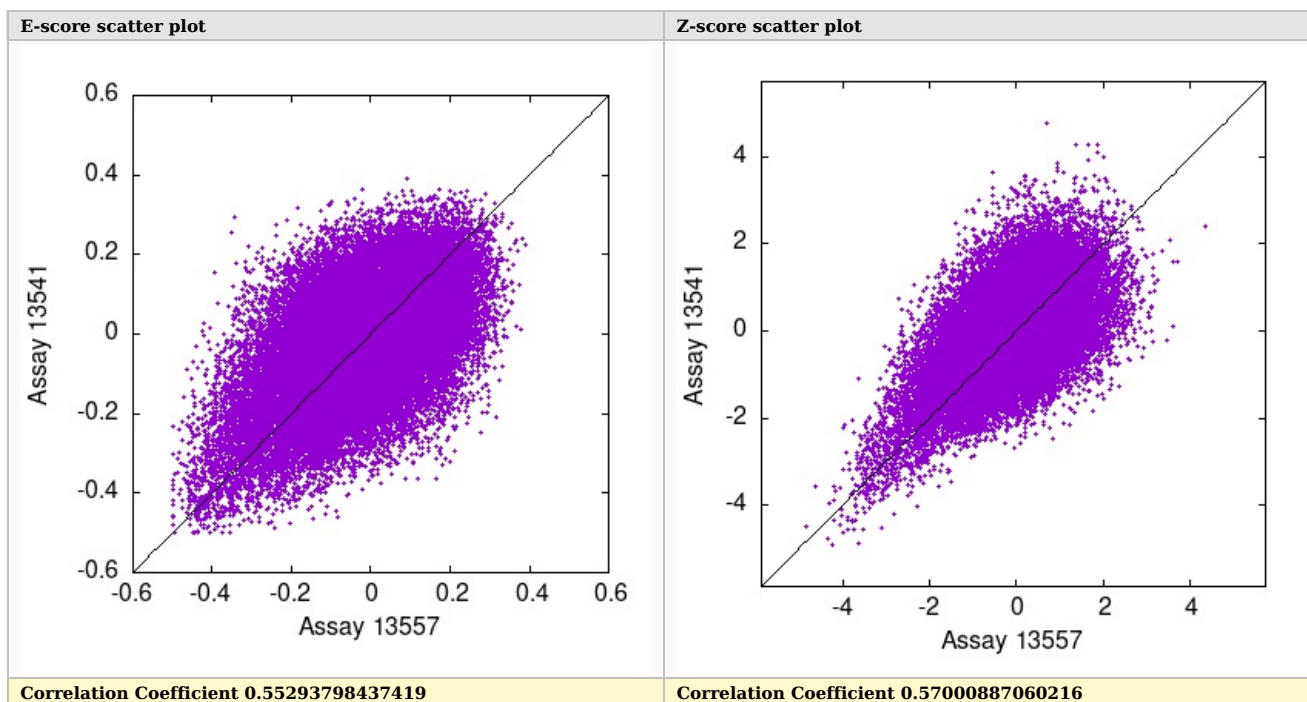


QC report





Top scoring motifs for Assay 13557

Protein ID: pTH14288.2 Gene: CENPT.FL Domain: Unknown Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:		Reverse:	
Top 10	Scores	Alignment	
AAGTACTT	0.38890	AAGTACTT-----	
TACGCGTA	0.38234	---TACGCGTA---	
GAAATTTT	0.37691	--GAAATTTT---	
TCGCGCGA	0.37565	---TCGCGCGA---	
TTGCGCAA	0.37157	-----TTGCGCAA	
CGTCGGAA	0.36867	----CGTCGGAA--	
TAGTTGCA	0.36505	---TAGTTGCA---	
ATAATGTA	0.36428	---ATAATGTA---	
GTACAGTA	0.35961	-----GTACAGTA	
GTATACTA	0.35768	---TAGTATAC---	

8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
TACGCGTA	4.35058	-----TACGCGTA	
TTGCGCAA	3.69983	---TTGCGCAA---	
ATAATGTA	3.60263	ATAATGTA-----	
CACAGTAG	3.58399	CTACTGTG-----	
ATATGTGC	3.53777	-----GCACATAT-	
TCGCGCGA	3.45373	---TCGCGCGA---	
ATTGTACA	3.35293	--ATTGTACA---	
TATTTGCA	3.25446	----TGCAAATA--	
GTATTGTA	3.17908	GTATTGTA-----	
GAATATTC	3.15232	-GAATATTC-----	



Top scoring motifs for Assay 13541

Protein ID: pTH14288.1 Gene: CENPT.FL Domain: Unknown Flag: Reject Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:	Reverse:	
Top 10	Scores	Alignment
TGTGCACA	0.39149	---TGTGCACA--
AAAGTACA	0.36464	-----TGTACTTT
ACGTAAACA	0.36305	---ACGTAAACA---
ATATTGTA	0.36116	----ATATTGTA-
AACAAAGC	0.36001	---AACAAAGC---
ATGTACAT	0.35918	----ATGTACAT-
ATATATGA	0.35887	--TCATATAT---
ACAACAAT	0.35445	---ACAACAAT---
ATTCGTTA	0.35364	TAACGAAT-----
ACATATAC	0.35255	---ACATATAC---

8 mer Z-scores for probeset 'all'

Forward:		Reverse:
		
Top 10	Scores	Alignment
ATATTGTA	4.75439	ATATTGTA---
ATTCGTTA	4.27385	-ATTCGTTA--
ACAACAAT	4.27157	-ATTGTTGT--
TGTGCACA	4.26695	--TGTGCACA-
ATGTACAT	4.09826	---ATGTACAT
ATTTATAA	4.00548	-ATTTATAA--
ATTTGTAC	3.91244	-ATTTGTAC--
AACAAAGC	3.84111	GCTTTGTT---
AACATGTT	3.72651	AACATGTT---
ATGTTAAT	3.67272	-ATGTTAAT--