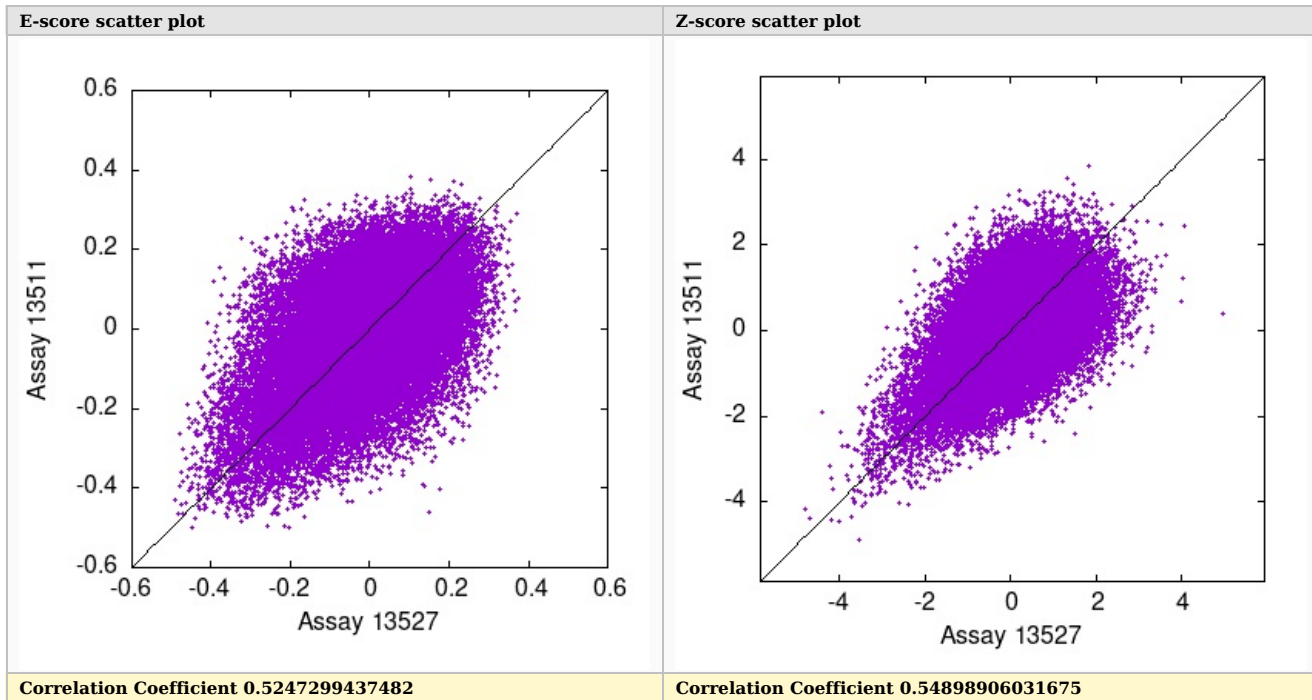


QC report



Top scoring motifs for Assay 13527

Protein ID: pTH14241.2 Gene: ADNP2.DBD.1 Domain: Homeobox Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

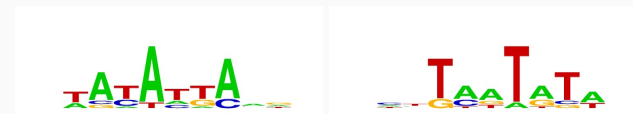
Reverse:



8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10	Scores	Alignment	Top 10	Scores	Alignment
ACAGTAGA	0.37383	-----ACAGTAGA	ATTATAAT	4.94634	--ATTATAAT--
ATATTAAG	0.37221	--ATATTAAG----	ATATTAAG	4.04129	--ATATTAAG--
ATTATAAT	0.36654	--ATTATAAT----	ACAACAG	4.03452	--ACAACAG--
GACAAACA	0.36596	-GACAAACA-----	ATATGAGA	3.97196	--ATATGAGA--
ACAACAG	0.36358	--ACAACAG-----	CTGTATTA	3.63147	CTGTATTA---
TACGCGTA	0.35761	-TACGCGTA-----	AACACTAA	3.50674	--AACACTAA--
AAAACTTA	0.35589	---AAAACTTA----	ATAATGTA	3.50137	-TACATTAT--
ACAGCATA	0.35175	-----ACAGCATA	ATACTACA	3.42207	--ATACTACA--
ATACAAAT	0.35020	ATACAAAT-----	CAATGATG	3.38134	--CAATGATG--
ATTCACAG	0.34918	--ATTCACAG----	TATGCATA	3.37304	---TATGCATA

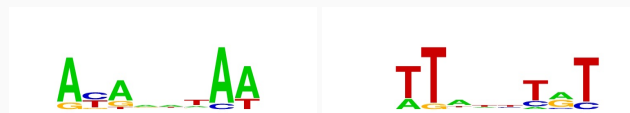
Top scoring motifs for Assay 13511

Protein ID: pTH14241.1 Gene: ADNP2.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	Top 10	Scores	Alignment
CAGAAACA	0.38405	CAGAAACA---	ATACAACA	3.85240	--ATACAACA---
GTACTTAA	0.37672	-GTACTTAA--	ATGTATAA	3.56877	TTATACAT----
ACAATAA	0.37080	-ACAATAA--	ATGGTTTA	3.40301	---TAAACCAT--
TAACGAAA	0.36236	--TAACGAAA--	GTAATGGA	3.28268	---TCCATTAC--
ATTATCTG	0.35296	--CAGATAAT--	AAATTAAG	3.24984	----AAATTAAG--
ATGTATAA	0.34681	-ATGTATAA--	AAGAATAA	3.23054	----AAGAATAA--
GATGAGTA	0.34025	GATGAGTA---	CAGAAACA	3.21385	-----CAGAAACA
ACAGAACT	0.33979	-ACAGAACT--	TAACGAAA	3.19964	---TAACGAAA--
ACAGTAAT	0.33797	-ACAGTAAT--	TACATAAA	3.15980	---TACATAAA--
CGTTAATA	0.33654	---TATTAACG	ACAACTAT	3.15965	----ACAACTAT--