

User Guide Sanger Sequencing Fragment Analysis

Ver_2.0



How to use "Copy" function



[6/6] Filtered Row: 6	6 Filtered Row: 6		
No.	Sample Name *	Primer*	
1	1	785F Copy	2
2	2	Copy entire rows Q	2
3	3	Copy entire rows below Q	2
4	4	Q	2
5	5	Q	۲
6	6	Q	٦.

① Copy

- When: Want to fill certain (selected) rows with a single value
- **How :** Select the rows you want to fill with the single value. One row should be filled with the value you want to copy. Right-click and select "Copy". The value fills the selected rows.



① Copy entire rows

- When: Want to fill entire rows with a single value
- **How**: Select the row with the value you want to copy, right-click and select "Copy entire rows". The value fills the entire rows.

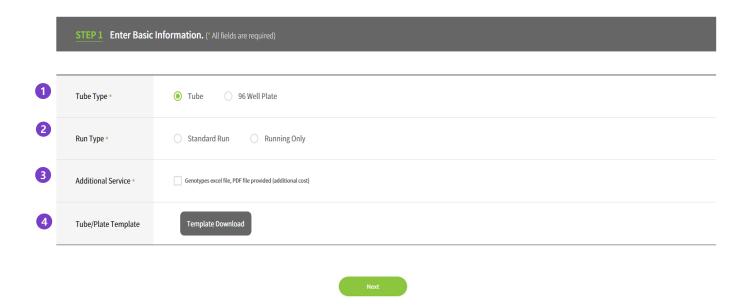


Copy entire rows below

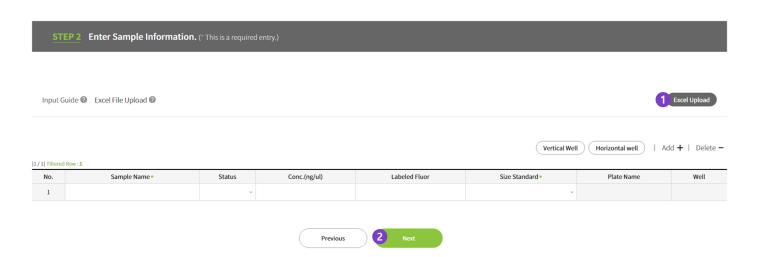
- When: Want to fill certain (here to the end) rows with a single value
- **How**: Select the row with the value you want to copy, right-click and select "Copy entire rows below". The value fills the below rows.

Fragment Analysis





- 1 Tube Type: Choose tube type of your samples.
- 2 Run Type: Size marker necessary to be included choose standard Run, unnecessary to be included choose Running Only
- 3 Additional Service : You don't have gene mapper program, we will provide PDF, excel file with additional charge
- 4 Tube / Plate Template: You can download and fill out the order sheet



- ① Excel Upload: Upload your order sheet that you filled out
- 2 Next: Click the next for next step
- * You may download your order sheet from first page, or you may fill out the information directly at step 2

Fragment Analysis



STEP 5 Enter Billing information. (* Required field)		
BLAST Service *	Yes No This is a service that searches for a sequence that is highly similar to the sequence of DNA obtained in The Basic Local Alignment Search Tool and NCBI database. We will provide you with data files retrieved from NCBI and no analysis services will be provided.	
Storage Period *	Immediate Disposal 1 Month 3 Month If immediate disposal is selected, additional orders and re-sequencing cannot be ordered as the sample is immediately discarded after the primary experiment.	
Ordered by * @	Gildong Hong	
Actual Orderer Contact	01012341234	
Primary E-mail	moonkun86@macrogen.com	
Additional E-mail for Result *	moonkun86@macrogen.com	
Comment		
Institution for Billing	Macrogen Korea Find	
Mobile Number *	01012341234	
Country *	Korea (the Republic of)	
	254, Beotkkot-ro, Geumcheon-gu, Seoul, Republic of Korea	
	11th floor, 1111	
Billing Address *	Geumcheon-gu	
	Seoul 08511	
PO Number		
ATTN	Macrogen	
Coupon	Choose ~ Apply	
Promotion Code	Please Enter. Promotion Code Apply	
Request on payment		
	Previous Save 1 Submit	

① Input all the information for billing and select the option

for "Blast service" and "Storage Period" and click [Submit]

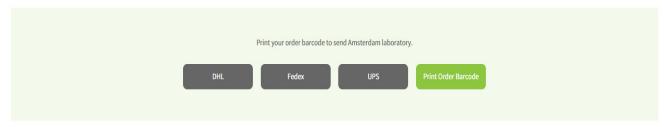
Fragment Analysis



Your order has been placed successfully.

Order Number | EC00000595

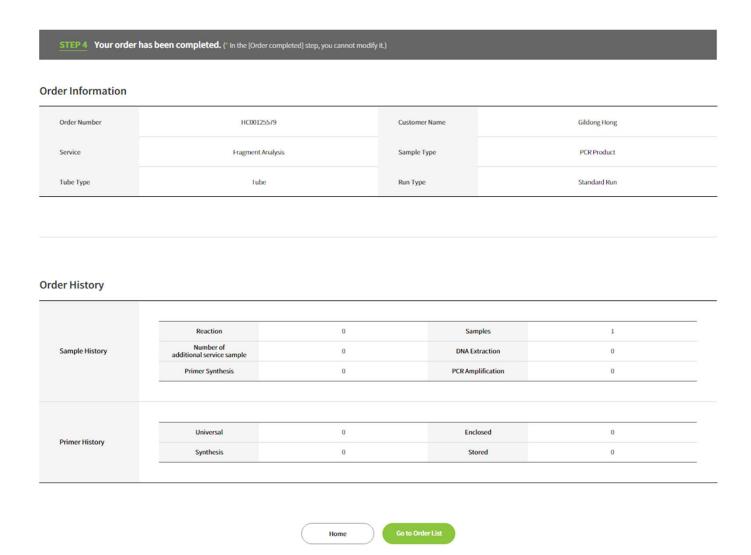
 $Please\ double\ check\ your\ order\ information\ then\ send\ your\ samples\ accordingly.$



^{*} Click DHL/FedEx/UPS Invoice button to download shipping documents.

* You can check the order number and get your commercial [invoice for DHL / Fedex/ UPS shipping].

Please [print order barcode] and send it with the samples.



^{*} Lastly, you can check the order information

^{*} Print all shipping documents and call DHL/FedEx/UPS to pick up your samples.

^{*} You will be notified of the status of your order by email.

^{*} For other inquiries, please use [Support > Contact us].