Measurement error model for correlation coefficient estimation

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Introduction

The `MeasurementError.cor` package fits a two-stage measurement error model for estimating correlation between two random variables under bivariate normality. It's application is perhaps most relevant for the gene expression data where both point and standard estimates are available. We have shown that the proposed measurement error corrected correlation estimate has lower bias compared with the usual sample pearson correlation. For details, refer to Ding and Gentleman (2003) as well as R help pages associated with each function.

The `cor.me.vector` and `cor.me.matrix` functions

The `cor.me.vector` calculates the measurement error model estimate of correlation between two observed vectors whereas `cor.me.matrix` calculates all pairwise measurement error model estimate of correlation in the matrix.

```
> library(MeasurementError.cor)
> exp <- matrix(abs(rnorm(100,1000,20)),ncol=10)
> se <- matrix(abs(rnorm(100,50,5)),ncol=10)
> cor.me.vector(exp[1,],se[1,],exp[2,],se[2,])

$estimate
     corr.me    corr.true          mu1          mu2          s1          s2
  -0.9622231    0.8314547 1006.2633828 1004.0943104    6.8922861    5.7536148
```

```
$counts
function gradient
     106       100

$convergence
[1] 1

> cor.me.matrix(exp,se)

$corr.true
            [,1]        [,2]        [,3]        [,4]        [,5]        [,6]
 [1,]   1.0000000   0.8314547   0.4667938   0.6732750 -0.46078867   0.54161400
 [2,]   0.8314547   1.0000000 -0.8937722   0.9498708   0.87775989 -0.81349604
 [3,]   0.4667938 -0.8937722   1.0000000   0.9374714   0.95999267   0.29308404
 [4,]   0.6732750   0.9498708   0.9374714   1.0000000   0.82975236 -0.85905524
 [5,]  -0.4607887   0.8777599   0.9599927   0.8297524   1.00000000 -0.03581306
 [6,]   0.5416140 -0.8134960   0.2930840 -0.8590552 -0.03581306   1.00000000
 [7,]  -0.8921477 -0.9601379   0.6458049   0.9357602 -0.61425596   0.37881552
 [8,]  -0.6953253   0.6200524 -0.7710386   0.8837656 -0.60299780 -0.27183599
 [9,]  -0.7712654 -0.1719785   0.3015360 -0.8386522   0.51686444 -0.89239506
[10,]   0.6569188   0.7132322   0.7854675   0.7041110   0.81234193   0.75792045
            [,7]        [,8]        [,9]       [,10]
 [1,]  -0.8921477 -0.6953253 -0.7712654   0.6569188
 [2,]  -0.9601379   0.6200524 -0.1719785   0.7132322
 [3,]   0.6458049 -0.7710386   0.3015360   0.7854675
 [4,]   0.9357602   0.8837656 -0.8386522   0.7041110
 [5,]  -0.6142560 -0.6029978   0.5168644   0.8123419
 [6,]   0.3788155 -0.2718360 -0.8923951   0.7579204
 [7,]   1.0000000   0.5049864 -0.7245576 -0.6570030
 [8,]   0.5049864   1.0000000 -0.7695131   0.3442996
 [9,]  -0.7245576 -0.7695131   1.0000000 -0.7567006
[10,]  -0.6570030   0.3442996 -0.7567006   1.0000000

>
```

the quantity of interest, i.e. the model estimate of the correlation between the true value of two random variables whereas cor.me is the model estimate of correlation between the measurement errors of the two random variables. The second quantity may not be of interest. mu1,mu2 and s1, s2 are the estimated mean and standard deviation of the two random variables.

`cor.me.matrix` only returns the estimated correlation matrix.

References

Beiying Ding and Robert Gentleman. Measurement error model for correlation coefficient estimation and its application in microarray analysis. 2003.