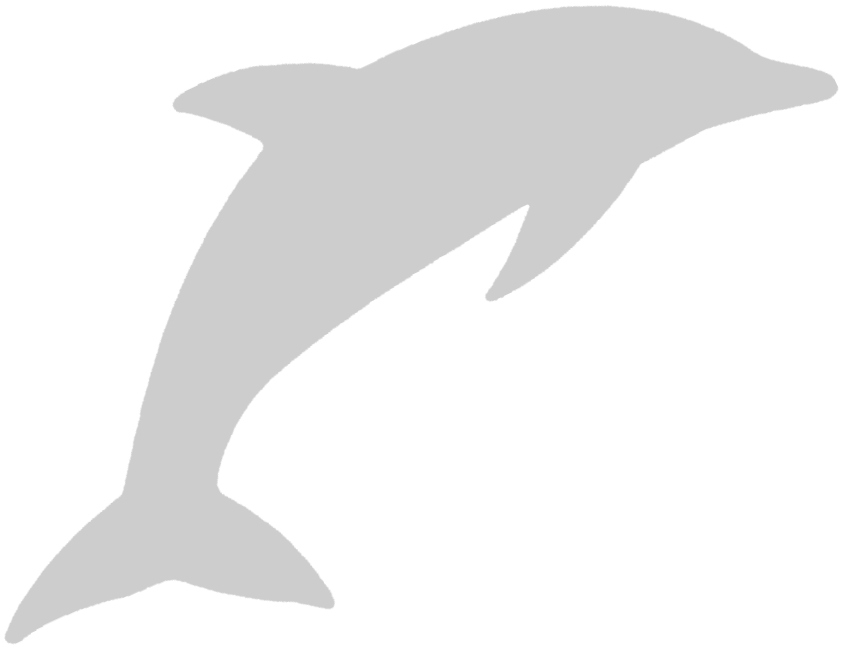


**Mo Bi Tec**

# pSKAN 8 Phagemid Vector

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Product Information and Instructions  
May 2000



**Mo Bi Tec**  
MOLECULAR BIOLOGISCHE TECHNOLOGIE

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The pSKAN Phagemid Display System has been developed by Prof. Dr. J. Collins and Dr. P. Röttgen at the GBF Braunschweig, Germany (US patent no 5,925,559).

Use of the pSKAN Phagemid Display System is covered by U.S. Patents 5,223,409, 5,403,484, 5,571,698 and 5,837,500; and EPO Patent 436,597. Any commercial use of the pSKAN Phagemid Display System, including the discovery or development of commercial products, requires a license from Dyax Corp.

For research use only. Not for use in diagnostic or therapeutic procedures.

pSKAN8 5/2000

## 1. Introduction

pSKAN8 is a phagemid vector used to prepare your own phagemid display library.

Below you find the protocol we used for constructing the library HyC, which is available as part of our pSKAN Phagemid Display System with ready-to-use libraries. This protocol can be used for the construction of your own library as well.

Note: The system is designed for the presentation of a hypervariable amino acid loop of 6 - 8 amino acids. We have no experience with inserting larger loops. Thus, this will have to be determined empirically.

## 2. Preparation of Your Own DNA Library with pSKAN8

### 2.1. Amplification of DNA Oligos

#### 2.1.1. Amplification Primers

primer F (# 4403)     5' GCTACAACGAGCTCAACGGTTGC 3'

primer B (# 4404)     5' GGGTAAGTGTCCACCGTCGGTACCGCA 3'

Note: These primers are **not available** at **MoBiTec**.

#### 2.1.2. Protocol

In a 0.5 ml reaction tube for PCR\* (polymerase chain reaction) fill:

- 1 pmole of the hypermutated oligo
- 40 pmole of primer F (# 4403)
- 40 pmole of primer B (# 4404)
- 10  $\mu$ l DNA-polymerase buffer (+ MgCl<sub>2</sub>) (10x)
- 5  $\mu$ l MgCl<sub>2</sub> (25 mM)
- 10  $\mu$ l dNTP solution (2 mM, each)
- 0.25  $\mu$ l of Taq-DNA polymerase (5 U/ $\mu$ l)

\*PCR is a process covered by patents owned by Hoffmann-La Roche. Use of this process requires a license.



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Add H<sub>2</sub>O to a final volume of 100 µl. If necessary, cover the solution with 50 µl mineral oil to prevent evaporation and place the reaction tubes in a PCR thermocycler. Carry out 3 to 5 subsequent cycles of 1 minute at 94°C (DNA denaturation), 2 minutes at 50°C (primer annealing) and 3 minutes at 72°C (DNA synthesis). We highly recommend the Quick Step PCR Purification Kit from Edge Bio Systems (distributed by **MoBiTec** in Germany). Alternatively, purify the DNA by phenol/chloroform extraction and precipitate with ethanol. Dissolve in 20µl TE buffer.

