

2004

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<http://www.healthsystem.virginia.edu/internet/biomolec/>**

Page of
Total Number of
Samples:

DNA SEQUENCING REQUEST FORM

Date: _____ P.I.: _____ User: _____ E-Mail: _____ (Please print)
Department: _____ Rm #/Bldg: _____ PTAO#: _____
Phone #: _____ FAX #: _____ Cancer Center Member? New PTAO?
Data delivery desired: (Please check one): E-mail Mac-to-Mac ABI File Attachment Zip Disk (Please bring it along with the form)

Smp No.	Log # 04-official use	Name of Sample (<= 8 letters!!!)	Name of Vector	Host Strain	Method of Purification	Type of Template*	Primer Name	Primer Length	From Bank ^a	Insert Site	Insert Length	Comment Specify if BACs, Lambdas etc or high GC content or repeats
# 1						S D P			Y / N			
# 2						S D P			Y / N			
# 3						S D P			Y / N			
# 4						S D P			Y / N			
# 5						S D P			Y / N			
# 6						S D P			Y / N			
# 7						S D P			Y / N			
# 8						S D P			Y / N			
# 9						S D P			Y / N			
# 10						S D P			Y / N			

Note: * Circle one: S -- single stranded; D -- double stranded or plasmid; P -- PCR product.
+ A -- Qiagen kit;; B -- Promega kit; C -- alkaline/PEG, Others -- please specify.
Facility use only. The log-in number will serve as a unique identifier for all files related with this particular sample.

**For a complete list of the available primers please visit: <http://www.people.virginia.edu/~dnaseq/primers.html>
Please submit samples with primer and template premixed. The amount of primer required is 8.0 pmole. The amount of template required: dsDNA : 500-1000 ng; ssDNA: 400-800 ng; PCR: 20 ng/100 bp. Samples should be in dH₂O qs to 14 ul. If you are requesting primers from our bank, please submit the template in the required amount in 10 ul of dH₂O and mark the primer requested on the tube. All tubes should be labeled, preferably on the top, with the sample numbers that appear in the first column.**