

Sample Submission Form for Affymetrix Microarrays

(Please print, fill out ALL fields and attach to submitted samples)



- ❖ Submission date: _____
- ❖ Name of submitter: _____
- ❖ Name of PI: _____
- ❖ Institute: _____
- ❖ Phone number/s: _____
- ❖ e-mail: _____
- ❖ UserID on SusanC/Noys: _____
- ❖ **(For non-WIS users): Service Agreement^(*) valid until:** _____
- ❖ Order Number: _____
- ❖ RNA/DNA extraction method: _____
- ❖ For every submitted sample please specify (start with the controls):

For internal use	#	Sample name (as on tube)	Amount submitted (ng)	Concent. (ng / μ l)	Volume submitted (μ l)	260/280 ratio	260/230 ratio
	1						
	2						
	3						
	4						
	5						
	6						
	7						
	8						
	9						
	10						
	11						
	12						

- ❖ OD measured using: Nanodrop / Qbit / Other (specify): _____

❖ **Array type** (check relevant box):

- | | |
|--|---|
| <input type="checkbox"/> Human Gene 2.0 ST | <input type="checkbox"/> Arabidopsis ATH1 Genome |
| <input type="checkbox"/> Human Gene 1.0 ST | <input type="checkbox"/> Arabidopsis Tiling |
| <input type="checkbox"/> Human PrimeView | <input type="checkbox"/> Barley Genome |
| <input type="checkbox"/> Human Exon 1.0 ST | <input type="checkbox"/> <i>C. elegans</i> Genome |
| <input type="checkbox"/> Human Promoter 1.0R | <input type="checkbox"/> <i>C. elegans</i> Tiling |
|
 | <input type="checkbox"/> Chicken Genome |
| <input type="checkbox"/> Mouse Gene 2.0 ST | <input type="checkbox"/> Drosophila Genome 2.0 |
| <input type="checkbox"/> Mouse Gene 1.0 ST | <input type="checkbox"/> Citrus Genome |
| <input type="checkbox"/> Mouse Exon 1.0 ST | <input type="checkbox"/> Drosophila Genome |
| <input type="checkbox"/> Mouse Promoter 1.0R | <input type="checkbox"/> Drosophila Tiling |
|
 | <input type="checkbox"/> <i>E. coli</i> Genome 2.0 |
| <input type="checkbox"/> Rat Gene 1.0 ST | <input type="checkbox"/> Maize Genome |
| <input type="checkbox"/> Rat Exon 1.0 ST | <input type="checkbox"/> Tomato Genome |
|
 | <input type="checkbox"/> <i>V. Vinifera</i> |
| <input type="checkbox"/> Human SNP 250K | <input type="checkbox"/> <i>Xenopus laevis</i> Genome |
| <input type="checkbox"/> Human SNP 6.0 | <input type="checkbox"/> Yeast Genome 2.0 |
|
 | <input type="checkbox"/> <i>S. cerevisiae</i> Tiling |
| <input type="checkbox"/> miRNA | <input type="checkbox"/> <i>S. pombe</i> Tiling |
| <input type="checkbox"/> miRNA 2.0 | <input type="checkbox"/> Zebrafish Genome |
| |
 |
| | <input type="checkbox"/> Other: _____ |

❖ **Microarray data analysis will be carried out by:**

- Weizmann Inst. Bioinfo Unit
 Submitter
 Other (provide name, telephone and email):

❖ **Any other issues we should be aware of:**

(*) Details concerning Service Agreement: www.weizmann.ac.il/Biological_Services/