

# Lab: Preprocessing and quality control

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For this lab, you will use the `estrogen` package, the `simpleaffy` package and the `affyPLM` package.

## Exercise 1.

```
> library(estrogen)
> openVignette("estrogen")
```

This contains exercises for Affymetrix preprocessing, rudimentary quality control, up to the production of a gene list. Working through these exercises should take you 20-60 min, depending on your experience with R and the `affy` package. If you have brought your own data, process it in a similar way.

**Exercise 2.** Load the `simpleaffy` package and repeat the QA procedures there. First use the `qc` function to compute the necessary values and then try out some of the quality measures. Are there any unusual arrays?

**Exercise 3.** Load the `affyPLM` package and compute RLE and NUSE. Plot these and again determine if there are any unusual arrays. Do the results from this exercise agree with those from Exercise 2?