

Sample Submission Form for SNP Microarrays

(Please print and attach to submitted samples)



- ❖ Submission date: _____
- ❖ Name of submitter: _____
- ❖ Name of PI: _____
- ❖ Affiliation (if outside WIS): _____
- ❖ Phone number/s: _____
- ❖ e-mail: _____
- ❖ **UserID on SusanC/Noys:** _____
- ❖ Order Number: _____
- ❖ Experiment title: _____

- ❖ DNA extraction method: _____

❖ **For every submitted sample please fill out:**

For internal use		Sample name (as on tube)	Sex (M/F)	Amount submitted (µg)	Concent. (ng /µl)	Volume submitted (µl)	260/280 ratio	260/230 ratio
	1							
	2							
	3							
	4							
	5							
	6							
	7							
	8							

	9							
	10							
	11							
	12							

❖ **Type of array/s needed** (check relevant box/es): **(Provided by WIS / us)**

- Mapping 10K 2.0 Array
- Mapping 50K Xba 240 Array
- Mapping 50K Hind 240 Array
- Mapping 250K Nsp Array
- Mapping 250K Sty Array
- Mapping 500K Array
- _____

❖ The aim of the experiment is:

- Copy number analysis. Suspected region is: _____
- Genotyping / Haplotype determination

❖ If you are interested in homozygous regions shared by several samples (for example individuals sharing a phenotype), please specify the samples:

Group 1: _____

Group 2: _____

❖ Please provide a family tree, if applicable (important for QC of results).

❖ **Any other comment, or anything you think we should be aware of?**
