Modifying existing sequencing job information

This is applicable to jobs other than bacterial metagenomics only. Bacterial metagenomics projects cannot be altered after submission.

If you do not have a bacterial metagenomics project:

The Return to Job Request link allows a user to edit information they provided about a sequencing job request. This can be used only after the user has created a sequencing request and has been provided a request ID. Modifications cannot be done after a sequencing request has been approved and accepted by the GSAF.

1. Enter the request ID associated with your sequencing job request and press ‘submit’.

2. A view with the price quote for the current job request will be displayed with buttons to modify your main job information and sample information. If you haven’t submitted your job to the GSAF, then you’ll also see submit and delete buttons.

3. If you would like to modify any of the fields, click “Edit” to open the page for editing your job information. Change the values in the fields which you wish to edit. Once you are done editing your information, click “Update”.

4. After Update you will be taken back again to the Price Quote page.

For descriptions of the different job-related fields, see Glossary of terms Return to Job Request to do this now.