



DBT-Supported Genomic Facility at UDSC, CIIDRET
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w w.e.f 1st Jan 2022
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Order form for DNA sequencing by primer walking(Service code D7) For Details read technical information) (Please fill out one form for each DNA-Primer combination)

Name and Postal address

Phone (with code)..... **Mobile No****Email**.....
Payment Receipt in name of..... **Signature**.....

Payment: Rs-----**Transaction No./UTR No.** with date -----
Name of the Bank-----

Service code

D	7
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Sample ID

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Type of sample (Plasmid/PCR product)----- Host -----
 Plasmid backbone (pUC, pBR, pBluescript, pGEM etc.)----- Antibiotic marker -----
 Size of plasmid (kb)----- Size of insert / PCR product (kb/bp) -----
 Concentration of DNA/PCR product----- Cloning sites for insert -----
 Expected GC Content (%)----- Expected Homopolymer (Yes/No) -----

Forward primers for ends (T7, T3/SP6/M13R/M13F) -----
Specific Primer (if not provided by facility) name, length, sequence -----
 Primer sequence 5' to 3' (18-23 mer)

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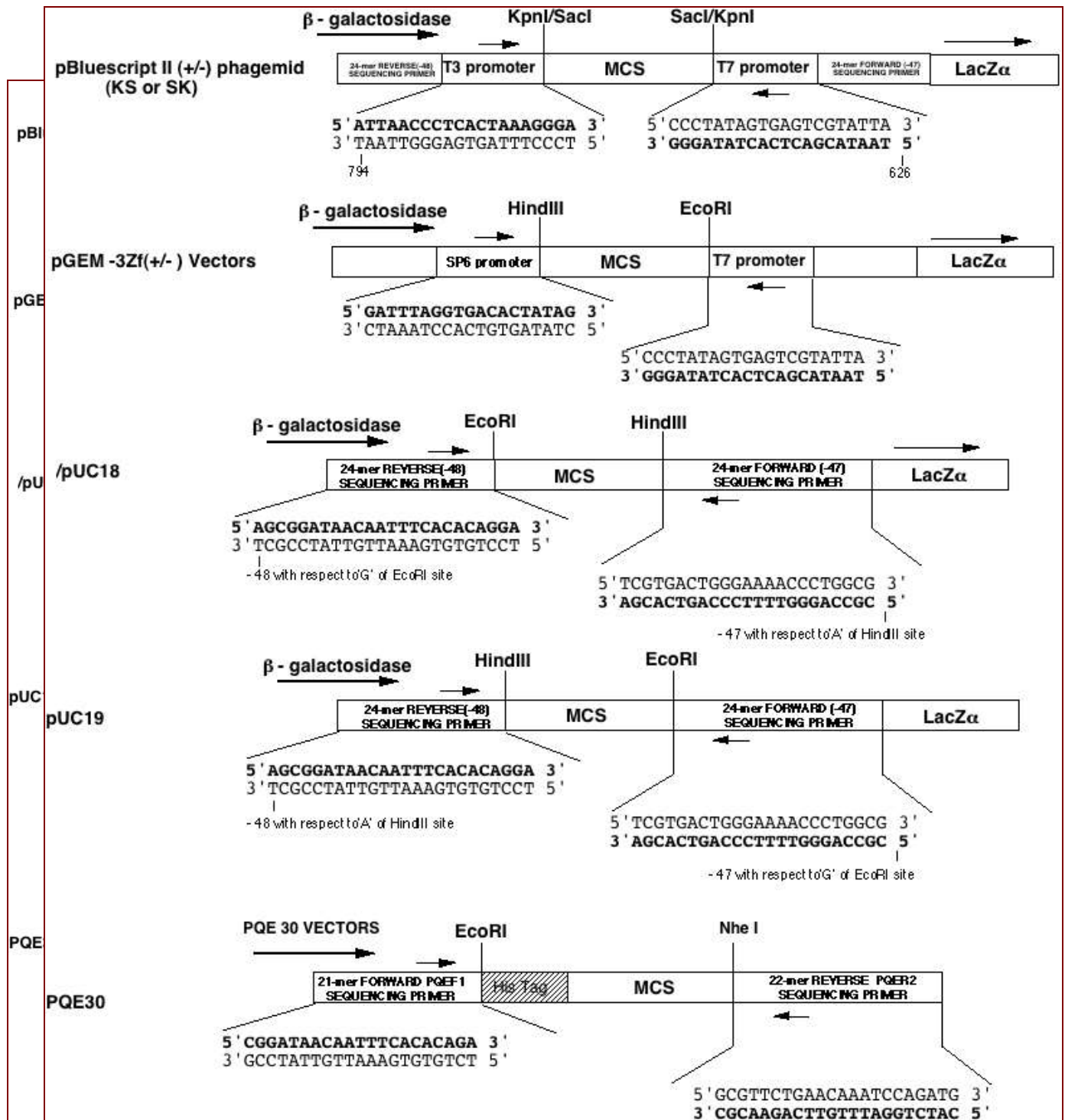
 Annealing temperature of specific primer -----
 conc.of specific primer-----pmole/ul
 Concentration of specific primer-----pmol/ μ l.,
 Purification grade (OPC/HAP/HPLC/Desalted) -----

Reverse primers for ends (T7, T3/SP6/M13R/M13F) -----
Specific Primer (if not provided by facility) name, length, sequence -----
 Primer sequence 5' to 3' (18-23 mer)

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 Annealing temperature of specific primer -----
 conc.of specific primer-----pmole/ul
 Concentration of specific primer-----pmol/ μ l.,
 Purification grade (OPC/HAP/HPLC/Desalted) -----

For Office use
 Date of receipt..... Date of result.....
 Comments.....



Primer sequences are shown in bold letters

T7TN TERMINATER PRIMER 20 MER NEB	: 5' GTCGGTTGAGTCGAAGGAAA 3'
T7 PROMOTER PRIMER 20MER=NEB1248	: 5' TAATACGACTCACTATAGGG 3'
T3 PROMOTER PRIMER 20MER=NEB1228	: 5' ATTAACCCCTCACTAAAGGGA 3'
SP6 PROMOTER PRIMER 19MER=PROMEGA Q5011:	5' GATTTAGGTGACACTATAG 3'
M13 REVERSE(-48) PRIMER 24MER=NEB1233	: 5' AGCGGATAACAATTTACACAGGA 3'
M13 FORWARD(-47) PRIMER 24MER=NEB1224	: 5' CGCCAGGGTTTTCCCAGTCACGAC 3'
PQE.F1 (FORWARD PRIMER 21MER)=QIAGEN	: 5' CGGATAACAATTTACACAGA 3'
PQE.R2 (REVERSE PRIMER 22MER)=QIAGEN	: 5' CGCAAGACTTGTTTAGGTCTAC 3'