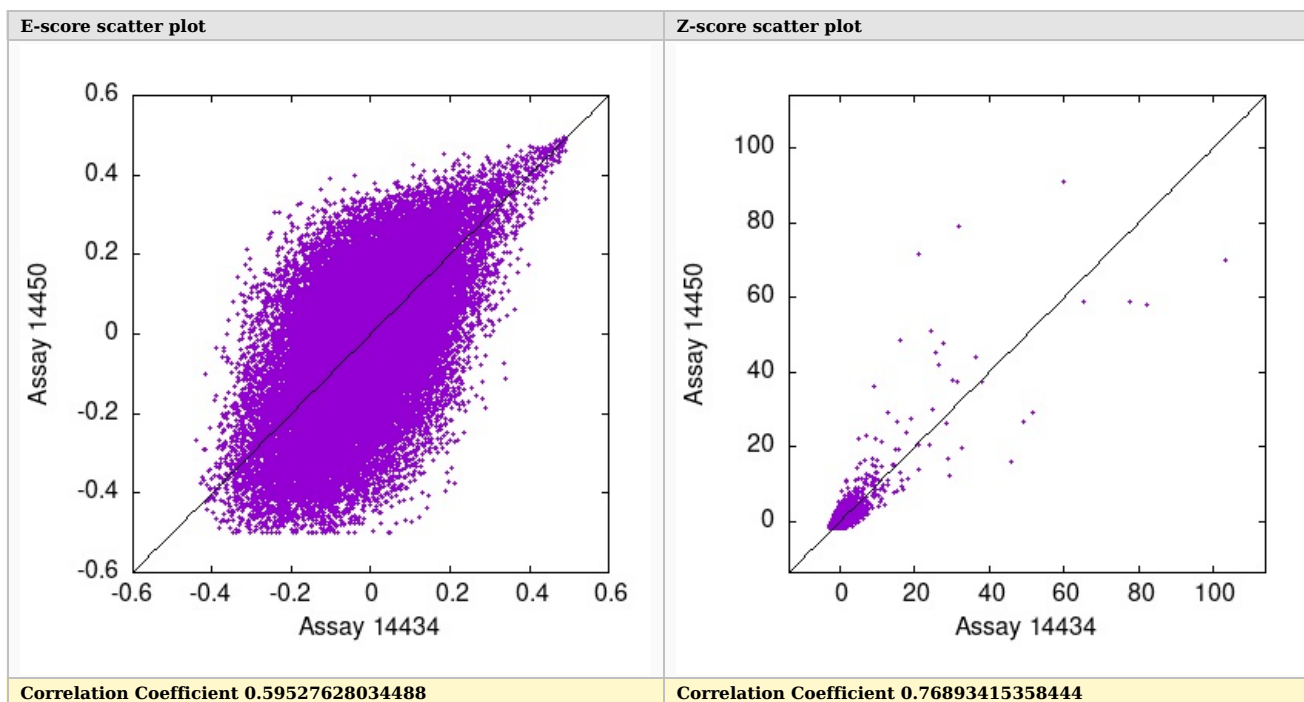




QC report





Top scoring motifs for Assay 14434

Protein ID: pTH15561.1 Gene: SOX5 Domain: HMG_box Flag: Pass matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:	Reverse:	
		
Top 10	Scores	Alignment
ATTGTTAA	0.49254	---ATTGTTAA
AACAATAG	0.49253	-CTATTGTT--
ATAACAAT	0.49231	---ATTGTTAT
ACAATTGT	0.49118	ACAATTGT---
AGAACAAT	0.49101	---ATTGTTCT
AATTGTTA	0.49093	--AATTGTTA-
AACAATTG	0.49063	--CAATTGTT--
GACAATA	0.49046	--TATTGTTC-
TAACAATA	0.48933	--TATTGTTA-
AACAATG	0.48908	--CATTGTTT-



8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
AACAATAG	103.38600	-CTATTGTT--	
ATTGTTAA	82.31556	---ATTGTTAA	
AGAACAAT	77.51613	--ATTGTTCT	
TAACAATA	65.08125	--TATTGTTA-	
ATAACAAT	60.05820	---ATTGTTAT	
AAACAATG	51.56551	--CATTGTTT-	
AACAATTG	49.16070	--CAATTGTT--	
ACAATTGT	45.86954	ACAATTGT---	
AATTGTTC	38.19368	--AATTGTTC-	
AATTGTTA	36.20242	--AATTGTTA-	



Top scoring motifs for Assay 14450

Protein ID: pTH15561.2 Gene: SOX5 Domain: HMG_box Flag: Pass matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:	Reverse:	
		
Top 10	Scores	Alignment
AACAATGG	0.49365	- - AACAAATGG
ATAACAAT	0.49239	ATAACAAT - -
AACAATAG	0.49207	- - AACAAATAG
AGAACAAT	0.49144	AGAACAAT - -
TAACAATA	0.48969	- TAACAATA -
CATTGTTA	0.48963	- TAACAATG -
CATTGTTC	0.48953	- GAACAATG -
AACAATTA	0.48923	- - AACAAATTA
AATTGTTA	0.48917	- TAACAATT -
GAACAATA	0.48897	- GAACAATA -

8 mer Z-scores for probeset 'all'

Forward:	Reverse:	
		
Top 10	Scores	Alignment
ATAACAAT	91.09606	--ATTGTTAT
AACAATGG	78.88061	CCATTGTT--
CATTGTTC	71.65770	-CATTGTTC-
AACAATAG	69.93455	CTATTGTT--
AGAACAAT	58.90226	--ATTGTTCT
TAACAATA	58.75110	-TATTGTTA-
ATTGTTAA	58.22485	--ATTGTTAA
CATTGTTA	50.94506	-CATTGTTA-
AACAATGT	48.37368	ACATTGTT--
GAACAATA	47.72583	-TATTGTTC-