IO17 | Large Scale Bioinformatics for Immuno-Oncology

Deconvolution: exercise 2

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Implement an R function that computes the root-mean-square error (RMSE) between two **vectors** *yest* and *ytrue*

```r
getrmse <- function (yest,ytrue) {

    rmse <- …
    return(rmse)
}
```