

The CWRU Genomics Core Facility

Microsatellite/Genotyping Request Form

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Date: _____	Your Name: _____
Primary Investigator: _____	Email: _____
Department: _____	Account Number: _____
Phone Number: _____	Affiliation: <input type="checkbox"/> CWRU <input type="checkbox"/> CCCC <input type="checkbox"/> CCF <input type="checkbox"/> OTHER
Project Title: _____	Funding: <input type="checkbox"/> NIH <input type="checkbox"/> NSF <input type="checkbox"/> CTSC <input type="checkbox"/> OTHER

Sample Submission:

- Samples should be brought to the Core post-PCR for analysis in 96-well plate format.
- Clearly label the plate with your name, date, and project title.
- **Email an excel file with sample name, well location, and size standard to the Core.**
- The Core is capable of analyzing dye sets 6-FAM, HEX, NED, ROX, VIC, and PET using a LIZ500-bp or LIZ600-bp standard. We also run SNaPshot reactions using a LIZ120-bp standard.

#	Sample Name	Size Standard	Dye set	Application kit (Mouse marker, SNAPshot)
1				
2				
3				
4				
5				
6				
7				
8				
9				
10				
11				
12				
13				
14				
15				
16				

Genotyping Service: # reactions _____ @ \$5.00/rxn Full plate (96 or more) # reactions _____ @ \$3.00/rxn	Already prepared reactions: # reactions _____ @ \$1.00/rxn Full plate (96 or more) # plates _____ @ \$48/plate	Total:
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