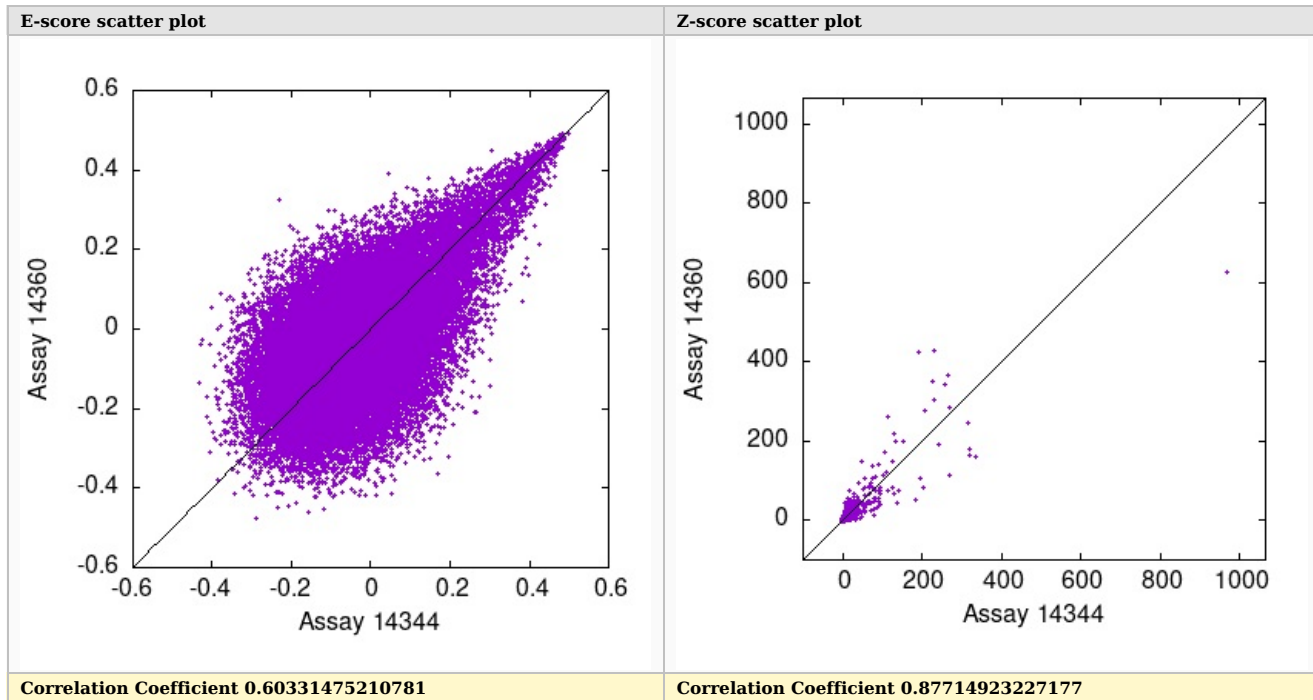


QC report



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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
AAAGGTCA	0.49720	-AAAGGTCA-	AAAGGTCA	967.45363	-AAAGGTCA-
AAAGGCA	0.48661	-AAAGGCA-	ATAGGTCA	336.42034	-ATAGGTCA-
ATAGGTCA	0.48570	-ATAGGTCA-	GAAGGTCA	319.13427	-GAAGGTCA-
AAAGGTTA	0.48562	-AAAGGTTA-	AAGGTCAG	319.02657	-AAGGTCAG
GACCTTTA	0.48406	TAAAGGTC--	AAAGGTTA	314.94963	-AAAGGTTA-
AAATGTCA	0.48373	-AAATGTCA-	AAAGGCA	269.47171	-AAAGGCA-
AAGGTCAA	0.48352	--AAGGTCAA	GACCTTTA	269.05222	TAAAGGTC--
AAGGTCAG	0.48338	--AAGGTCAG	AAAGGTCG	264.97656	-AAAGGTCG-
AAAGTTCA	0.48281	-AAAGTTCA-	AAATGTCA	257.02256	-AAATGTCA-
CAAAGGTC	0.48245	CAAAGGTC--	AAGGTCAA	241.13958	--AAGGTCAA

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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
AAAGGTCA	0.49348	-AAAGGTCA-	AAAGGTCA	624.83983	-AAAGGTCA-
AAGGTCAC	0.48971	--AAGGTCAC	AAGGTCAT	427.19946	--AAGGTCAT
TAAGGTCA	0.48872	-TAAGGTCA-	TAAGGTCA	425.62836	-TAAGGTCA-
AAAGGTCG	0.48826	-AAAGGTCG-	AAAGGTCG	365.65746	-AAAGGTCG-
AAGGTCAT	0.48741	--AAGGTCAT	AAGGTCAC	351.90498	--AAGGTCAC
AAATGTCA	0.48695	-AAATGTCA-	AAATGTCA	344.43462	-AAATGTCA-
AAAGGCA	0.48544	-AAAGGCA-	CAAAGGTC	305.31815	CAAAGGTC--
CAAAGGTC	0.48357	CAAAGGTC--	AAAGGCA	284.44688	-AAAGGCA-
GAAGGTCA	0.48088	--GAAGGTCA-	AAAAGGTC	277.08247	AAAAGGTC--
AAGGTCAG	0.48061	--AAGGTCAG	GACATTTA	261.68568	TAAATGTC--