

**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<sup>(\*)</sup> valid until:** \_\_\_\_\_
- ❖ Order Number: \_\_\_\_\_
- ❖ RNA extraction method: \_\_\_\_\_
- ❖ For every submitted sample please specify: (start with the controls)

<b>For internal use</b>	<b>#</b>	<b>Sample name (as on tube)</b>	<b>Amount submitted (ng)</b>	<b>Concent. (ng /<math>\mu</math>l)</b>	<b>Volume submitted (<math>\mu</math>l)</b>	<b>260/280 ratio</b>	<b>260/230 ratio</b>
	1						
	2						
	3						
	4						
	5						
	6						
	7						
	8						
	9						
	10						

- ❖ Measurements were done using: Nanodrop / Other (specify): \_\_\_\_\_

❖ **Type of array/s needed** (check relevant box/es):

- |  |   |
|--|---|
| <input type="checkbox"/> Human Genome U133A 2.0          | <input type="checkbox"/> Arabidopsis ATH1 Genome      |
| <input type="checkbox"/> Human Genome U133 plus 2.0      | <input type="checkbox"/> Arabidopsis Tiling           |
| <input type="checkbox"/> Human Gene 1.0 ST               | <input type="checkbox"/> <i>C. elegans</i> Genome     |
| <input type="checkbox"/> Human Exon 1.0 ST               | <input type="checkbox"/> <i>C. elegans</i> Tiling     |
| <input type="checkbox"/> Human Promoter 1.0R             | <input type="checkbox"/> Chicken Genome               |
| <input type="checkbox"/> Human Tiling 1.0R Set (PM/MM)   | <input type="checkbox"/> Drosophila Genome 2.0        |
| <input type="checkbox"/> Human Tiling 2.0R Set (PM only) | <input type="checkbox"/> Citrus Genome                |
| <input type="checkbox"/> Human Tiling 2.0R Array _____   | <input type="checkbox"/> Drosophila Genome            |
| <br>   | <input type="checkbox"/> Drosophila Tiling            |
| <input type="checkbox"/> Mouse Genome 430A 2.0           | <input type="checkbox"/> <i>E. coli</i> Genome 2.0    |
| <input type="checkbox"/> Mouse Genome 430 2.0            | <input type="checkbox"/> Maize Genome                 |
| <input type="checkbox"/> Mouse Gene 1.0 ST               | <input type="checkbox"/> Tomato Genome                |
| <input type="checkbox"/> Mouse Exon 1.0 ST               | <input type="checkbox"/> <i>V. Vinifera</i>           |
| <input type="checkbox"/> Mouse Promoter 1.0R             | <input type="checkbox"/> <i>Xenopus laevis</i> Genome |
| <input type="checkbox"/> Mouse Tiling 1.1R Set (PM/MM)   | <input type="checkbox"/> Yeast Genome 2.0             |
| <input type="checkbox"/> Mouse Tiling 2.0R Set (PM only) | <input type="checkbox"/> <i>S. cerevisiae</i> Tiling  |
| <input type="checkbox"/> Mouse Tiling 2.0R Array _____   | <input type="checkbox"/> <i>S. pombe</i> Tiling       |
| <br>   | <input type="checkbox"/> Zebrafish Genome             |
| <input type="checkbox"/> Rat Genome 230 2.0              | <br>  |
| <input type="checkbox"/> Rat Gene 1.0 ST                 | <input type="checkbox"/> Other: _____                 |
| <input type="checkbox"/> Rat Exon 1.0 ST                 |   |

❖ **Arrays are provided by:**

- User  
 Weizmann Inst. MicroArray Unit  
 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/](http://www.weizmann.ac.il/Biological_Services/)