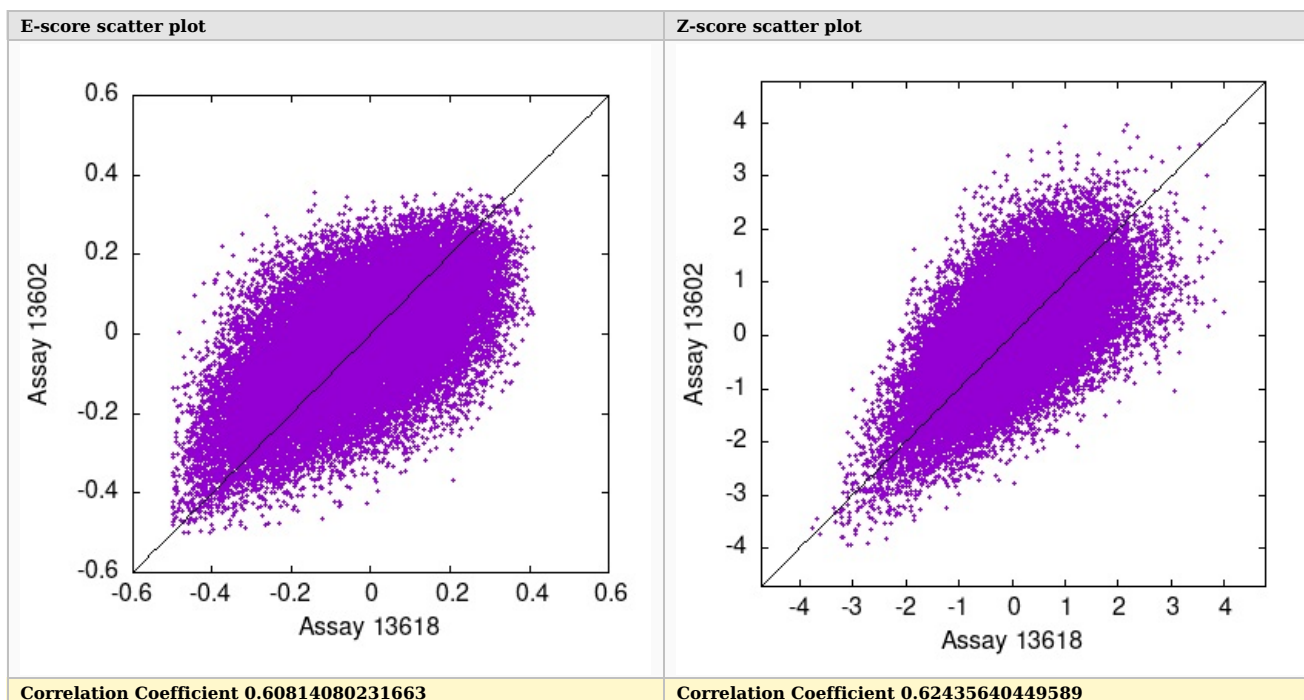


QC report



Top scoring motifs for Assay 13618

Protein ID: pTH14299.2 Gene: DACH2.DBD Domain: Unknown Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

AATAAATT	0.40900	-AATTATT-
GATTTATA	0.40896	-GATTTATA-
GATATGTA	0.40639	-GATATGTA-
AATTTATA	0.40235	-AATTTATA-
AATTTAAA	0.40052	-AATTTAAA-
ATTATTTT	0.39732	GAAATAAT--
ATTAAAT	0.39498	--ATTAAAT
ATAAATTA	0.39472	TAATTTAT--
TATCAACA	0.39444	-TGTTGATA-
ATATTAAG	0.39351	--CTTAATAT

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

ATTATTTT	3.98078	--ATTATTTT--
CTATATTA	3.93379	--CTATATTA--
GAAATTTT	3.82215	--GAAATTTT--
GATATGTA	3.80010	-GATATGTA--
CTGTATTA	3.77616	TAATACAG--
AATTTAAA	3.68574	-TTTAAATT--
ATAATGTA	3.67093	--TACATTAT
GATTTATA	3.66407	-TATAAATC--
TATTAATA	3.61840	-TATTAATA--
AATAAATT	3.59110	-AATAAATT--

Top scoring motifs for Assay 13602

Protein ID: pTH14299.1 Gene: DACH2.DBD Domain: Unknown Flag: Reject Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

ATGTATAA	0.36374	ATGTATAA--
ACTAAGGA	0.35526	--ACTAAGGA--
TATAAATA	0.35392	---TATAAATA
AATAATAG	0.35081	--CTATTATT--
CCAAAACA	0.34914	--CCAAAACA--
ATTATCTG	0.34856	-ATTATCTG--
ACTATCAC	0.34838	-ACTATCAC--
ACTATTAT	0.34834	-ACTATTAT--
AATATCAC	0.34448	-AATATCAC--
GAGTGCTA	0.34448	---TAGCACTC

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

ATGTTAAC	3.95362	----ATGTTAAC
TATAAATA	3.94459	--TATAAATA--
ACTATTAT	3.85745	----ATAATAGT
CGTAATAA	3.72237	CGTAATAA----
ATACATGA	3.62058	--ATACATGA--
TAATAATA	3.57360	--TAATAATA--
ATACATTA	3.53723	--ATACATTA--
ATTAAGTG	3.51752	--ATTAAGTG--
AACATGCA	3.47254	---AACATGCA--
AATAGTTC	3.43942	---AATAGTTC--