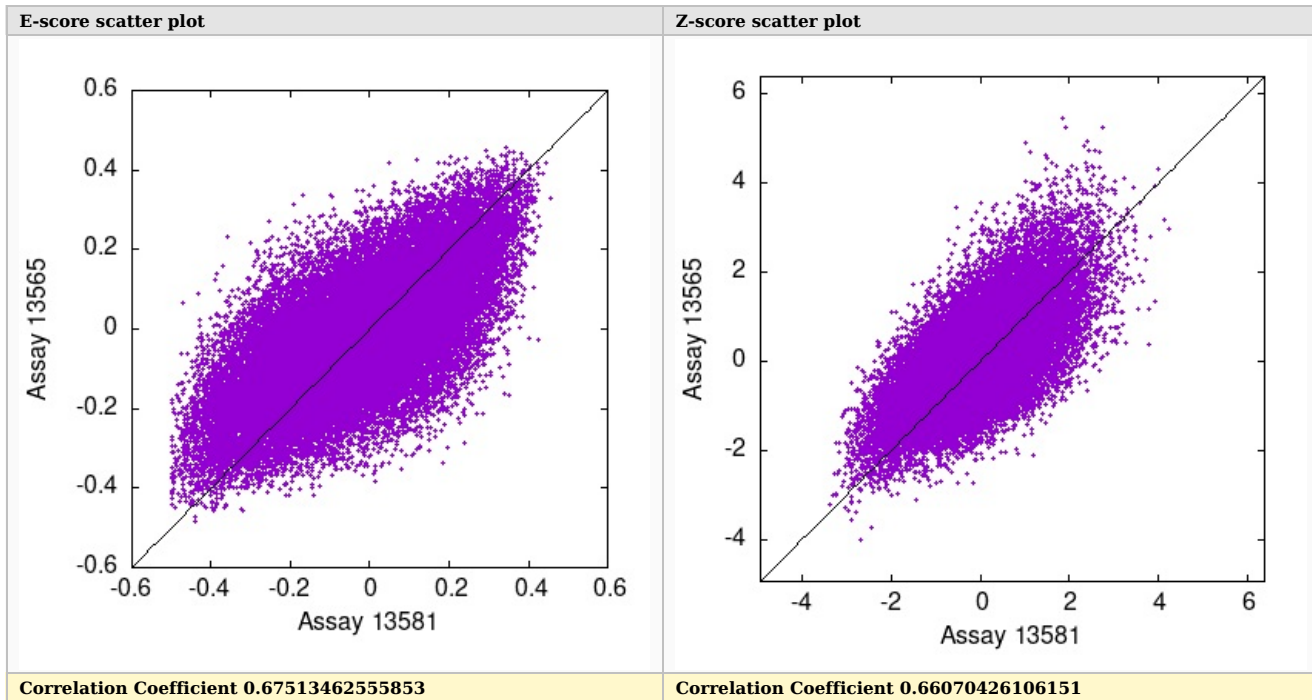


## QC report



### Top scoring motifs for Assay 13581

Protein ID: pTH13957.2 Gene: CHAMP1.DBD Domain: zf-C2H2 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
AAATAATAA	0.45602	--AATAATAA--	AATAATAA	4.25128	--AATAATAA--
ACTAATTA	0.44618	--ACTAATTA--	AGTAATTA	4.13176	--AGTAATTA--
ACTATAAT	0.43729	---ACTATAAT	ACTAATTA	4.00186	--ACTAATTA--
ATATTAAG	0.42970	---ATATTAAG	ACTATAAT	3.92271	---ACTATAAT
ACTAACAT	0.42635	--ACTAACAT--	ATATCACA	3.91826	---ATATCACA
TAATAATA	0.42584	-TAATAATA--	ATATAACA	3.82133	-ATATAACA--
ATATAACA	0.42535	-ATATAACA--	ATTATTTC	3.78685	GAAATAAT---
ATTATTTC	0.42493	GAAATAAT---	ATATACAA	3.78323	--ATATACAA--
AGTAATTA	0.42441	--AGTAATTA--	ATATTAAG	3.72001	--ATATTAAG--
AAATAATA	0.42221	-AAATAATA--	ATTAATCA	3.63989	--ATTAATCA--

### Top scoring motifs for Assay 13565

Protein ID: pTH13957.1 Gene: CHAMP1.DBD Domain: zf-C2H2 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
ATAATATA	0.45767	---TATATTAT-	ACTAATAA	5.45107	TTATTAGT----
ACTAATAA	0.44891	TTATTAGT----	GTTAGTTA	5.26544	--GTTAGTTA--
AACTAACG	0.44567	-CGTTAGTT---	ATAATATA	5.25693	---TATATTAT-
GTTAGTTA	0.44439	--GTTAGTTA--	AACTAACG	4.94130	-CGTTAGTT---
AACTAACT	0.44214	-AGTTAGTT---	ACTAACGG	4.92239	CCGTTAGT----
AAAGTAAT	0.43872	--ATTACTTT--	ATATGTTA	4.85238	--ATATGTTA--
GTTAACTA	0.43644	----TAGTTAAC	AAGTAATT	4.74237	-AATTACTT---
ATTAGTTG	0.43482	--ATTAGTTG--	GTTAACTA	4.73338	----TAGTTAAC
AAGTAACA	0.43339	-TGTTACTT---	AACTAACT	4.70248	-AGTTAGTT---
AACTAATG	0.43212	-CATTAGTT---	CGTTAGTC	4.68527	-CGTTAGTC---