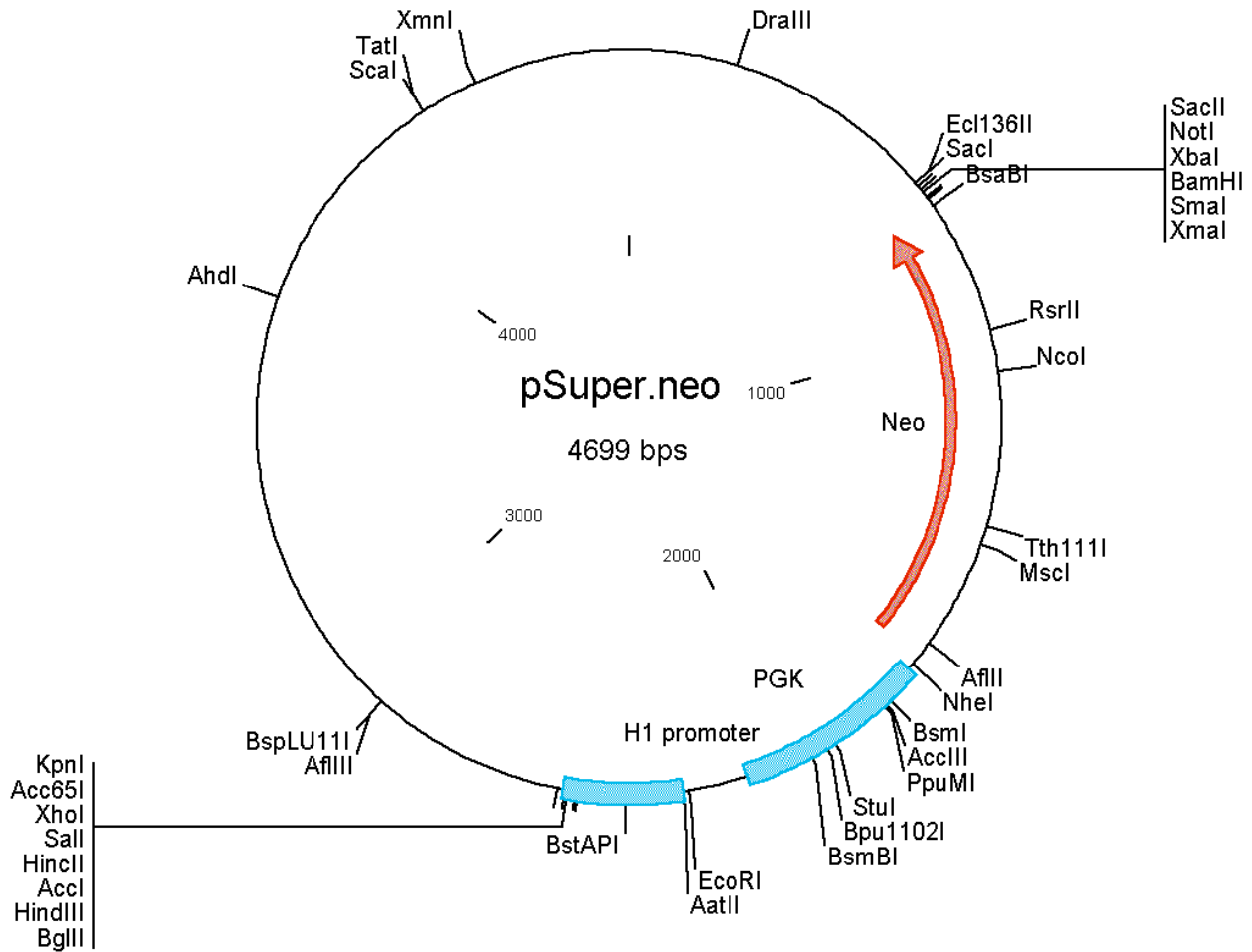
Key Sites

BglII: 2451
HindIII: 2457
EcoRI: 2230
Sall: 2472
XhoI: 2478

Vector Features

f1(+) origin: 135-441
PGK promoter: 2110-1712
Neo ORF: 1684-715
H1 promoter: 2235-2483
Ampicillin resistance ORF: 3714-4193

