

Form associated with:

Service Project 6551018 at Genomics Core

Owner: Matthew Brown (Brown, Matthew (MSU) Lab)

LIMS Form

*** Sample Information**

	Sample Name	Primer Name	DNA Type	Primer Added	Anneal At 60	Fragment Length	High GC or Secondary Structure
1	GvSwiss	Gv5SSU	PCR	no	no	>800 bp	
2	GvSwiss	MedlinB	PCR	no	no	>800 bp	
3	GvSwiss	RoscGutt45F	PCR	no	no	>800 bp	
4	GvSwiss	RoscGutt571F	PCR	no	no	>800 bp	
5	GvSwiss	RoscGutt629R	PCR	no	no	>800 bp	
6	GvSwiss	RoscGutt1321F	PCR	no	no	>800 bp	
7	GvSwiss	RoscGutt1456R	PCR	no	no	>800 bp	
8							
9							
10							
11							

IMPORTANT NOTES:

Our standard annealing temperature is 60 C. Please let us know if your samples need to be annealed at 55 C or 65 C instead!

TO AVOID DELAYS IN SAMPLE PROCESSING:

1. Make sure you have an account on our [LIMS site](#). Registration is only required once.
2. Complete all fields in the table above. Leave none blank.
3. Only use the following characters in the Sample Information table:
 - Uppercase letters
 - Lowercase letters
 - Numbers 1-9
 - Hyphen -
4. Enter your total number of samples in the Sequencing Charges question below.
5. Click the "Add Selected Services" icon after entering your sample quantity so a valid charge is added to your request.

*** Service Type**

- ☒ User Prep - DNA or PCR Products ready for sequencing
- ☐ Clean and Load - BigDye cycled reactions only
- ☐ PCR Purification and Sequence - PCR products for Purification prior to Sequencing

*** Sequencing Charges**

- Clean and Load Sample
- Post-PCR Clean and Sequence, 1-48 samples
- Post-PCR Clean and Sequence, 49+ samples

Purified DNA for Sanger Sequencing, 1-48 samples

Purified DNA for Sanger Sequencing, 49+ samples

Purified DNA for Sanger Sequencing, per plate