Note: Also available at **MoBiTec** in Germany are thermostable polymerases and tubes for thermocyclers, which can be used without mineral oil (see **MoBiTec** catalog chapter 7).

## 2.2. Modification of DNA

### 2.2.1. Restriction Reactions

for *Sst* I (= *Sac* I) and *Kpn* I restrictions:

PCR product	16 µl
<i>Kpn</i> I, <i>Sac</i> I	20 units
restriction buffer (10x)	3 µl
H <sub>2</sub> O ad	30 µl

Mix and incubate the tubes at 37°C for at least 2 hours and check the restriction products by polyacrylamide gel electrophoresis (ca.20%). Purify the DNA by phenol/chloroform extraction and ethanol precipitation. In order to be able to work with optimal buffer conditions, the two restriction reactions may also be performed in two steps. For a convenient buffer exchange inbetween the reactions, MobiSpin columns are available from **MoBiTec**.

### 2.2.2. Ligation of DNA Fragments

To ligate cleaved DNA fragments mix in an Eppendorf reaction tube:

- vector and insert DNA fragments at a molar ratio of 1 : 4
- 1:10 of end volume of ligation buffer (10x)
- 0.5 units ligase per µg DNA

add H<sub>2</sub>O to reach an end volume according to a DNA concentration of 0.01 µg/µl.

Mix and incubate overnight at 8°C or 3 - 4 hours at room temperature<sup>1</sup>. Add 1 volume of H<sub>2</sub>O and inactivate the enzyme by heating to 65°C for 10 minutes. Purify the DNA by phenol/chloroform extraction and ethanol precipitation and redissolve it in 10 µl of TE buffer.

## 2.3. Preparation of Competent Cells

Inoculate 10 ml of LB/Tc medium with a single colony of *E. coli* WK6λmutS<sup>2</sup> and incubate at 37°C and 180 rpm overnight.

The next day, inoculate 1000 ml of LB/Tc medium (2 x 500 ml Erlenmeyer flasks) at 1% with the overnight grown culture and incubate again at same conditions until an optical density of OD<sub>600</sub> = 0.6 has been reached.

Transfer 250 ml aliquots of the culture into centrifuge tubes (GS3), chill them on ice and centrifuge for 15 minutes at 6,000 rpm and 4°C (Sorvall RC5C centrifuge; GS3 rotor). Resuspend each pellet in 250 ml of ice-cold H<sub>2</sub>O and repeat the centrifugation step. Resuspend each pellet in 125 ml of ice-cold H<sub>2</sub>O, pour together two aliquots and centrifuge again. Resuspend each pellet in 10 ml of ice-cold glycerol (10%) collect both aliquots in an GSA centrifuge tube and centrifuge for 15 minutes at 8,000 rpm. Finally resuspend the bacterial pellet in 1 ml of glycerol (10%).

Fill 50 µl aliquots in precooled, sterile Eppendorf reaction tubes, freeze immediately in liquid nitrogen and store at -70°C until the transformation by electroporation.

<sup>1</sup> Ligation at 8°C is efficient, but slow.

<sup>2</sup> *E. coli* WK6λmutS

Genotype: *galE*, *strA*, *nalI*, Δ(*lac-proAB*), λ<sup>+</sup>, *mutS*::Tn10, F'*[lac<sup>q</sup>, lacZΔM15, proAB<sup>+</sup>*]

## 2.4. Transformation of *E. coli* Cells by Electroporation

Place frozen aliquots of competent *E. coli* WK6λmutS cells on ice and let them thaw. To each aliquot add 1 µl of ligated DNA (0.01 - 0.1 µg) and incubate on ice for 1 minute. Fill the suspension in a precooled electroporation cuvette (use a Pasteur pipette), place the cuvette in the electroporation sled and give a pulse at a voltage of 2.5 kV, a capacity of 25 µF and a resistance of 200 Ω (Gene Pulser and Puls Controller, Bio-Rad). Immediately add 1 ml of LB medium, mix with a



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Pasteur pipette and transfer the suspension in an Eppendorf reaction tube. Incubate for 1 hour at 37°C and plate on LB agar (1 ml/135 mm plate) containing ampicillin (100 µg/ml) and tetracycline (20 µg/ml). Incubate overnight at 37°C. In the same way carry out a transformation with and without pSKAN-DNA as a control and plate out on LB/Tc and LB/Amp/Tc plates. Also diluted aliquots of transformed cells should be plated in order to calculate the size of the final library.

