## Placing an order on UC Stratocore as CCHMC or external users

1. Once your account has been approved, log in to Stratocore using your account information and press "Order" at the top of the screen. From there, you may enter the services you are requesting for your samples.

samplest	Incidents Rights Training Projects Orders Settings Users Accou	unts Adm	ns Invoicing He	elp	
You may create a project for your order – however, this is for <b>your</b> <b>record only</b> . Creating a project is not placing an order. You may have multiple orders within a project but should only have one order per group of samples.	Order service or consumable Make this order on behalf of this user: Teke Ellen I Teke Ellen Attach this order to this project: Ino project selected I Teke Request/start a new project Select a financial account for the order: D600051-626120000-1-GESC D6 Dev Services and consumables available for order (select and enter the quantity	y values)			
	Lab service	Min M	ax Incr. Unit price	Quantity	Price
Select the Lab service you would like us	#090127 Amplicon-seq/ChIP-seq/DNA-seq (per Sample) add a comment	1	- 1 1+:40.00	0	0.00
to provide and enter the quantity	#090126 DNA/RNA/Library QC (per Sample) show description add a comment	1	- 1 1+:25.00	0	0.00
(number of samples). We will add the	#090128 Exome/Methylation Sequencing service (per Sample) add a comment	1	· 1 1+:250.00	0	0.00
Pass-through (reagents, consumables,	#090133 Library Only Sequencing service (per Sample) add a comment	1	- 1 1+:25.00	0	0.00
	#090129 Methylation EPIC Bead Chip Assay Service add a comment	1	- 1 1+:70.00	0	0.00
etc.) later to finalize the <b>total price</b> ,	#090130 RNA/DNA Isolation (per Sample) add a comment	1	- 1 1+:40.00	0	0.00
which will match the one we provided	#090198 Sequencing servicemiRNA/Small RNA-seq (per Sample) add a comment	1	- 1 1+:120.00	0	0.00
to you	#090132 Sequencing serviceRNA-seq (per Sample) add a comment	1	- 1 1+:120.00 - 1 1+:120.00	0	0.00
to you.	#200100			0	
	#090199 Sequencing serviceSingle Cell Sequencing (per Sample) add a comment	1		0	0.00
Press " <b>Save quote</b> " ( <mark>not Order</mark> ) once	#090131 Special RNA/DNA Isolation 2 (per Sample) add a comment	1	- 1 1+:25.00	0	0.00
	#090131 Special RNA/DNA isolation 2 (per Sample) add a comment   #090134 TaqMan assay (CNV, SNR qPCR, etc) show description add a comment	1		0	0.00
· Press " <b>Save quote</b> " ( <mark>not Order</mark> ) once	#090131 Special RNA/DNA Isolation 2 (per Sample) add a comment	1 1 1	- 1 1+:25.00	· · ·	
· Press " <mark>Save quote</mark> " ( <mark>not Order</mark> ) once you have entered all of the	#090131 Special RNA/DNA isolation 2 (per Sample) add a comment   #090134 TaqMan assay (CNV, SNR qPCR, etc) show description add a comment   Pass-through Pass-through show description add a comment		- 1 1+:25.00	· · ·	

2. A PO number is **required** for invoicing. Therefore, after you save the quote (do not press the Order button) we will update the services to finalize the **total price** including the pass-through. Then we will email you with the **total price** for you to submit to your purchasing office for a PO number.

3. Please email us the PO number as well as a sample information file (please fill out the excel file titled **Manifest file template** which can be downloaded at <u>https://med.uc.edu/depart/eh/cores/genomics/services-and-order</u>).

4. We will update your quote with the PO number and create an order from it. After this is completed, please bring your samples to the Core.