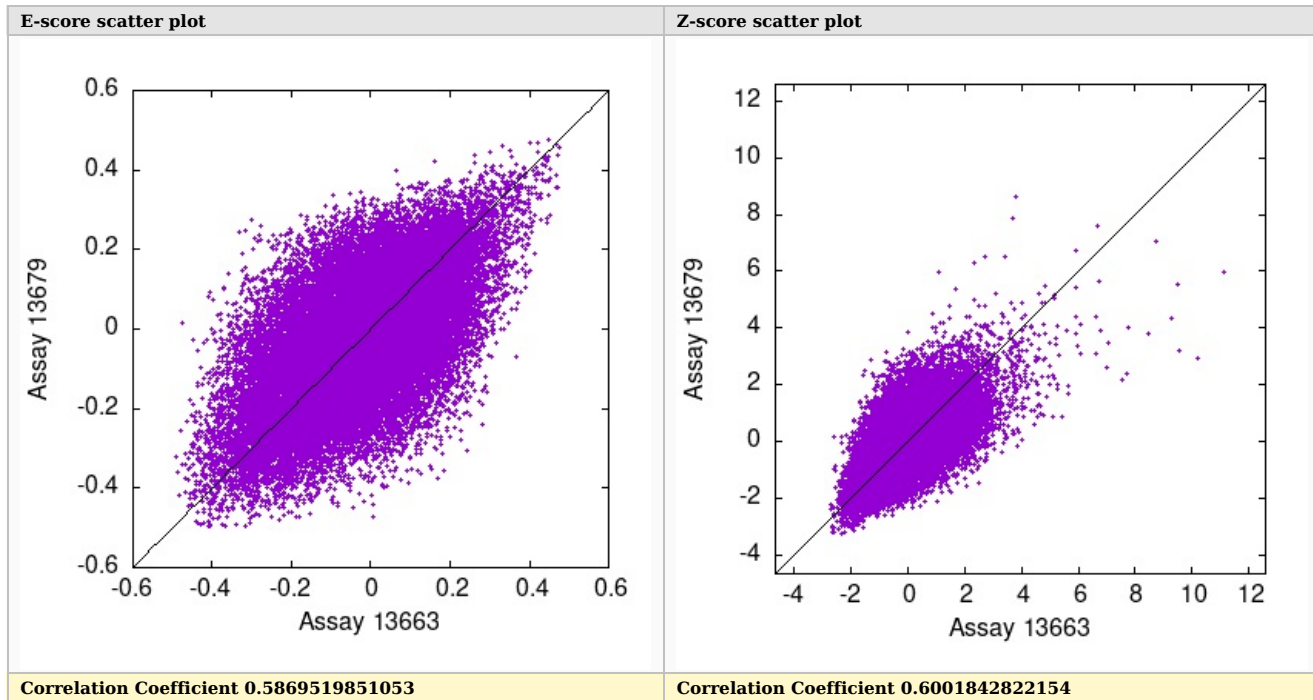


## QC report



### Top scoring motifs for Assay 13663

Protein ID: pTH14272.1 Gene: MSANTD4.FL Domain: Myb\_DNA-binding Flag: Pass\_matched\_pair 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
TACTACTA	0.47493	-TAGTAGTA---	TACTACTA	11.14040	TAGTAGTA---
GTGTAGTA	0.47270	-GTGTAGTA---	ATGAGTAG	10.18475	-ATGAGTAG--
AGTAGTAA	0.47124	--AGTAGTAA--	AGTAGTAA	9.52683	-AGTAGTAA--
TGTAGTAA	0.46960	--TGTAGTAA--	GTGTAGTA	9.47186	GTGTAGTA---
GTAGTAAA	0.46806	---GTAGTAAA--	GTAGTAAA	9.28286	--GTAGTAAA--
ATGAGTAG	0.46639	--ATGAGTAG--	TGTAGTAA	8.74612	-TGTAGTAA--
ACTACTCA	0.45285	TGAGTAGT----	GAGTAAC	8.44185	--GTAGTAAC-
GAGTAAC	0.45247	---GTAGTAAC-	TACTACGA	7.75126	TCGTAGTA---
ATTACTA	0.45136	---TAGTAAAT	ATTACTA	7.72856	---TAGTAAAT
ACTACTAT	0.44997	ATAGTAGT----	GATGAGTA	7.54830	GATGAGTA---

### Top scoring motifs for Assay 13679

Protein ID: pTH14272.2 Gene: MSANTD4.FL Domain: Myb\_DNA-binding Flag: Pass\_matched\_pair 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
ACTACTAC	0.47514	---ACTACTAC-	TACTAGTA	8.62524	--TACTAGTA-
CTGTAGTA	0.46669	--TACTACAG--	CTGTAGTA	7.87323	--TACTACAG--
CTTACTAC	0.46650	CTTACTAC----	ACTACTAC	7.57222	---ACTACTAC
TGTAGTAA	0.46585	-TTACTACA---	TGTAGTAA	7.06375	-TTACTACA--
TACTAGTA	0.46142	--TACTAGTA--	ATGTAGTA	6.71759	--TACTACAT-
TACTACTA	0.45781	--TACTACTA--	ACTACTAG	6.53245	---ACTACTAG
ACTACTAG	0.44834	---ACTACTAG-	TAGTAGCA	6.49744	--TGCTACTA-
ATGTAGTA	0.44685	--TACTACAT--	GTAATTAC	6.31956	GTAATTAC---
ACTAGTAG	0.44371	----CTACTAGT	TACTACTA	5.99335	--TACTACTA-
TGAGTAAA	0.43703	---TTTACTCA-	GTGTACAC	5.96874	--GTGTACAC-