```
\begin{array}{l} 1 \leq \\ j \leq \\ d \\ j \\ x_1, \dots, x_M \end{array}
        density
     density stats X_1, \dots, Y_N X_1, \dots, X_N 
        \begin{array}{c} 0.516\ 1.036\ 0.956\\ 0.6550\ 722\ 2.15\\ 0.698\ 1.92\ 2.72\\ 0.588\ 1.80\ 3.34\\ mRNA.pngInblack, kerneldensity estimates of the densities of mRNA gene expression for each level of polyQlength (Q80, Qandalan expression of the polyQlength). \\ \end{array}
           andusingalog(1+
        {}_m^{\circ}iRNA.pngInblack, kerneldensity estimates of the densities of miRNA gene expression for each level of polyQlength (Q80, and using a log (1+
 \begin{cases} \omega \overset{\hookrightarrow}{=} \\ (+)^M | \sum_{m \in M} \omega_m = \\ 1 \} \\ (M - 1) \\ \omega \overset{\rightleftharpoons}{=} \\ M^{-1} \mathbf{1}_M \\ \mathbf{1}_M \overset{\longleftrightarrow}{=} \\ \Omega \\ \Pi(\omega) := \\ \{P \in \\ (+)^{M \times N} | P \mathbf{1}_N = \\ \omega, P^{\top} \mathbf{1}_M = \end{cases} 
  (+)^{M \times N} | P \mathbf{1}_{N} = (+)^{M \times N} | P \mathbf{1}_{N} 
           c(x,y) :=
           ||x-
  \ddot{y} \parallel_2^2
        C_{X,Y} \in {}^{M \times N} 
 (C_{X,Y})_{m,n} :=
        c(x_m, y_n)
(m, n) \in M \times N
N
  N_{\min_{P \in \Pi(\bar{\omega})} \langle C_{X,Y}, P \rangle_F} \\ \langle C_{X,Y}, P \rangle_F := \\ \sum_{(m,n) \in M \times N} (C_{X,Y})_{mn} P_{mn} \\ Y \\ P \\ E(P) :=
```