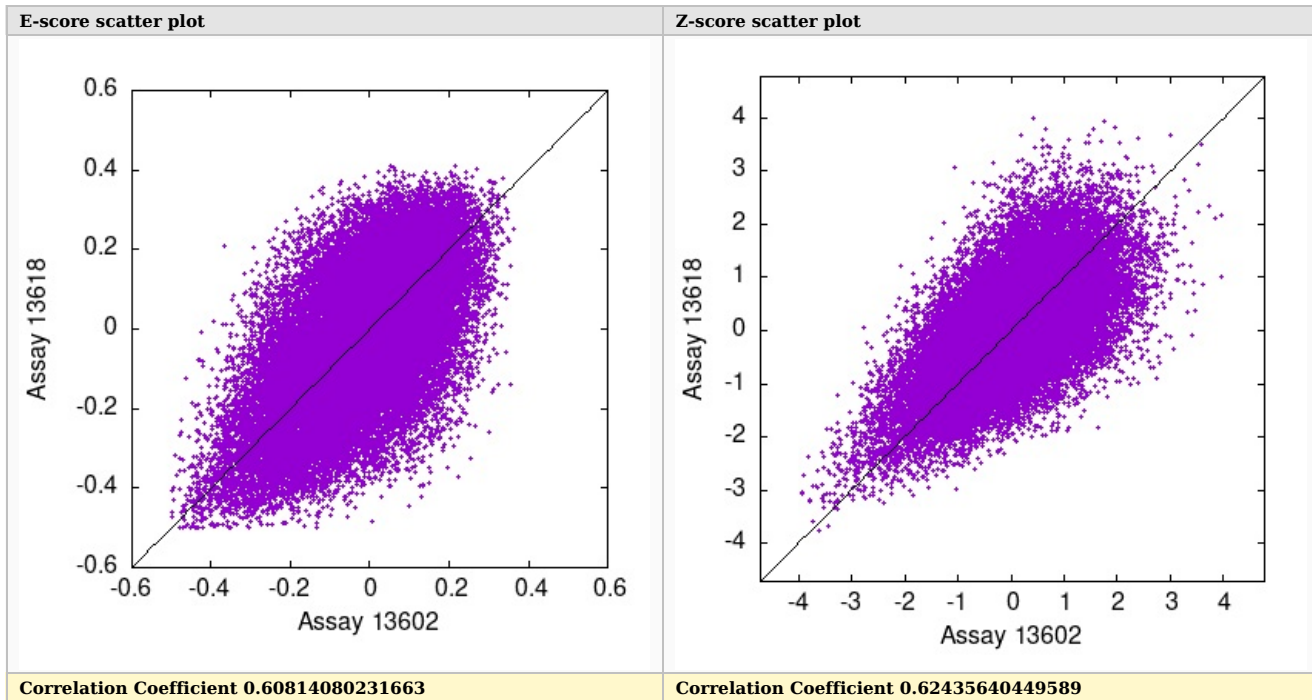


QC report



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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
ATGTATAA	0.36374	ATGTATAA---	ATGTTAAC	3.95362	----ATGTTAAC
ACTAAGGA	0.35526	-ACTAAGGA--	TATAAATA	3.94459	--TATAAATA--
TATAAATA	0.35392	---TATAAATA	ACTATTAT	3.85745	---ATAATAGT
AATAATAG	0.35081	--CTATTATT-	CGTAATAA	3.72237	CGTAATAA----
CCAAAACA	0.34914	--CCAAAACA-	ATACATGA	3.62058	--ATACATGA--
ATTATCTG	0.34856	-ATTATCTG--	TAATAATA	3.57360	--TAATAATA--
ACTATCAC	0.34838	-ACTATCAC--	ATACATTA	3.53723	--ATACATTA--
ACTATTAT	0.34834	-ACTATTAT--	ATTAAGTG	3.51752	---ATTAAGTG-
AATATCAC	0.34448	-AATATCAC--	AACATGCA	3.47254	--AACATGCA--
GAGTGCTA	0.34448	---TAGCACTC	AATAGTTC	3.43942	--AATAGTTC-

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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
AATAAATT	0.40900	-AATTTATT-	ATTATTTTC	3.98078	--ATTATTTTC-
GATTTATA	0.40896	-GATTTATA-	CTATATTA	3.93379	--CTATATTA--
GATATGTA	0.40639	-GATATGTA-	GAAATTTTC	3.82215	--GAAATTTTC
AATTTATA	0.40235	-AATTTATA-	GATATGTA	3.80010	-GATATGTA--
AATTTAAA	0.40052	-AATTTAAA-	CTGTATTA	3.77616	TAATACAG---
ATTATTTTC	0.39732	GAAATAAT--	AATTTAAA	3.68574	-TTTAAATT--
ATTTAAAT	0.39498	--ATTTAAAT	ATAATGTA	3.67093	--TACATTAT
ATAAATTA	0.39472	TAATTTAT--	GATTTATA	3.66407	-TATAAATC--
TATCAACA	0.39444	-TGTTGATA-	TATTAATA	3.61840	-TATTAATA--
ATATTAAG	0.39351	--CTTAATAT	AATAAATT	3.59110	-AATAAATT--