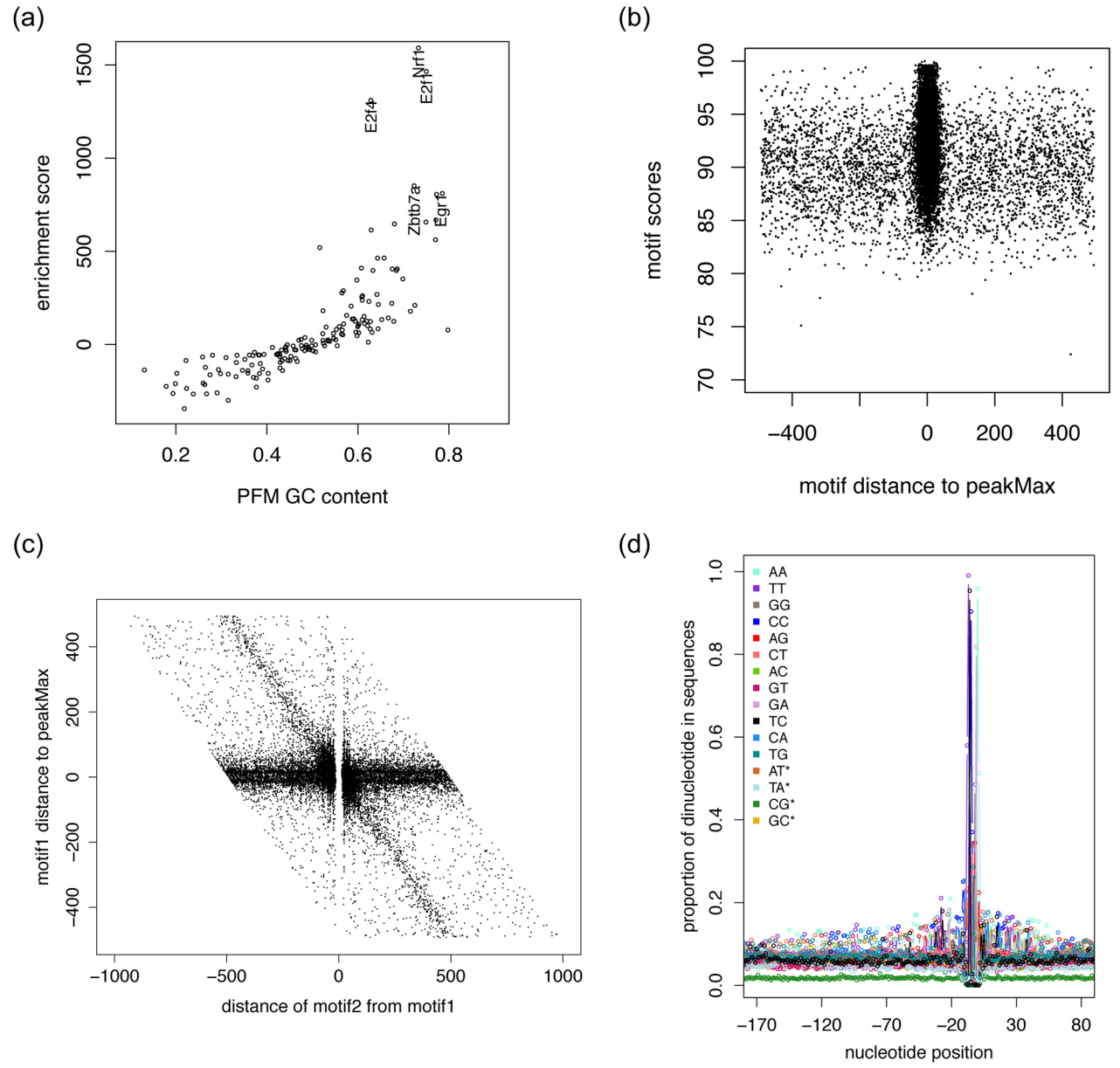
Figure 11


(a)
enrichment score
PFM GC content
E2f4
E2f1
Nf1
Zbtb7a
Egr6
(b)
motif scores
motif distance to peakMax
(c)
motif1 distance to peakMax
distance of motif2 from motif1
(d)
proportion of dinucleotide in sequences
nucleotide position
AA
TT
GG
CC
AG
CT
AC
GT
GA
TC
CA
TG
AT*
TA*
CG*
GC*