Solutions for chapter Fold-Changes, Log-Ratios, Background Correction, Shrinkage Estimation, and Variance Stabilization

Exercise 1

c > hist(log2(r1/g1), breaks=33, col="azure")

> hist(log2(r2/g2), breaks=33, col="azure")

Exercise 2

For the kidney data:

For the CCl_4 data:

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Exercise 3
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a > px = seq(-100, 500, length=50)
> f = function(x, b) log2(x+b)
> h = function(x, a) log2((x+sqrt(x^2+a^2))/2)
> matplot(px, y=cbind(h(px, a=50), f(px, b=50)),
        type="l", lty=1:2, xlab="x", ylab="f, h")
```

b Repeat the plot command above, with

> px = seq(0, 1e8, length=50)

c This question is very nicely explored in a paper by ?.

Exercise 4

The comparison of the dashed line with the data shows that there are systematic dye-related differences between the two channels. The dotted line shows that they can be fitted by an affine transformation, that is, a shift and a scaling of the data from one dye to adjust them to the other.