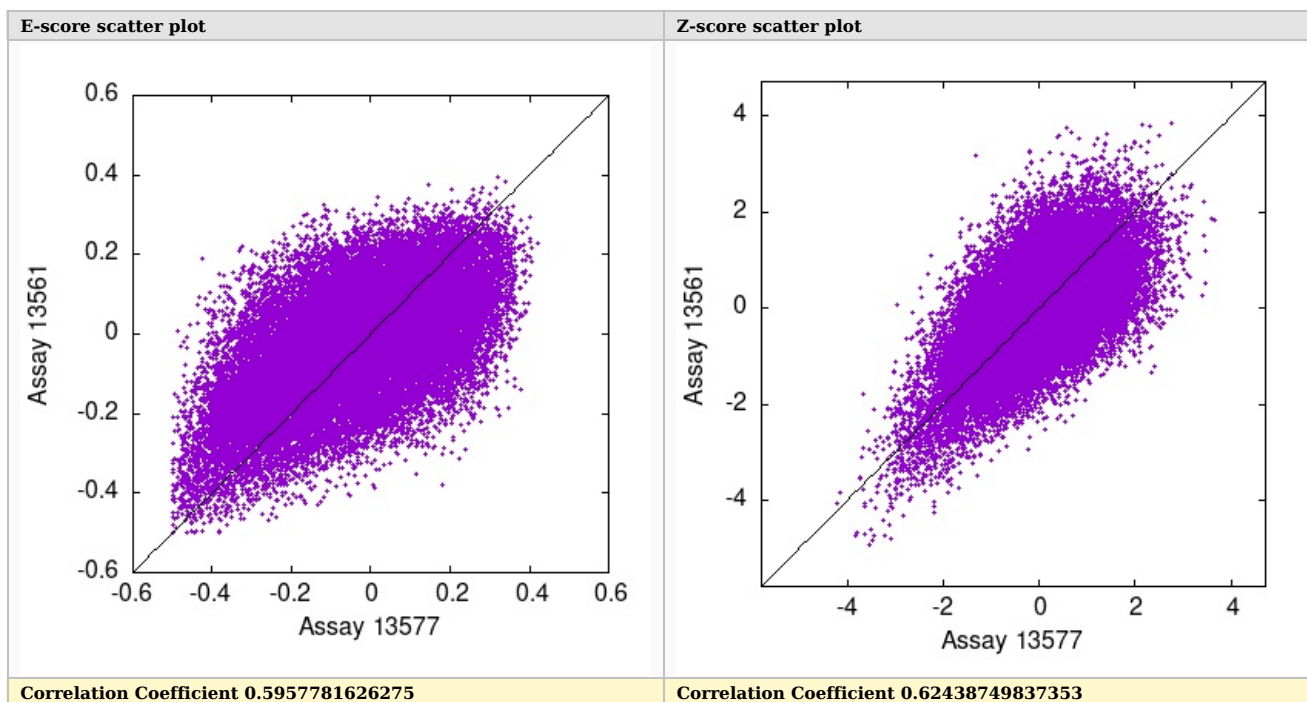


QC report



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'

Forward:

Reverse:



Top 10

Scores

Alignment

ACTAATTA	0.42042	-ACTAATTA-
CTGTATTA	0.40728	CTGTATTA--
GAATATTC	0.40397	GAATATTC--
GATATGTA	0.40293	-GATATGTA-
TAATAATA	0.40267	TAATAATA--
TTCAACAA	0.40243	--TTGTTGAA
AATATTAA	0.40011	-AATATTAA-
ATAGTAAT	0.40006	--ATAGTAAT
ATATCACA	0.39712	--ATATCACA
AAACGTTT	0.39501	AAACGTTT--

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

CTGTATTA	3.67092	--TAATACAG-
GAATATTC	3.58962	--GAATATTC-
ACTAACAT	3.47709	---ACTAACAT
ATATCACA	3.46030	---TGTGATAT
TTCAACAA	3.42498	TTCAACAA---
TATTTGCA	3.41715	--TGCAAAATA-
TATTAATA	3.41432	--TATTAATA-
AATATTAA	3.18637	---AATATTAA
TAATAATA	3.16052	--TAATAATA-
ATTATAAT	3.14394	-ATTATAAT--

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'

Forward:

Reverse:



Top 10

Scores

Alignment

AATGCATT	0.39303	--AATGCATT-
AAATGCAT	0.38275	-AAATGCAT--
AGCAATTG	0.37590	CAATTGCT---
ACAACAAT	0.36915	-ACAACAAT--
TACAACAA	0.36847	TTGTTGTA--
AACATGCA	0.36550	AACATGCA---
AAGCAATT	0.36549	-AATTGCTT--
TAACGAAA	0.35580	--TAACGAAA--
ATACAACA	0.35082	---ATACAACA
ATAATGCA	0.34657	ATAATGCA---

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

AAATGCAT	3.84282	-AAATGCAT-
ACAACAAT	3.81978	-ACAACAAT-
AATGCATT	3.77419	--AATGCATT
CAATTGCG	3.76626	CAATTGCG--
TGAAGACA	3.66135	TGAAGACA--
TAACGAAA	3.62723	-TAACGAAA-
TACAACAA	3.60219	TTGTTGTA--
AACATGCA	3.58763	AACATGCA--
ACAACGCA	3.57475	ACAACGCA--
AGCAATTG	3.53843	CAATTGCT--