

Clustering Sheet for Flowcell

22KLGVLT3

Name Bartlomiej Gebarski	Sequencer NovaSeqX1B	Run Type NovaSeqX 10B XP flowcell PairedEnd 150 - single lane (r1: 150, r2: 150)
Clustering Date 2024-05-21	Sequencing Date 2024-05-21	Status Started
1st Index Length 20	2nd Index Length 15	
Comments no comment	Reagents Checked Out <input type="checkbox"/>	

num	Id	Scientist	Custom	PR	Comments
1	293535_R17174_1		<input type="checkbox"/>	Standard	100pM 5% PhiX - 26.69µl of buffer (10mM Tris); - 6.46µl of 2.5nM library; - 0.85µl of 1nM PhiX; CRISPR_Screen
2	293628_R17179_1		<input type="checkbox"/>	Standard	- 20.84µl of buffer (10mM Tris); - 11.63µl of 2.5nM library; - 1.53µl of 1nM PhiX; WGS
3	293629_R17179_2		<input type="checkbox"/>	Standard	- 20.84µl of buffer (10mM Tris); - 11.63µl of 2.5nM library; - 1.53µl of 1nM PhiX; WGS
4	293630_R17179_3		<input type="checkbox"/>	Standard	- 20.84µl of buffer (10mM Tris); - 11.63µl of 2.5nM library; - 1.53µl of 1nM PhiX; WGS
5	293632_R17179_5		<input type="checkbox"/>	Standard	- 20.84µl of buffer (10mM Tris); - 11.63µl of 2.5nM library; - 1.53µl of 1nM PhiX; WGS
6	293633_R17179_6			Standard	120pM 1% PhiX - 23.88µl of buffer (10mM Tris); - 8.08µl of 2.5nM library; - 2.04µl of 0.1nM PhiX RNAseq
7	M18553_R17180_1		<input type="checkbox"/>	Standard	120pM 1% PhiX - 23.88µl of buffer (10mM Tris); - 8.08µl of 2.5nM library; - 2.04µl of 0.1nM PhiX [various]
	⌕M18560_R17187_1				Solexa-Control
	⌕M18571_R17197_1				Solexa-Control
	⌕M18583_R17209_1				Solexa-Control
	⌕293927_R17194_1				DNA amplicon sequencing
8	M18562_R17189_1		<input type="checkbox"/>	Standard	120pM 1% PhiX - 23.88µl of buffer (10mM Tris); - 8.08µl of 2.5nM library; - 2.04µl of 0.1nM PhiX ChIP-Seq
	⌕M18574_R17200_1				Solexa-Control
	⌕M18584_R17210_1				Solexa-Control
	⌕M18585_R17211_1				Solexa-Control

Instructions

1. Check, if instrument is idle. Check free disc space, delete old runs if necessary.
2. Thaw the reagents in water for 4h or in a 2°C to 8°C refrigerator for 48 hours. Inspect the position #12 well on the underside of the cartridge to make sure that the contents are free of ice, which indicates that the reagents are thawed. If the reagent cartridge cannot be loaded into the instrument within 24 hours, store at 2°C to 8°C for up to 72 hours or return to -25°C to -15°C storage for up to 7 days. After thawing, do not refreeze more than one time.
3. Prepare Cluster Sheet, Samples and all other reagents.
4. Take out the flowcell from the fridge to equilibrate it to room temperature (5-10min).
5. Thaw Lyo-insert and Pre-load-buffer at room temperature for 10 minutes.
6. Denature samples in Eppendorf tubes:
 - **21.35 µl** of buffer (10mM Tris);
 - **10.1 µl** of **2.5nM** library
 - **2.55 µl** of **0.1nM** PhiX
 - Add **8.5µl** of **0.2N** NaOH to each sample;
 - Incubate for 5 minutes;
 - Neutralize by adding **127.5µl** of **Pre-load Buffer**;
7. Uncap the library tube strip. Do not pierce the library tube strip foils.
8. Dispense **165µl** denatured library or denatured library with PhiX into each sample tube.
9. After dispensing libraries, cap the library tube strip. Make sure that no air gaps are present at the bottom of the tubes (if yes, spin down briefly).
10. Insert the library tube strip into the reagent cartridge and press down.
11. Insert the lyo insert into the reagent cartridge and press down.
12. Log in to the software and start new run.
13. Follow instructions on the screen.