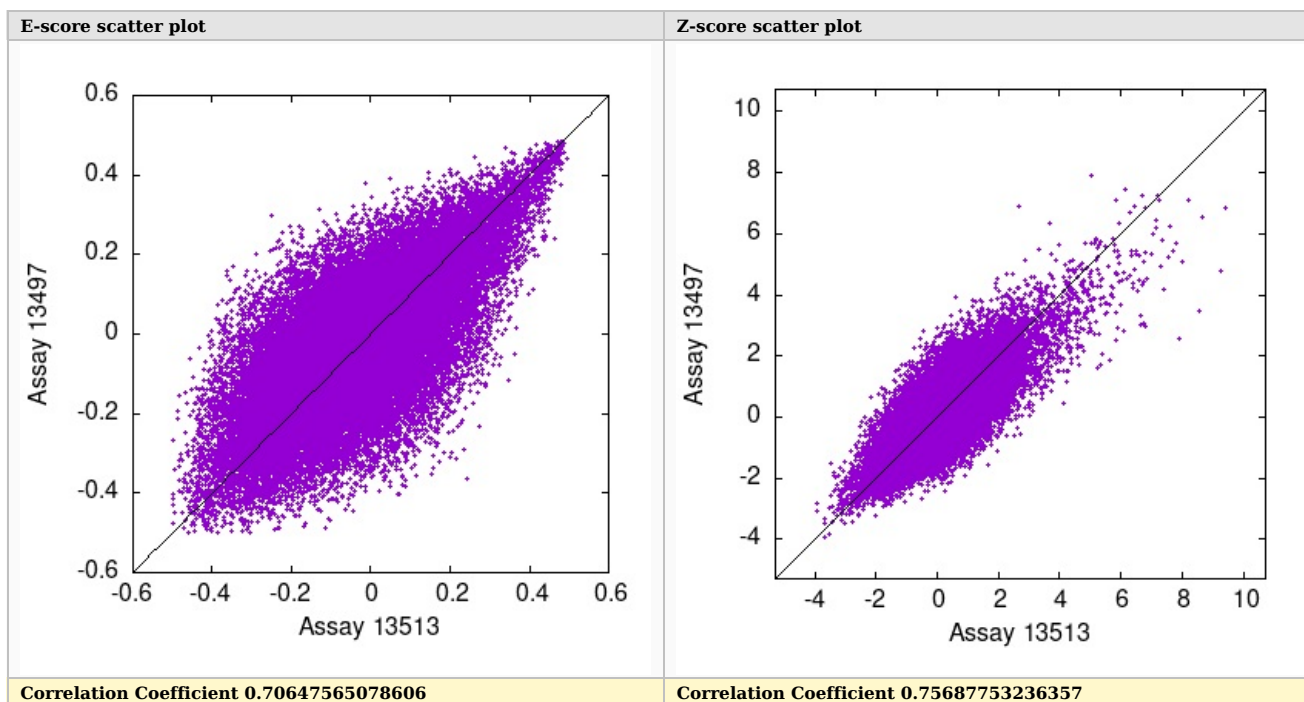




QC report



Top scoring motifs for Assay 13513

Protein ID: pTH13913.2 Gene: AHCTF1.DBD Domain: AT_hook Flag: Pass_matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:	Reverse:	
		
Top 10	Scores	Alignment
AAATAAAT	0.49357	--AAATAAAT--
AATTAATT	0.48760	--AATTAATT--
AATAAATT	0.48584	---AATAAATT
AATTTAAA	0.48509	---TTTAAATT
AAAAAATA	0.48473	---AAAAAATA
AAAAATAT	0.48382	---AAAAATAT--
AAAAATAA	0.48167	AAAAATAA----
AATAATAA	0.48042	---AATAATAA
ATTTAAAT	0.47921	--ATTTAAAT--
AAAAAATA	0.47898	---AAAAAATA



8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
Top 10	Scores	Alignment	
AATTAATT	9.39101	-AATTAATT-	
AAATAAAT	9.28751	AAATAAAT--	
AAAAATAT	8.66712	--ATATTTTT	
AATTTAAA	8.54261	-TTTAAATT-	
AAAAAATA	8.21743	-AAAAAATA-	
AATATATT	8.01288	-AATATATT-	
ATTTTTTC	7.91754	GAAAAAAT--	
AATAATAA	7.81197	-AATAATAA-	
AATAAATT	7.70647	-AATAAATT-	
AAAATATT	7.60208	-AAAATATT-	

Top scoring motifs for Assay 13497

Protein ID: pTH13913.1 Gene: AHCTF1.DBD Domain: AT_hook Flag: Pass_matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:	Reverse:	
		
Top 10	Scores	Alignment
AAATATTT	0.48374	--AAATATTT--
AAAAAATA	0.48304	--AAAAAATA--
ATTTAAAT	0.48291	--ATTTAAAT--
AAAAATAA	0.48082	AAAAATAA----
AAAAAAT	0.48026	--AAAAAAT--
AATTAATT	0.47835	--AATTAATT--
ATAATATA	0.47644	----ATAATATA
ATTATTTA	0.47640	-TAAATAAT--
AAATAAAT	0.47460	--AAATAAAT--
AAAATATT	0.47356	---AATATTTT-

8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
Top 10	Scores	Alignment	
AAAAAAT	7.89610	-AAAAAAT-	
AAATATTT	7.43353	-AAATATTT-	
AAAAATAA	7.25444	-AAAAATAA-	
ATTTAAAT	7.22518	-ATTTAAAT-	
AAAAAATA	7.11045	--AAAAAATA	
AATTTAAT	7.10444	-AATTTAAT-	
ATTATTTA	7.07541	-TAAATAAT-	
CAAAAAAA	6.91058	CAAAAAAA - -	
AAATAAAT	6.90748	-AAATAAAT-	
AATTAAT	6.86585	-AATTAAT-	