Only showing significant correlations (<0.05)

	Only sh	owing s	ignifican	t correla	ations (<	(0.05)									
Epidish_DWLS.Vanderbilt.scRNAseq -	0.95	0.96		0.86	0.84	0.98	0.93	0.99	0.98	0.99	0.96	0.99	0.9	0.97	0.96
DWLS_DWLS.Vanderbilt.scRNAseq -		0.95		0.9	0.84	0.95	0.92	0.99	0.97	0.98	0.95	0.99	0.92	0.94	0.95
CBSX_DWLS.Vanderbilt.scRNAseq -	0.93	0.95		0.85	0.83	0.95	0.88	0.99	0.98	0.99	0.97	1	0.9	0.91	0.92
DWLS_CBSX.Vanderbilt.scRNAseq -		0.93		0.86	0.75	0.98	0.91	0.99	0.99	0.98	0.9	0.99	0.86	0.95	0.74
CBSX_CBSX.Vanderbilt.scRNAseq -		0.89		0.79	0.84	0.99	0.91	0.98	0.99	0.98	0.91	0.99	0.85	0.93	0.75
DWLS_MOMF.Vanderbilt.scRNAseq -		0.98		0.8	0.63	0.99	0.91	0.95	0.98	0.96	0.8	0.99	0.89	0.93	0.93
Epidish_CBSX.Vanderbilt.scRNAseq		0.89		0.79	0.68	0.98	0.92	0.98	0.98	0.98	0.88	1	0.88	0.96	0.74
MuSic_Vanderbilt	0.00	0.99		0.84	0.69	0.98	0.99	0.07	0.96	0.96	0.89	0.99	0.81	0.87	0.00
BayesPrism_Vanderbilt	0.00	0.97		0.81	0.52	0.99	0.88	0.97	0.99	0.98	0.83	1	0.9	0.92	0.86
DeconRNASeq_DWLS.Vanderbilt.scRNAseq		0.83		0.72	0.7	0.91	0.86	0.92	0.97	0.96	0.95	0.99	0.73	0.86	0.72
Epidish_BSeqSC.Vanderbilt.scRNAseq - DWLS_BSeqSC.Vanderbilt.scRNAseq -	0.00	0.91		0.78	0.54	0.97	0.97	0.86	0.98	0.96	0.81	0.99	0.83	0.85	0.61
CBSX_MOMF.Vanderbilt.scRNAseq -	0.00	0.94		0.77	0.54	0.97 0.96	0.97	0.88	0.96	0.96	0.81 0.79	0.99	0.83	0.87 0.86	0.81
DeconRNASeq_CBSX.Vanderbilt.scRNAseq		0.83		0.9	0.64	0.90	0.84	0.85	0.93	0.79	0.79	0.04	0.73	0.80	0.62
CBSX_CBSX.Melanoma.scRNAseq		0.74		0.54	0.76	0.94	0.04	0.03	0.90	0.91	0.87	0.90	0.77	0.77	0.02
DWLS_CBSX.HNSCC.scRNAseq	0.0	0.55		0.62	0.76	0.98		0.98			0.07				
DeconRNASeq_MOMF.Vanderbilt.scRNAseq	0.0	0.97		0.02	0.00	0.87	0.91	0.88	0.93	0.9	0.75	0.99	0.82		
SCDC_Vanderbilt		0.93		0.81	0.54	0.92	0.98	0.93	0.91	0.92	0.89	0.97	0.85	0.73	
Epidish_MOMF.Vanderbilt.scRNAseq		0.8		0.81	0.01	0.86	0.69	0.94	0.97	0.73	0.81	0.79	0.7	0.82	0.84
Epidish_CBSX.HNSCC.scRNAseq		0.54		0.58	0.86	0.97	0.00	0.97	0.0.	00	0.0.			0.02	0.0.
