

# On the computation of the $p$ -d multivariate normal density

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### Parameter settings

```
N = 10000
p = 100
# Setting mu and sigma
mu = rep(0, p)
set.seed(20231025)
sigma = cov(matrix(rnorm(N * p), N, p))
```

### Evaluating $f(x)$ using cholesky decomposition

```
# Evaluating f(x) using cholesky decomposition
mn_cholesky = function(x, mu, sigma) {
  R = chol(sigma)
  det_sigma = prod(diag(R))^2
  quad_term = crossprod(solve(t(R)), x - mu))
  log_den1 = (-p/2) * log(2 * pi) - 0.5 * log(det_sigma) - 0.5 * (quad_term)
  return(log_den1)
}
```

### Evaluating $f(x)$ directly

```
mn_direct = function(x, mu, sigma) {
  log_den2 = (-p/2) * log(2 * pi) - 0.5 * log(det(sigma)) - 0.5 * t(x - mu) %*%
    solve(sigma) %*% (x - mu)
  return(log_den2)
}
```

### Let's evaluate $f(x)$

```
set.seed(20231025)
x = rnorm(p)
```

```

print(mn_cholesky(x, mu, sigma))

##          [,1]
## [1,] -136.9197

print(mn_direct(x, mu, sigma))

##          [,1]
## [1,] -136.9197

print(mvtnorm::dmvnorm(x, mu, sigma, log = TRUE))

## [1] -136.9197

```

## Comparison of computing time

```

library(microbenchmark)
microbenchmark(mn_cholesky(x, mu, sigma), mvtnorm::dmvnorm(x, mu, sigma, log = TRUE),
               mn_direct(x, mu, sigma))

## Unit: microseconds
##                                         expr      min       lq      mean
##   mn_cholesky(x, mu, sigma) 396.846 453.2685 612.9111
##   mvtnorm::dmvnorm(x, mu, sigma, log = TRUE) 366.059 436.9205 774.9825
##   mn_direct(x, mu, sigma) 1013.850 1087.1035 1421.1759
##   median      uq      max neval
##   495.2430 546.7740 10258.09    100
##   581.2935 721.6605 13752.34    100
##   1187.7070 1313.0580 20625.22    100

```