```
1.
mylm <- function(y,X){</pre>
    # Transform to matrix
   Y = as.matrix(Y); Xnew = as.matrix(X)
    # Check if numeric
   if(!is.numeric(Y) | !is.numeric(Xnew))
    stop("Y or X is not numeric!\n")
    # Check for dimensions
   dy = dim(Y); dx = dim(Xnew)
   if(dy[2] != 1 | dy[1] != dx[1])
   stop("Y or X has wrong dimensions\n")
    # Check for ill conditioned elements.
    # We can use is.finite to response only to finite real numbers
   if(FALSE %in% is.finite(Y) | FALSE %in% is.finite(Xnew))
   warning("Y or X is ill conditioned\n")
    # Check if of full rank
   D = cbind(1, Xnew)
   DtD = t(D) \% \% D
    if(det(DtD) == 0)
    stop("Design matrix is not full rank\n")
   # Regressing
   DtD.inv = solve(DtD)
   hat.matrix = D %*% DtD.inv %*% t(D)
   beta = DtD.inv %*% t(D) %*% Y
   fitted = D %*% beta
   residuals = Y - fitted
   SS.tot = sum((Y - mean(Y))^2)
   if(SS.tot == 0)
   warning("Y is constant!\n")
   SS.res = sum((Y - fitted)^2)
   SS.reg = SS.tot - SS.res
   R2 = SS.reg / SS.tot
    # Return result
   result = list(beta = beta,
                fitted = fitted,
                residuals = residuals,
                R2 = R2
                hatdiag = diag(hat.matrix))
   return(result)
}
  2.
```

```
data = read.table("HW2ex.txt")
shape = dim(data)

Y = data[,1]; X = data[,2:shape[2]]

results = mylm(Y,X)
plot(results$fitted, results$residuals, pch='.')
```