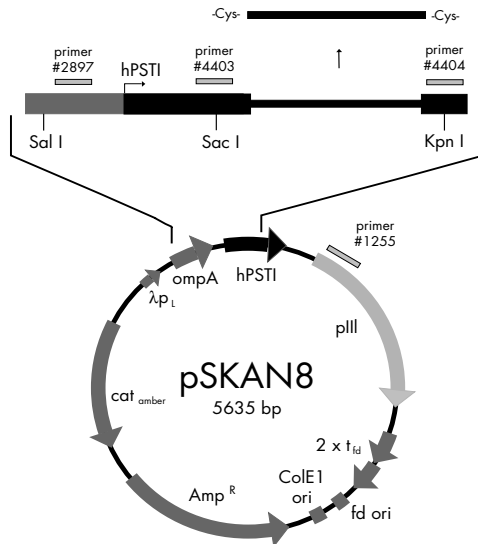
As a control, the individual clones of the library can be sequenced using the following primers:

primer # 1255      5' GGGATTTGCTAAACAAC 3'

primer # 2897      5' GGAGGTCTAGATAACGAGG 3'

Double stranded sequencing is recommended. One primer may be sufficient.

### 3. Vector Map



#### Phagemid vector construction.

pSKAN8 is based on the vector pMAMPF-3. Between the two Cys a hypervariable amino acid loop (with (NNG/T)<sub>6-8</sub>) can be created using the protocol in chapter 2. (See also the ready-to-use pSKAN libraries.)

The phagemid contains the genetic elements:

- Amp<sup>R</sup>: selection marker (ampicillin resistance)
- ColE1 ori: plasmid origin of replication
- fd ori: origin of replication for the fd phage
- λp<sub>L</sub>: λ-promoter left
- ompA: signal sequence for secretion
- hPSTI: gene of the human pancreatic secretory trypsin inhibitor
- pIII: gene for the phage coat protein III
- 2 x t<sub>fd</sub>: two phage fd transcription terminator sequences
- cat<sub>amber</sub>: chloramphenicol resistance is inactive due to an amber mutation

## 4. Primer and Restriction Site Positions in pSKAN8

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<Sal I > _____ primer #2897
5' AAGTCGACCAAGGAGGTCTAGATAACGAGGGCAAAAAATGAAAAAGACAGCTATCGCGATTTCGGTACCTGGCTGTTTCGGCTACCCGTAGCGCAGGCC3'
(bp #5536)
      ↑
      hpSTI
      →
5' GAC TCT CTG GGT GAA GCT AAA TGC TAC AAC GAG CTC AAC GGT TGC ACT AAG ATC TAC GAC CCG GTT TGC
asp ser leu gly arg glu ala lys cys tyr asn glu leu asn gly cys thr lys ile tyr asp pro val cys
(bp #1)
      (primer #4403)
      ↑
      <Sac I >
      →
      variable region
      ↓
<Kpn I >
GGT ACC GAC GGC AAC ACT TAC CCG AAC GAA TGC GTT CTG TGC TTC GAA AAC CGT AAA CGT CAG ACT TCT ATC
gly thr asp gly asn thr tyr pro asn glu cys val leu cys phe glu asn arg lys arg gln thr ser ile
(primer #4404)
      ↓
      hpSTI
      ←
      - linker -
CTG ATC CAG AAA TCT GGT CCG TGC TTA ATT CAT GAA GAA GGT GAA TTC TCA GAA GCG CGC GAA GAT ATC AGA
leu ile gln lys ser gly pro cys leu ile his glu glu gly glu phe ser glu ala arg glu asp ile arg
      ↑
      pIII
      →
      primer #1255
GCT GAA ACT GTT GAA AGT TGT TTA GCA AAA TCC CAT ACA GAA AAT TCA TTT ACT AAC GTC TGG AAA GAC GAC3'
ala glu thr val glu ser cys leu ala lys ser his thr glu asn ser phe thr asn val trp lys asp asp
      (bp #288)

```

The sequence of pSKAN8 in this overview starts with the last bases of the linear sequence (bp # 5536 to # 5635). In the circular plasmid bp # 1 follows bp # 5635.



## 5. Literature

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## 6. Order Information, Shipping & Storage

order #	description	amount
<b>shipped at room temperature; store at +4°C:</b>		
pSKAN8	pSKAN8, lyophilized vector DNA	5µg
PSKAN4	pMAMPF-3-PSTI4, lyophilized vector DNA	5µg
PS1255	sequencing primer # 1255 (18 mer) lyophilized	500 pmol
PS2897	sequencing primer # 2897 (19 mer) lyophilized	500 pmol
PSKAN2	anti-hPSTI antibody, monoclonal mouse lyophilized	40 µg
PSKAN3	anti-pIII (g3p) antibody, monoclonal mouse lyophilized	100 µl
<b>shipped on dry ice; store at -70°C:</b>		
PSKAN	pSKAN Phagemid Display System HyA, HyB, HyC pSKAN libraries inclusive <i>E. coli</i> host strain <i>WK6λmutS</i>	3 x 1ml 1ml