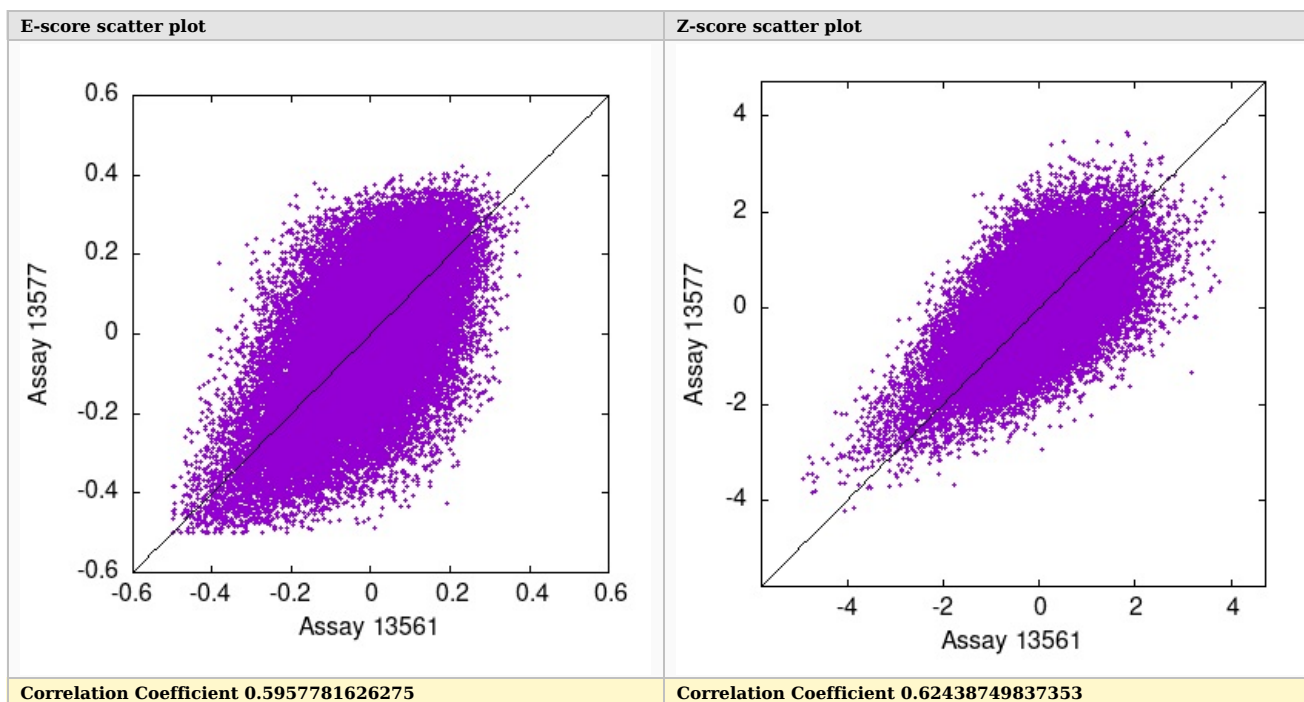


QC report



Top scoring motifs for Assay 13561

Protein ID: pTH13916.1 Gene: DNTTIP1.FL Domain: AT_hook Flag: Reject 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
AATGCATT	0.39303	--AATGCATT--	AAATGCAT	3.84282	-AAATGCAT-
AAATGCAT	0.38275	-AAATGCAT--	ACAACAAT	3.81978	-ACAACAAT-
AGCAATTG	0.37590	CAATTGCT--	AATGCATT	3.77419	--AATGCATT
ACAACAAT	0.36915	-ACAACAAT--	CAATTGCG	3.76626	CAATTGCG--
TACAACAA	0.36847	TTGTTGTA--	TGAAGACA	3.66135	TGAAGACA--
AACATGCA	0.36550	AACATGCA--	TAACGAAA	3.62723	--TAACGAAA-
AAGCAATT	0.36549	-AATTGCTT--	TACAACAA	3.60219	TTGTTGTA--
TAACGAAA	0.35580	-TAACGAAA--	AACATGCA	3.58763	AACATGCA--
ATACAACA	0.35082	---ATACAACA	ACAACGCA	3.57475	ACAACGCA--
ATAATGCA	0.34657	ATAATGCA--	AGCAATTG	3.53843	CAATTGCT--

Top scoring motifs for Assay 13577

Protein ID: pTH13916.2 Gene: DNTTIP1.FL Domain: AT_hook Flag: Reject 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
ACTAATTA	0.42042	-ACTAATTA-	CTGTATTA	3.67092	--TAATACAG-
CTGTATTA	0.40728	CTGTATTA--	GAATATTC	3.58962	--GAATATTC-
GAATATTC	0.40397	GAATATTC--	ACTAACAT	3.47709	---ACTAACAT
GATATGTA	0.40293	-GATATGTA-	ATATCACA	3.46030	---TGTGATAT
TAATAATA	0.40267	TAATAATA--	TTCAACAA	3.42498	TTCAACAA--
TTCAACAA	0.40243	--TTGTTGAA	TATTTGCA	3.41715	--TGCAAAATA-
AATATTAA	0.40011	-AATATTAA-	TATTAATA	3.41432	--TATTAATA-
ATAGTAAT	0.40006	--ATAGTAAT	AATATTAA	3.18637	---AATATTAA
ATATCACA	0.39712	--ATATCACA	TAATAATA	3.16052	--TAATAATA-
AAACGTTT	0.39501	AAACGTTT--	ATTATAAT	3.14394	-ATTATAAT--