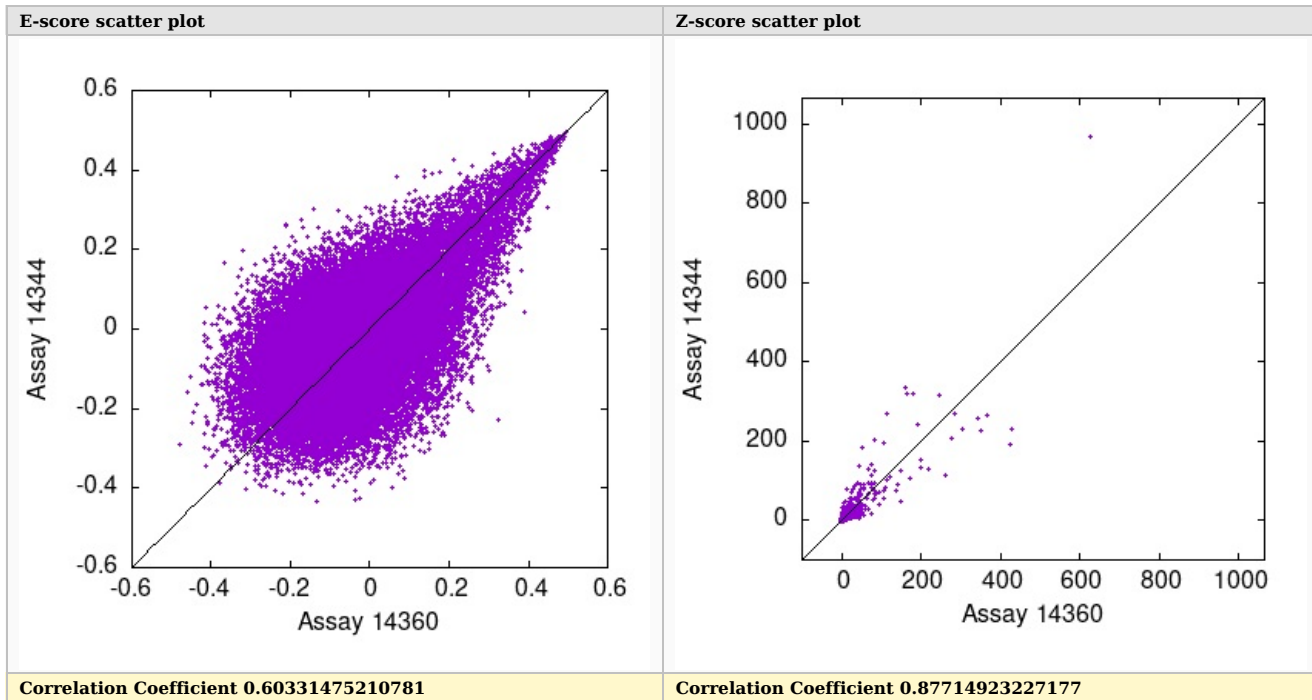


QC report



Top scoring motifs for Assay 14360

Protein ID: pTH15544.2 Gene: NR4A2 Domain: zf-C4 Flag: Pass_matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

Reverse:

AAAGGTCA

TGACCtTT

Top 10

Scores

Alignment

AAAGGTCA	0.49348	-AAAGGTCA-
AAGGTCAC	0.48971	--AAGGTCAC
TAAGGTCA	0.48872	-TAAGGTCA-
AAAGGTCG	0.48826	-AAAGGTCG-
AAGGTCAT	0.48741	--AAGGTCAT
AAATGTCA	0.48695	-AAATGTCA-
AAAGGGCA	0.48544	-AAAGGGCA-
CAAAGGTC	0.48357	CAAAGGTC--
GAAGGTCA	0.48088	-GAAGGTCA-
AAGGTCAG	0.48061	--AAGGTCAG

8 mer Z-scores for probeset 'all'

Forward:

Reverse:

AAAAGGTCA

TGACCtTT

Top 10

Scores

Alignment

AAAGGTCA	624.83983	-AAAGGTCA-
AAGGTCAT	427.19946	--AAGGTCAT
TAAGGTCA	425.62836	-TAAGGTCA-
AAAGGTCG	365.65746	-AAAGGTCG-
AAGGTCAC	351.90498	--AAGGTCAC
AAATGTCA	344.43462	-AAATGTCA-
CAAAGGTC	305.31815	CAAAGGTC--
AAAGGGCA	284.44688	-AAAGGGCA-
AAAAGGTC	277.08247	AAAAGGTC--
GACATTTA	261.68568	TAAATGTC--

Top scoring motifs for Assay 14344

Protein ID: pTH15544.1 Gene: NR4A2 Domain: zf-C4 Flag: Pass_matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:

Reverse:

AAAGGTCA

TGACCtTT

Top 10

Scores

Alignment

AAAGGTCA	0.49720	-AAAGGTCA-
AAAGGGCA	0.48661	-AAAGGGCA-
ATAGGTCA	0.48570	-ATAGGTCA-
AAAGGTCA	0.48562	-AAAGGTCA-
GACCTTTA	0.48406	TAAAGGTC--
AAATGTCA	0.48373	-AAATGTCA-
AAGGTCAA	0.48352	--AAGGTCAA
AAGGTCAG	0.48338	--AAGGTCAG
AAAGTTCA	0.48281	-AAAGTTCA-
CAAAGGTC	0.48245	CAAAGGTC--

8 mer Z-scores for probeset 'all'

Forward:

Reverse:

AAAAGGTCA

TGACCtTT

Top 10

Scores

Alignment

AAAGGTCA	967.45363	-AAAGGTCA-
ATAGGTCA	336.42034	-ATAGGTCA-
GAAGGTCA	319.13427	-GAAGGTCA-
AAGGTCAG	319.02657	--AAGGTCAG
AAAGGTCA	314.94963	-AAAGGTCA-
AAAGGGCA	269.47171	-AAAGGGCA-
GACCTTTA	269.05222	TAAAGGTC--
AAAGGTCG	264.97656	-AAAGGTCG-
AAATGTCA	257.02256	-AAATGTCA-
AAGGTCAA	241.13958	--AAGGTCAA