



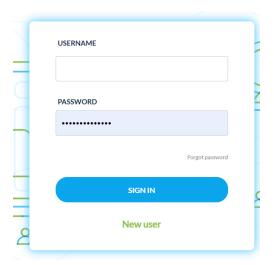
Sanger Sequencing Order Management in BookitLab

Introduction

This manual provides a step-by-step guide on how to <u>place Sanger sequencing orders</u> on the BookItLab system and how to <u>view results</u>.

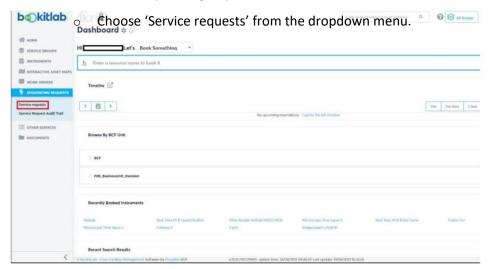
Placing a New Sequencing Order

1. **Log In**: Enter the BookltLab system using your username and password.



2. Navigate to Sequencing Requests:

o Go to 'Sequencing requests' on the left-hand side.







3. Create a New Order:

o Click the '+' sign on the right-hand side to initiate a new order.



o Select the order type (Plasmid / PCR / BAC) from the dropdown menu.



Click 'Select'.

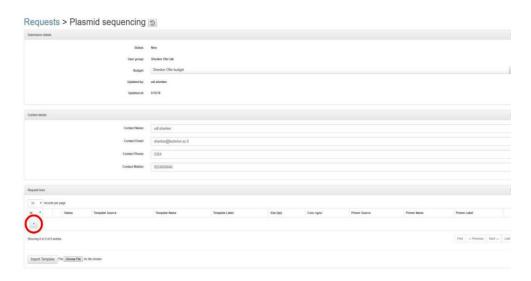






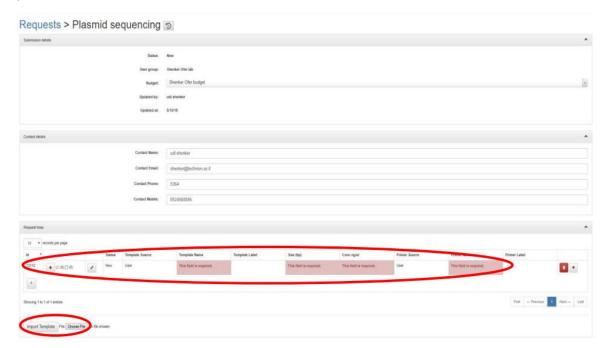
4. Fill-in Order Details:

o Add a row by clicking the '+' symbol on the left side of the table under 'Request lines'.



 Enter details such as template name, size, concentration, and primer name. Avoid using spaces or special characters; all fields are mandatory.

For large sample orders, download the CSV template by clicking 'Import template', complete it, and upload it back.

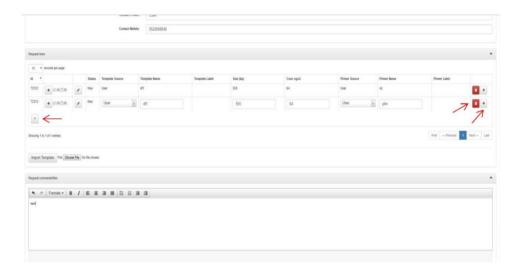






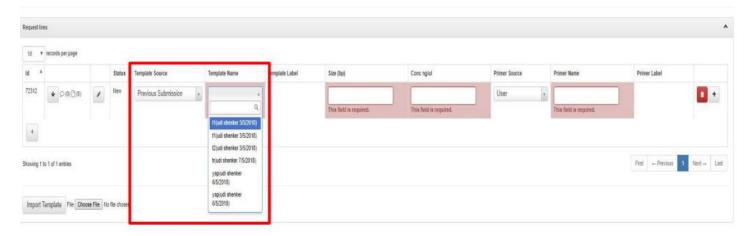
5. Add or Edit Rows:

- To duplicate a row, click the '+' symbol on the right of a filled line.
- For a blank row, click the '+' on the left.
- To delete a row, click the 'trash' symbol.



