



CIB Genomics Core Lab  
Blue Pippin size selection

Date Received: \_\_\_\_\_

Date \_\_\_\_\_  
 Project Name \_\_\_\_\_  
 Contact \_\_\_\_\_  
 Department \_\_\_\_\_  
 UMC \_\_\_\_\_  
 OCC # \_\_\_\_\_  
 E-mail \_\_\_\_\_  
 Phone \_\_\_\_\_ Fax \_\_\_\_\_

**Ninglin Yin, Genomics Group Leader**  
**Center for Integrated BioSystems**  
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 (435)797-8218 or (435)760-3688  
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 e-mail: [usudna@aggiemail.usu.edu](mailto:usudna@aggiemail.usu.edu)  
 Web site: <http://www.biosystems.usu.edu>  
**Billing Questions: Sue McCormick (435) 797-8780**

**Please sign/date acknowledgement below:**  
*While the CIB strives to maintain high quality results, if results are not found to be acceptable, investigator will still be billed for all runs and reruns of samples. Lab fees may be negotiated.*

Principal Investigator:  
 Name (print): \_\_\_\_\_

(Sign): \_\_\_\_\_ Date: \_\_\_\_\_

**Important points regarding samples**

- The maximum volume that can be loaded is 30 µl. Samples delivered in volumes larger than 30 µl will be divided between multiple lanes and charged accordingly.
- Samples should be in a low salt buffer with any DNA-binding proteins/enzymes removed. Please clean up enzyme digests, PCR products, etc. first.

Please check box to have the samples analyzed on the TapeStation after size selection. Additional charge is \$5 per sample plus \$5 per TapeStation run.

**Charge is \$15/sample.**

							<b>Lab Use Only</b>		
	Sample Name	Type of DNA (PCR product, seq library, etc.)	buffer	Conc. (ng/µl)	Total Volume	Size range desired			
1									
2									
3									
4									
5									
6									
7									
8									

\_\_\_\_\_ **Estimated Price for Project**