

CUSTOMER REQUISITION FORM (GN)

OFFICE USE ONLY

ID No:

Date:

Received by:

Time:

PERSONAL DETAILS

Name:

Designation:

PI/Research Guide:

PI/RG Designation:

Department:

Institute/ University with address:

Phone No:

Email ID:

SERVICE DETAILS

Study Title:

Service Availed/Requested:

Sample Name/s:

Sample Type & number of samples:

Species/Strain/cell line:

No of Cells (in RNA Later):

Storage Temperature:

References (if any):

Transportation Temp:

SERVICE AVAILING

<ul style="list-style-type: none"> <input type="checkbox"/> Expression Analysis (Real Time PCR based) <ul style="list-style-type: none"> <input type="checkbox"/> Absolute quantification <input type="checkbox"/> Syber Green <input type="checkbox"/> Relative quantification <input type="checkbox"/> Taqman probe based <input type="checkbox"/> Microarray Based Analysis <input type="checkbox"/> Chromosomal Microarray (array CGH (8X60k format)) <ul style="list-style-type: none"> <input type="checkbox"/> Microarray based Expression analysis <input type="checkbox"/> Microarray based SNP analysis <input type="checkbox"/> Microarray based CHIP on Chip <input type="checkbox"/> NGS services <ul style="list-style-type: none"> <input type="checkbox"/> Microbial sequencing <input type="checkbox"/> Whole Genome Resequencing <input type="checkbox"/> Mitochondrial sequencing <input type="checkbox"/> Whole transcriptome analysis <input type="checkbox"/> Amplicon Sequencing <input type="checkbox"/> Small RNA analysis <input type="checkbox"/> RNA sequencing <input type="checkbox"/> MicroRNA analysis <input type="checkbox"/> Exome sequencing <input type="checkbox"/> Custom Gene Synthesis <input type="checkbox"/> Oligo Purification <ul style="list-style-type: none"> <input type="checkbox"/> PAGE purification <input type="checkbox"/> HPLC purification <input type="checkbox"/> Affinity Column purification 	<ul style="list-style-type: none"> <input type="checkbox"/> DNA Cloning Service <ul style="list-style-type: none"> <input type="checkbox"/> Cloning in standard Vector and confirmation <input type="checkbox"/> Cloning in Vector for Eukaryotic (fungal) system <input type="checkbox"/> Cloning in expression vector and confirmation <input type="checkbox"/> Cloning in Vector for Eukaryotic (mammalian) system <input type="checkbox"/> Cloning in Vector for bacterial system <input type="checkbox"/> Identification and Phylogenetic services <ul style="list-style-type: none"> <input type="checkbox"/> Bacterial identification services(16s RNA sequencing) <input type="checkbox"/> Animal identification services (Based on COI sequencing) <input type="checkbox"/> Yeast identification services (Based on ITS2 region) <input type="checkbox"/> Algae identification service (Based on 18s RNA sequencing) <input type="checkbox"/> Fungi identification services (Based on ITS1&2 region) <input type="checkbox"/> Primer synthesis service <ul style="list-style-type: none"> <input type="checkbox"/> Oligo 10nM 11-40mer <input type="checkbox"/> Oligo 20nM 11-40mer <input type="checkbox"/> Oligo 20nM 41-60mer <input type="checkbox"/> Oligo 20nM 61-80mer
<ul style="list-style-type: none"> <input type="checkbox"/> DNA sequencing services <ul style="list-style-type: none"> <input type="checkbox"/> Single pass sequencing upto 1KB <input type="checkbox"/> Purification of PCR products before sequencing <input type="checkbox"/> Plasmid isolation from stab/culture plates 	<ul style="list-style-type: none"> <input type="checkbox"/> Cell Line Authentication <ul style="list-style-type: none"> <input type="checkbox"/> Human Cell Line Authentication Services <input type="checkbox"/> Pharmacogenomics Services <ul style="list-style-type: none"> <input type="checkbox"/> DMET Services <input type="checkbox"/> Protein Services <ul style="list-style-type: none"> <input type="checkbox"/> Peptide Synthesis Services <input type="checkbox"/> Conjugations Services <input type="checkbox"/> N-terminal Sequencing

REMARKS

****If sample would be found damaged on receiving, then sample won't be processed ahead and would be intimated.
The concentration mentioned would be considered final throughout the experiment.
Methodology suggested would be executed if feasible otherwise best alternative method would be done on your consent.
Once the experiment is initiated, it cannot be terminated but minor feasible changes would be accepted.**

Kindly send your sample on the below mentioned address:

GeneXplore Diagnostics and Research Centre Pvt. Ltd.,
1201-1210, 12th Floor, Iconic Shyamal, Shyamal Cross Road,
Satellite, Ahmedabad, Gujarat, India-380015
Phone: +91 79-26584044/55, Mobile: +91-9824999084
E-mail: academic@genexplore.co.in; research@genexplore.co.in,
Web: www.genexplore.co.in

Signature of Research Student/RG/PI