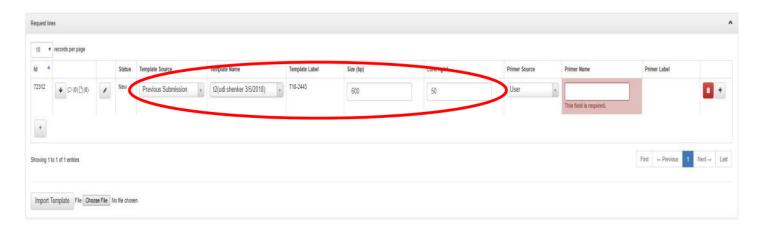
6. Use Previous Samples or Primers:

- Select 'Previous Submission' in the 'Template/Primer source' dropdown.
- o Choose the sample from the 'Template/Primer name' list based on the submission date.
- Once you choose a previously submitted sample the 'Template name', 'Size' and 'Concentration' fields will be filled automatically.
- Note: Samples/primers are stored for 6 months only. Ensure the label appears; if not, contact the ATGC team.



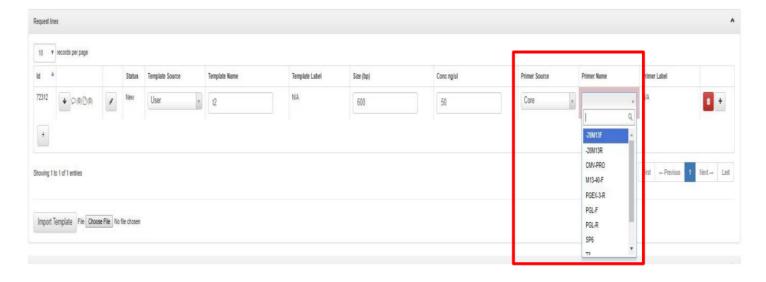






7. Core Primers:

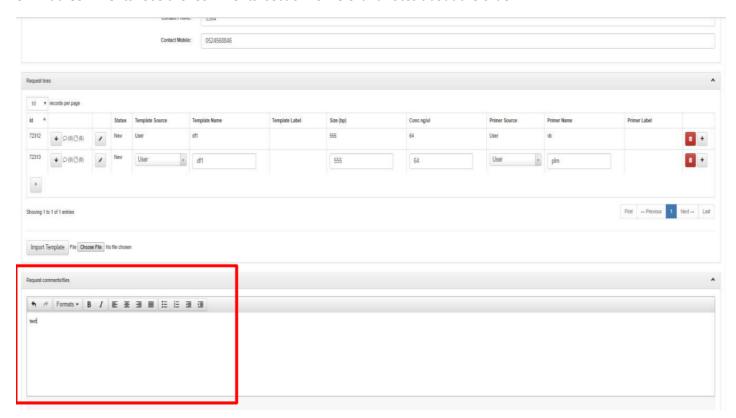
- o Choose 'Core' under 'Primer source' and select the primer from the 'Primer name' dropdown.
- Labels will not be printed for core primers.



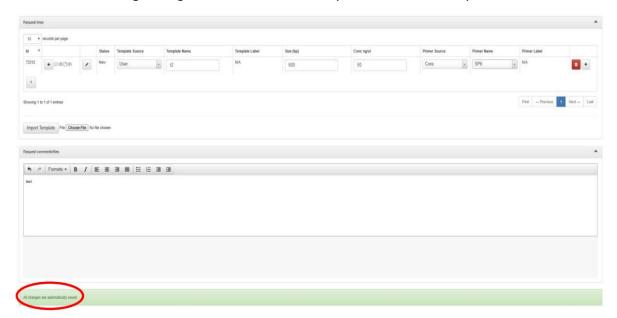




8. Add Comments: Use the 'Comments' section for relevant notes about the order.



9. **Automatic Saving**: Changes are saved automatically; no need to manually save.