CBSX_BSeqSC.Vanderbilt.scRNAseq		0.89		0.76		0.89	0.84	0.8	0.92	0.98	0.81	0.99	0.82	0.58	
Bisque_Vanderbilt -		0.95		0.72	0.82	0.84	0.75	0.91	0.97	0.91	0.79	0.99	0.69	0.53	0.76
DeconRNASeq_TIL10	0.76	0.7		0.68	0.87						0.81				0.74
CPM_Vanderbilt -	0.74	0.85		0.77	0.62		0.91	0.83	0.87	0.95	0.79	0.95	0.82	0.84	0.6
DWLS_CBSX.Melanoma.scRNAseq -	0.73	0.86		0.69	0.69	0.95					0.82				
Epidish_BPRNACan	0.72	0.96	0.54		0.66						0.96				
Epidish_TIL10 -	0.72	0.86			0.84						0.71				0.71
Quantiseq -		0.78			0.88						0.71				
Epidish_CBSX.Melanoma.scRNAseq		0.86		0.62	0.66	0.95					0.82				
CBSX_BPRNACan_	~	0.95			0.69	0.00	0.00	0.07	0.70	0.00	0.97	0.00		0.00	
DeconRNASeq_BSeqSC.Vanderbilt.scRNAseq - DWLS_BPRNACan		0.85	0.54		0.58	0.86	0.86	0.67	0.79	0.96	0.79	0.98		0.68	
Epidish_BPRNACan3DProMet	0.69	0.96 0.95	0.51		0.69 0.65						0.96 0.96				
CBSX_BPRNACanProMet	0.69	0.95			0.65						0.96				
CBSX_CBSX.HNSCC.scRNAseq		0.55		0.63	0.83	0.98		0.99			0.90				
Epidish_BPRNACanProMet		0.96		0.03	0.7	0.30		0.55	l		0.96				
DWLS_LM22 -		0.50		_	0.54						0.50	0.97	0.54		0.63
DWLS_BPRNACanProMet		0.96		_	0.72						0.96	0.07	0.01		0.00
DWLS_TIL10		0.86			0.88						0.63				
Epidish_LM22 -												0.97	0.58		0.63
CBSX_BPRNACan3DProMet	0.65	0.91			0.65						0.96				
DWLS_BPRNACan3DProMet	0.65	0.94			0.69						0.96				
CBSX_LM22	0.64				0.52							0.92	0.52		0.57
CBSX_TIL10	0.00	0.78			0.9						0.8				0.57
AutogeneS_Vanderbilt -	0	0.6		0.64	0.71	0.87	0.99	0.75	0.89	0.84		0.85	0.72	0.59	
DeconRNASeq_LM22					0.77							0.86	0.63		
DeconRNASeq_CCLE.TIL10		0.69		0.72	0.9						0.81				0.76
DeconRNASeq_BPRNACanProMet	0.56	0.91			0.72						0.77				
DeconRNASeq_BPRNACan3DProMet_	0.56	0.91			0.74						0.75				
CBSX_CCLE.TIL10 - DeconRNASeq_BPRNACan		0.9			0.85						0.67				
DWLS_CCLE.TIL10		0.95			0.73						0.65				
DeconRNASeq_CBSX.HNSCC.scRNAseq		0.91			0.88	0.95		0.56			0.65				
Epidish_CCLE.TIL10		0.59			0.73	0.95		0.50			0.69				
DeconRNASeq_CBSX.NSCLC.PBMCs.scRNAseq		0.82			0.76						0.09				
DeconRNASeq_CBSX.Melanoma.scRNAseq	• • • •	0.83	-0.7		0.76	0.62					0.72				
CBSX_CBSX.NSCLC.PBMCs.scRNAseq		0.03	0.7		0.75	0.02					0.12				
DWLS_CBSX.NSCLC.PBMCs.scRNAseq		0.73			0.76										
Epidish_CBSX.NSCLC.PBMCs.scRNAseq		0.8			0.72										
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pearson's Correlation 1.0 0.5 0.0 -0.5 -1.0