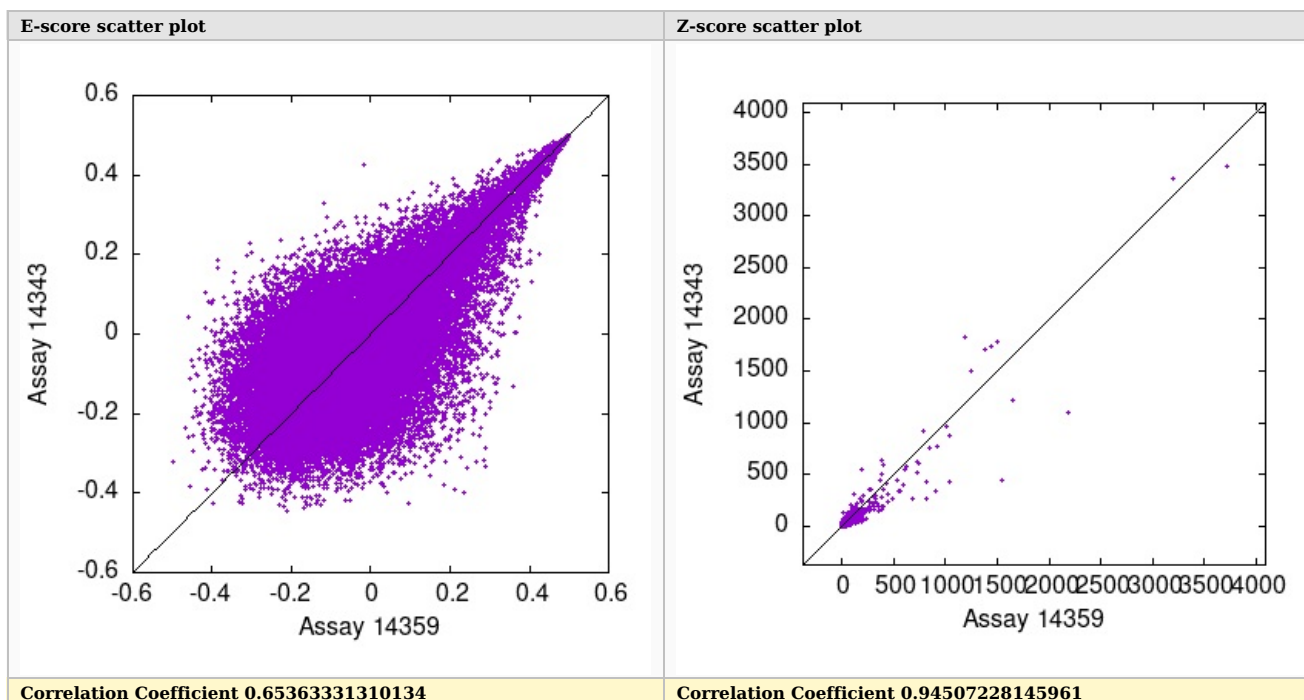


QC report



Top scoring motifs for Assay 14359

Protein ID: pTH15542.2 Gene: GCM1 Domain: GCM 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
ACCCGCAT	0.49928	-ACCCGCAT-	ACCCGCAT	3,717.42911	-ACCCGCAT-
ATGCGGGC	0.49883	-GCCCGCAT-	ATGCGGGC	3,189.91932	-GCCCGCAT-
CATGCGGG	0.49715	--CCCGCATG	ACCCGTAT	2,182.98721	-ACCCGTAT-
ACCCGTAT	0.49707	-ACCCGTAT-	CATGCGGG	1,653.81818	--CCCGCATG
CCCGCATC	0.49567	--CCCGCATC	CACCCGCA	1,552.19430	CACCCGCA--
AATGCGGG	0.49548	--CCCGCATT	CCCGCATC	1,501.46445	--CCCGCATC
TACCCGCA	0.49475	TACCCGCA--	ATACGGGC	1,445.96201	-GCCCGTAT-
CACCCGCA	0.49444	CACCCGCA--	AATGCGGG	1,381.99643	--CCCGCATT
CCCGCATA	0.49439	--CCCGCATA	CCCGCATA	1,242.82090	--CCCGCATA
ATACGGGC	0.49292	-GCCCGTAT-	TACCCGCA	1,185.32707	TACCCGCA--

Top scoring motifs for Assay 14343

Protein ID: pTH15542.1 Gene: GCM1 Domain: GCM 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
ACCCGCAT	0.49949	-ACCCGCAT-	ACCCGCAT	3,484.83414	-ACCCGCAT-
ATGCGGGC	0.49916	-GCCCGCAT-	ATGCGGGC	3,365.91227	-GCCCGCAT-
TACCCGCA	0.49820	TACCCGCA--	TACCCGCA	1,827.80248	TACCCGCA--
CCCGCATC	0.49764	--CCCGCATC	CCCGCATC	1,788.24019	--CCCGCATC
AATGCGGG	0.49711	--CCCGCATT	ATACGGGC	1,736.88696	-GCCCGTAT-
CCCGCATA	0.49693	--CCCGCATA	AATGCGGG	1,711.54305	--CCCGCATT
ATACGGGC	0.49573	-GCCCGTAT-	CCCGCATA	1,498.97053	--CCCGCATA
CATGCGGG	0.49569	--CCCGCATG	CATGCGGG	1,216.51835	--CCCGCATG
ACCCGTAT	0.49485	-ACCCGTAT-	ACCCGTAT	1,103.45295	-ACCCGTAT-
GACCCGCA	0.49330	GACCCGCA--	ATGCGGGA	964.55728	-TCCCGCAT-