

Square Root Rank Estimation

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The function 'sqrtFitMSE' tries to fit sqrt law to the eigenvalues distribution close to the maximal noise eigenvalue. It gets the vector of eigenvalues 'ell', and also a value q . It performs the fitting assuming that the signal rank is q . The function returns the MSE for the best fit it could find.

```
sqrtFitMSE <- function(q, ell, d=10) {
  p <- length(ell)
  i <- p - which.max(ell[p:1]>0) + 1 #index of the last non-zero value at ell
  range <- (ell[q+1] - ell[i]) / d #we want to consider a range which is 1/d from the total range
  j <- which.min(ell > (ell[q+1]-range)) #we want to take eigenvalues that are between ell[q+1] and ell[j]
  # we should take ell from (q+1) to j
  if (j < q+10) {j <- q+10} #we want to use at least 10 eigenvalues
  x <- ell[(q+1):j]
  u <- ((p-q):(p-j+1))/(p)
  y <- (1-u)^(2/3)
  l <- (ell[q+1] + ell[q+2]) / 2
  if (q==0) {
    L <- Inf
  } else {
    L <- (ell[q] + ell[q+1]) / 2
  }
  constrainedFit <- nls(x ~ m + alpha*y, algorithm="port",
    start=c(m=x[1], alpha=(x[1]-x[2])/(y[1]-y[2])),
    lower=c(m=l,alpha=-Inf),
    upper=c(m=L,alpha=Inf)
  )
  return(mean(resid(constrainedFit)^2))
}
```

The function 'kEst' gets as input the vector of eigenvalues 'ell'. For each possible value 'q' of the rank (currently limited to q in 0:10) it uses 'sqrtFitMSE' to compute the MSE of fitting a sqrt law to the distribution of the eigenvalues, under the assumption of 'q' signals. It then estimate the rank by taking 'q' with the minimal MSE.

```
kEst <- function(ell, k_max=10) {
  mse <- rep(0,k_max+1)
  for (q in 0:max_k) {
    mse[q+1] <- sqrtFitMSE(q, ell)
  }
  k <- which.min(mse)-1
}
```

Testing the algorithm: we test the estimator with various values for p , where $n = p$, noise eigenvalues are $U[0.5, 1.5]$, and we have rank $k = 2$ with signal strengths $\lambda = (10, 4)$. For each value of p we make 100 iterations and show the fraction of correct estimations.

```

set.seed(265)
P <- 2^(5:9)
lam <- c(10,4)
k <- length(lam)
max_k <- 10
iter <- 100
frac <- rep(0,length(P))
for (i_P in 1:length(P)) {
  p <- P[i_P]
  n <- p
  count <- 0
  for (i_iter in 1:iter) {
    Sigma <- diag(c(lam, rep(0,p-k)) + runif(p,0.5,1.5))
    S <- rWishart(1,n,Sigma)
    S <- S[,1]/n
    ell <- eigen(S,TRUE,TRUE)$values
    k_hat <- kEst(ell)
    if (k_hat == k) {count <- count+1}
  }
  frac[i_P] <- count/iter
}
plot(P, frac, ylim=c(0,1), pch=16, main="Probability of Estimating the Correct Number of Signals as function of P",
abline(1, 0, col="red", lwd=3))

```

Probability of Estimating the Correct Number of Signals as function c

