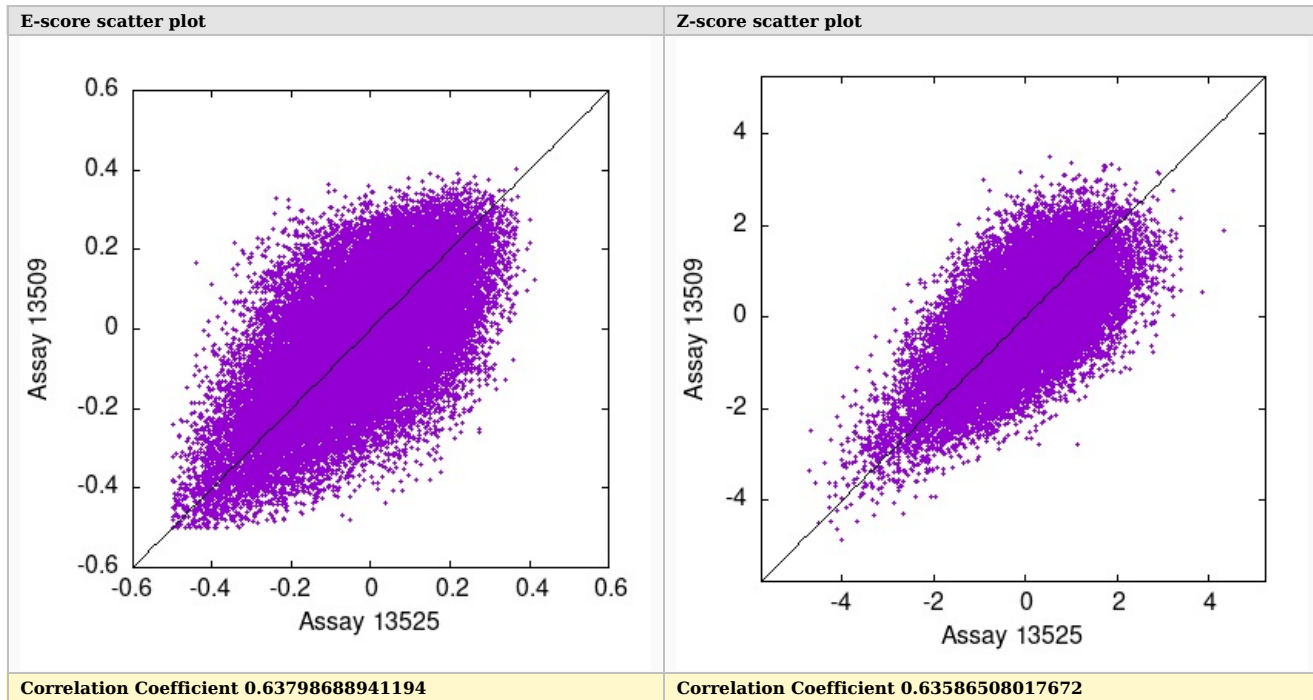


QC report



Top scoring motifs for Assay 13525

Protein ID: pTH14239.2 Gene: ADNP.DBD.1 Domain: Homeobox Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

Reverse:



8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

GTCATGAC	0.41536	-----GTCATGAC
TACTAGTA	0.40346	---TACTAGTA--
ATGCGCAT	0.40033	--ATGCGCAT---
AACGTTTG	0.39521	CAAACGTT-----
GATGCATC	0.39308	---GATGCATC--
CAACAATG	0.38459	---CAACAATG--
ATAGTAAT	0.37179	--ATAGTAAT---
AACGTATT	0.37145	--AACGTATT---
GACAAACA	0.36991	---GACAAACA--
TATAACGA	0.36955	---TCGTTATA--

Top 10

Scores

Alignment

TACTAGTA	4.31890	TACTAGTA-----
GTCATGAC	3.84475	---GTCATGAC---
AACGTTTG	3.39400	--AACGTTTG---
CGCAACAA	3.38606	---TTGTTGCG---
ATGTATGC	3.38206	---ATGTATGC---
CAACAATG	3.37589	--CAACAATG---
ATTATAAT	3.37101	---ATTATAAT---
AAATTACG	3.29589	---AAATTACG---
ATATGCTG	3.27850	-----ATATGCTG
CGTATAAA	3.23410	---TTTATACG---

Top scoring motifs for Assay 13509

Protein ID: pTH14239.1 Gene: ADNP.DBD.1 Domain: Homeobox Flag: Reject Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:

Reverse:



8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

ACAAACAT	0.40105	-ATGTTTGT--
AATTTATG	0.38986	--AATTTATG--
AATTTGTA	0.38943	--AATTTGTA--
AACATTAA	0.38340	TTAATGTT---
CAAATTTG	0.38283	CAAATTTG---
ATATATAA	0.37806	---ATATATAA
AATTGTGA	0.37670	--AATTGTGA--
ATAATGCA	0.37562	--ATAATGCA--
TATAATAA	0.37389	--TTATTATA--
AACATGTT	0.37184	--AACATGTT--

Top 10

Scores

Alignment

AATTTATG	3.50058	--AATTTATG--
CAAATTTG	3.36253	CAAATTTG----
AACATTAA	3.31713	---TTAATGTT---
AATGCATT	3.30075	--AATGCATT--
AATTTGTA	3.28358	--AATTTGTA--
AATATTTT	3.25227	--AATATTTT--
ACATGTAA	3.20706	-ACATGTAA---
AACAAAAT	3.20424	--ATTTTGT---
AACTTTAT	3.14421	--AACTTTAT---
AGAATTCT	3.14349	-AGAATTCT---