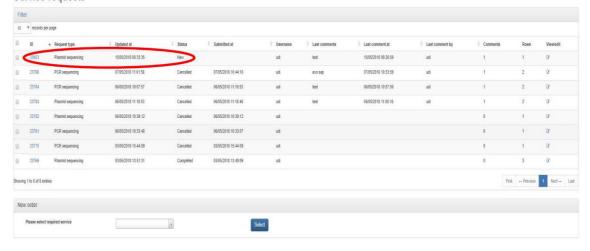






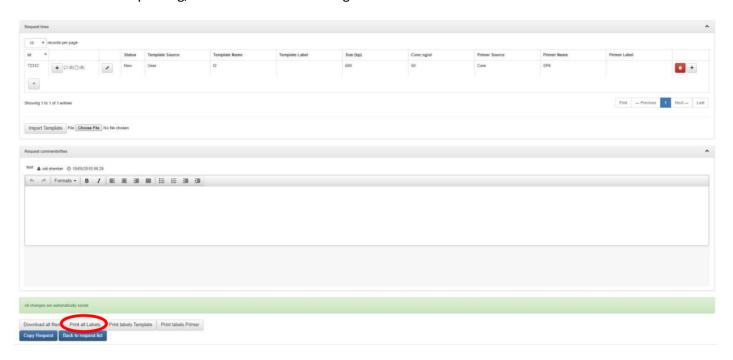
Your order will now be visible in the main table under 'Sequencing requests'. The order status will be 'New'
until labels are printed.

Service requests



10. Finalize Order:

- o Print labels using the BCF computer connected to the label printer.
- o Log in to BookItLab, select your order, and click 'Print all labels' at the bottom of the page.
- o **Important**: Use Google Chrome. Do not highlight lines when printing, as this may cause errors.
- After printing, the order status will change to 'Started'.



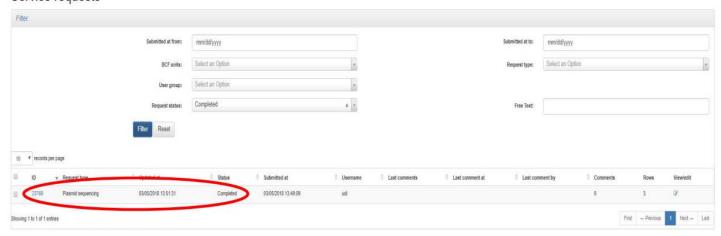




Viewing Results

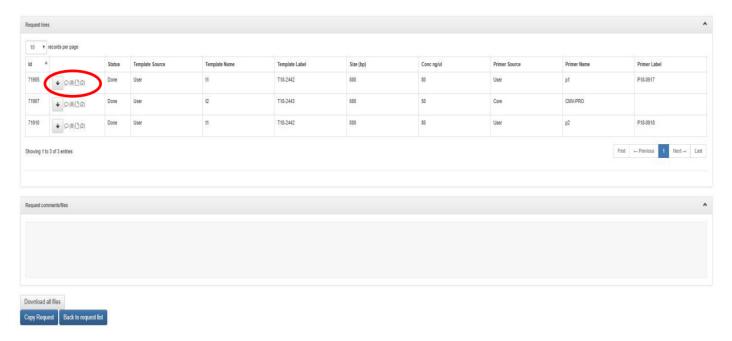
1. Check Order Status: Once the reaction is complete, the order status changes to 'Completed'.

Service requests



2. Access Results:

- Open the order by clicking the order ID.
- Results will be visible on the left side of each row.

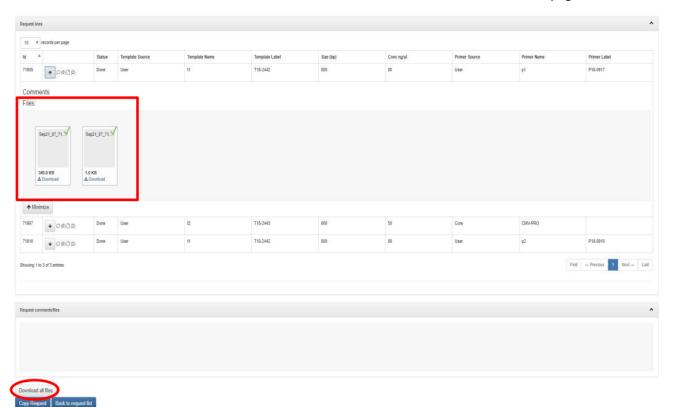






3. Download Results:

- o Download individual files by clicking 'Download' at the bottom of each row.
- o To download all files at once, click 'Download all files' at the bottom of the page.



4. View Results:

- For efficient data analysis, we recommend downloading the free Chromas software from <u>Technelysium's website</u>.
- o After installation, import your `.ab1` files into Chromas to view and analyze the electropherograms.
- For a visual guide on using Chromas, you might find the following tutorial helpful: <u>Sequence Analysis</u>
 Protocol Open and View Sequence Files Chromas PC.

Notes and Tips

- Labels can only be printed using Google Chrome.
- Ensure all fields are completed accurately to avoid delays.
- Contact the ATGC team if any issues arise with labels or previous sample usage.