STRING	hESC	0.9632 0.9607	0.9567	0.9380	0.9421	0.8593	0.8378	0.6837	0.5996		
	hHEP	0.9701 0.9654	0.9601	0.9191	0.9275	0.8354	0.7898	0.6400	0.5200		
	mDC	0.9747 0.9716	0.9720	0.9227	0.9379	0.8877	0.7846	0.6457	0.5056		
	mESC	0.9762 0.9722	0.9722	0.9196	0.9362	0.8584	0.8453	0.5855	0.5338		- 0.9
	mHSC-E	0.9455 0.9468	0.9368	0.9519	0.9337	0.8906	0.6002	0.6996	0.5647		
	mHSC-GM	0.9316 0.9392	0.9074	0.9384	0.9305	0.9024	0.7596	0.7311	0.7058		
	mHSC-L	0.8307 0.8283	0.7775	0.8146	0.8093	0.6575	0.6942	0.6361	0.6729		
											- 0.8
Cell-type- specific ChIP-seq	hESC	0.9842 0.9840	0.9827	0.9557	0.9486	0.9677	0.6746	0.5088	0.5701		
	hHEP	0.9858 0.9857	0.9847	0.9768	0.9534	0.9829	0.6538	0.4735	0.5854		
	mDC	0.9809 0.9815	0.9815	0.8879	0.8622	0.9801	0.6635	0.5126	0.3761		
	mESC	0.9847 0.9842	0.9824	0.9565	0.9075	0.9788	0.6970	0.5018	0.6145		- 0.7
	mHSC-E	0.9775 0.9770	0.9748	0.9519	0.8927	0.9546	0.6408	0.4715	0.5020		
	mHSC-GM	0.9726 0.9730	0.9663	0.9475	0.9244	0.9409	0.7042	0.5238	0.4395		
	mHSC-L	0.9483 0.9482	0.9420	0.9203	0.8978	0.9367	0.7259	0.5423	0.5447		
											- 0.6
Non-specifi ChIP-seq	hESC	0.9139 0.9125	0.9089	0.7745	0.7855	0.7315	0.6644	0.5481	0.5174		
	hHEP	0.9210 0.9184	0.9156	0.8179	0.8357	0.8424	0.7358	0.5686	0.5283		
	mDC	0.9357 0.9337	0.9312	0.8516	0.8608	0.8107	0.7101	0.6038	0.5134		
	mESC	0.9374 0.9327	0.9263	0.8520	0.8817	0.6055	0.7421	0.5678	0.5426		- 0.5
	mHSC-E	0.9142 0.9135	0.8984	0.7982	0.8043	0.8268	0.7179	0.6222	0.5947		0.5
	mHSC-GM	0.9187 0.9225	0.9137	0.7928	0.7698	0.8705	0.6361	0.6404	0.6276		
	mHSC-L	0.9028 0.8959	0.8989	0.7994	0.6841	0.8562	0.6438	0.6490	0.5982		
											- 0.4
LOF/GOF	mESC	0.9914 0.9913							0.6469		0.4
		CRITICO CRITICO CO	GRINGERE (DEED ED)	GRANTINX	CHATLUX	GENELINY	GNE	8CC	'u		
		GRACE GRACE GRACE CONTROL OF THE STORY	(Jeeg	Ø [*]		G*					