T7 primer (AATACGACTCACTATAG): 627-643
M13 (-20) primer (GTAAAACGACGGCCAGT): 600-616
M13 reverse primer (AACAGCTATGACCATG): 2561-2546





pSUPER RNAi System™

VECTOR: pSUPER.neo+GFP
CATALOG#: VEC-PBS-0005/0006

Length: 5429 bp

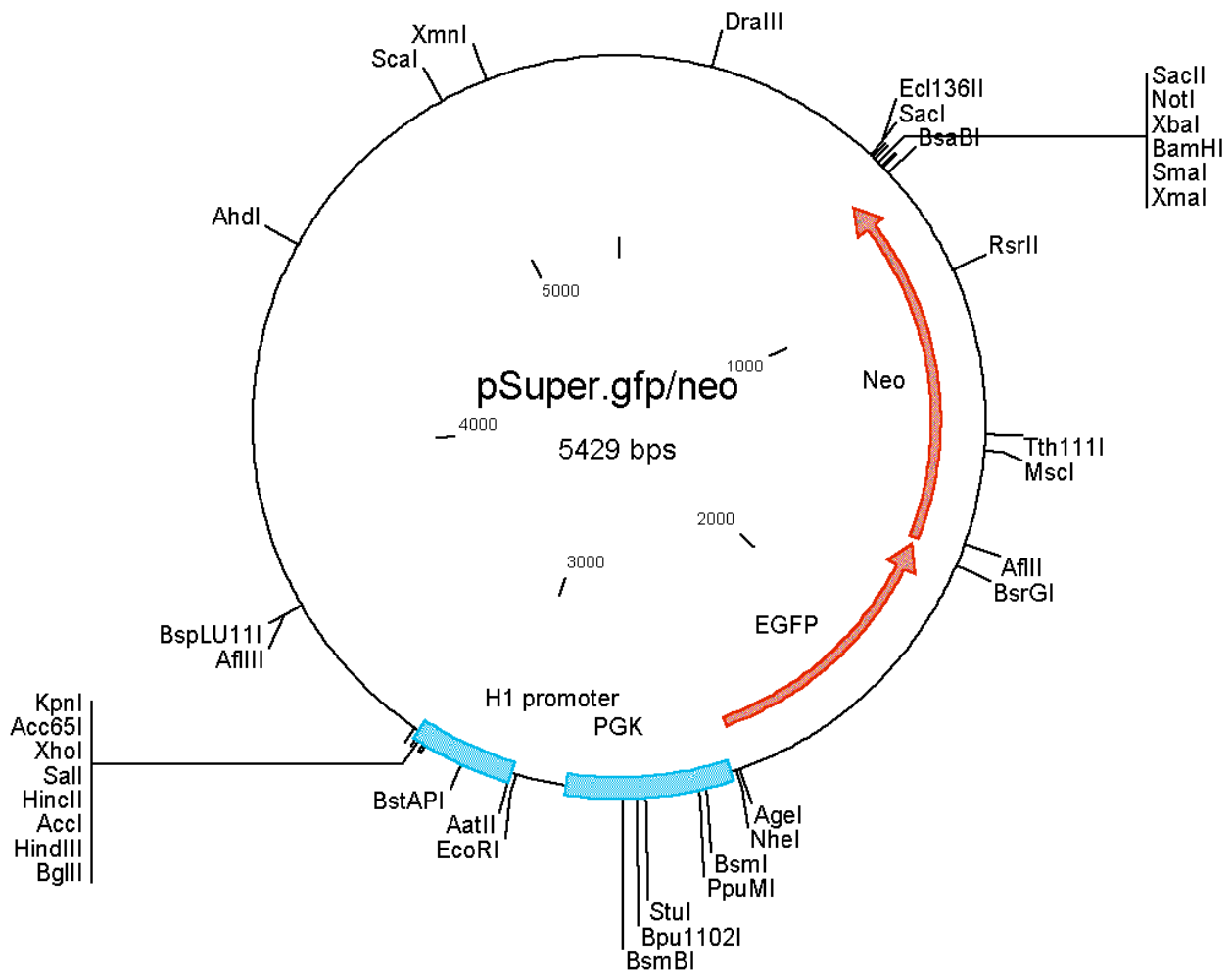
Key Sites

BglII: 3181
HindIII: 3187
EcoRI: 2960
Sall: 3202
XhoI: 3208

Vector Features

f1(+) origin: 135-441
PGK promoter: 2840-2442
Neo ORF: 1684-715
EGFP ORF: 2424-1691
H1 promoter: 2965-3213
Ampicillin resistance ORF: 5301-4444

