AT-rich interaction domain family
basic region + helix-loop-helix motif + leucine zipper; 1.3
basic region + helix-loop-helix motif; 1.2
basic region + helix-span-helix domain; 1.6
basic region + leucine zipper; 1.1
Factor family with homology to the ets-protooncogenes; 3.5.2
fork head domain; 3.3
heat stress transcription factor; 3.4
helix-loop-helix motif; 1.2.8
high-mobility group protein-like factors; 4.7
homeodomain; homeobox protein; 3.1
LIM domain plus homeo domain; 3.1.3.1
LIM domain; 3.1.3.2
MCM1-agamous-deficiens-SRF; 4.4
paired domain, paired box + homeo domain; 3.2.1
paired domain, paired box; 3.2
POU domain; 3.1.2
Rel-related factor; 4.1
RING finger
runt homology domain; 4.11
signal transducers and activators of transcription
T domain, T-box proteins
tryptophan cluster; 3.5
zinc finger + homeo domain; 3.1.4
zinc finger; 2.2
zinc finger; 2.3
zinc finger; zinc twist; 2.1

TFBS Motif Hits in pre-miRs
Excluding Primate-Specific

Hits in pre-miR (z=2.33)  Randomized control