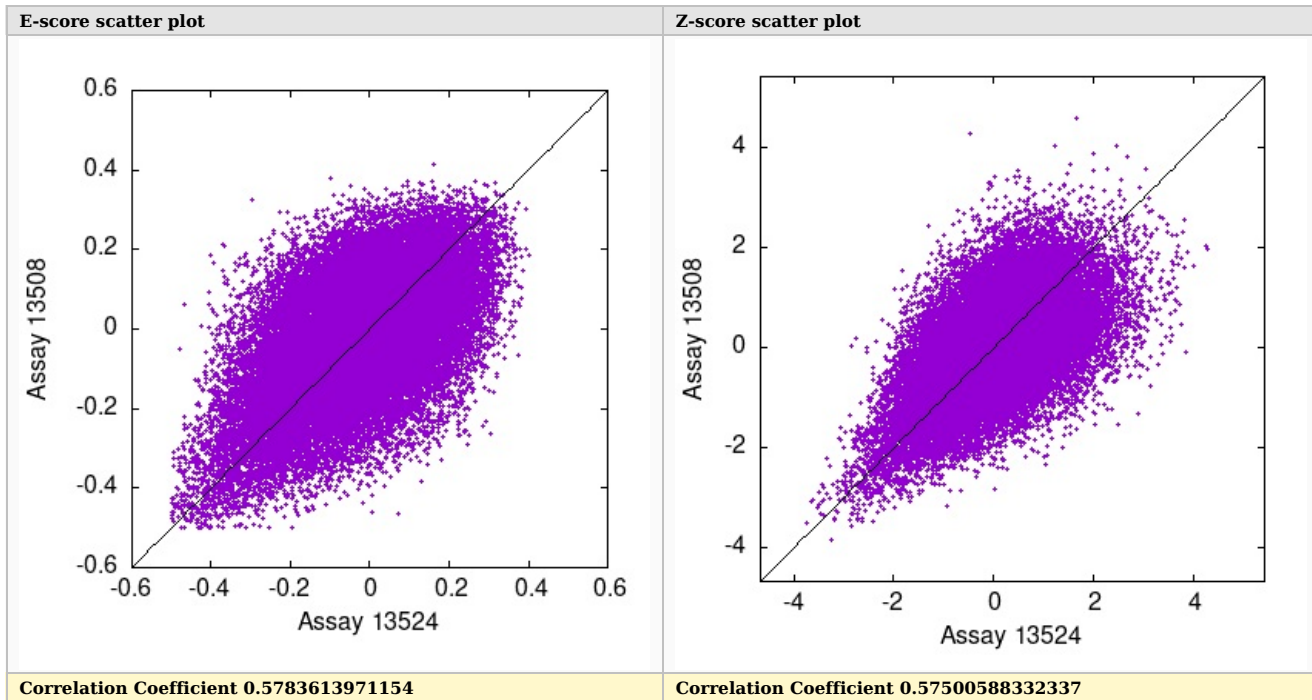


## QC report



## Top scoring motifs for Assay 13524

Protein ID: pTH13953.2 Gene: ATMIN.DBD Domain: zf-C2H2 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
GATATGTA	0.40239	----TACATATC	TATTTGCA	4.28604	TGCAAAATA-----
ACAAAATA	0.39254	--ACAAAATA--	ACAAAATA	4.26013	-----TATTTTGT-
ATGTAGTA	0.39249	-TACTACAT---	GTACAATA	4.01885	-----GTACAATA--
AATAAATT	0.38777	--AATAAATT--	AATACATT	3.88079	-----AATACATT--
TATCAACA	0.38364	TATCAACA----	ACCACTAA	3.84938	----ACCACTAA---
ATAATAGA	0.38117	--ATAATAGA--	ATGTAGTA	3.83061	---TACTACAT----
TGACAAAA	0.38075	TGACAAAA----	GTAATACA	3.82250	--GTAATACA-----
AGAGATAT	0.38013	---AGAGATAT--	ACATTACT	3.78291	-----ACATTACT--
TATTTGCA	0.37787	--TGCAATAA--	AACACTTA	3.75513	----AACACTTA---
ATATTAGC	0.37590	----ATATTAGC	AAAATAAA	3.74851	--AAAATAAA-----

## Top scoring motifs for Assay 13508

Protein ID: pTH13953.1 Gene: ATMIN.DBD Domain: zf-C2H2 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
TATAAATA	0.41382	---TATAAATA	TATAAATA	4.58360	TATTTATA---
ATTTACTA	0.38050	--TAGTAAAT--	ATTTACTA	4.27085	-ATTTACTA--
ATGTATAA	0.37306	--TTATACAT--	ATAATGCA	4.04240	--TGCAATTAT--
ACTACTAT	0.37265	--ATAGTAGT--	AATAATAA	4.03150	---TTATTATT
AACGAAAT	0.37253	--AACGAAAT--	ACATACAA	3.89224	-ACATACAA--
ATATAAAT	0.36695	--ATATAAAT--	AACATACA	3.82662	AACATACA---
TAACGAAA	0.36681	-TAACGAAA--	AGAAAAGA	3.66445	TCTTTTCT---
AAACATAA	0.36577	AAACATAA---	ATATAGTA	3.57793	-ATATAGTA--
ACATACAA	0.36483	--ACATACAA--	ACGTAATA	3.57752	-ACGTAATA--
ATATATCA	0.36392	--ATATATCA--	ATATATCA	3.56020	-ATATATCA--