T7 primer (AATACGACTCACTATAG): 627-643
M13 (-20) primer (GTAAAACGACGGCCAGT): 600-616
M13 reverse primer (AACAGCTATGACCATG): 3291-3276





pSUPER RNAi System™

VECTOR: pSUPER.puro
CATALOG#: VEC-PBS-0007/0008

Length: 4353 bp

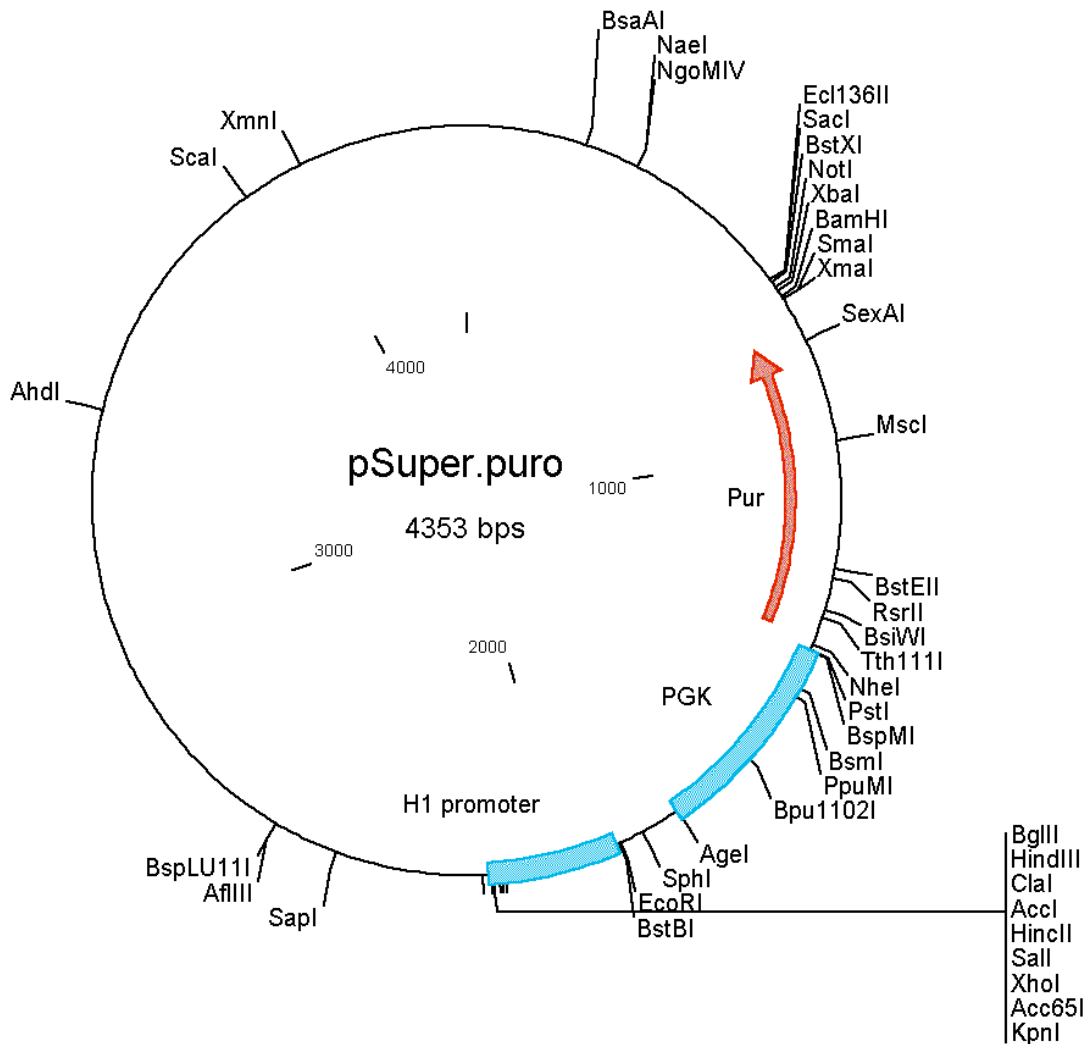
Key Sites

BglII: 2105
HindIII: 2111
EcoRI: 1884
Sall: 2126
XhoI: 2132

Vector Features

f1(+) origin: 135-441
PGK promoter: 1768-1370
Puro ORF: 1355-756
H1 promoter: 1889-2137
Ampicillin resistance ORF: 3368-4225

T7 primer (AATACGACTCACTATAG): 627-643
M13 (-20) primer (GTAAAACGACGGCCAGT): 600-616
M13 reverse primer (AACAGCTATGACCATG): 2215-2200



USER NOTIFICATION

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