STRING	hESC	0.9620	0.9594	0.9475	0.9221	0.9185	0.8765	0.7953	0.6867	0.5974		
	hHEP	0.9647	0.9603	0.9589	0.9038	0.9166	0.8340	0.7918	0.6442	0.5130		
	mDC	0.9729	0.9714	0.9706	0.9162	0.9371	0.8875	0.7361	0.6195	0.4868		
	mESC	0.9777	0.9741	0.9744	0.9267	0.9458	0.8789	0.8071	0.6035	0.5485		- 0.9
	mHSC-E	0.9498	0.9482	0.9221	0.9294	0.9335	0.8891	0.8307	0.7232	0.5938		0.9
	mHSC-GM	0.9319	0.9325	0.9223	0.9252	0.9135	0.9044	0.8169	0.7637	0.7000		
	mHSC-L	0.8378	0.8329	0.8456	0.8920	0.8192	0.8641	0.6990	0.6576	0.6406		
Cell-type- specific ChIP-seq	hESC	0.9854	0.9857	0.9845	0.9756	0.9651	0.9782	0.7135	0.5060	0.5963		- 0.8
	hHEP	0.9864	0.9862	0.9849	0.9789	0.9622	0.9853	0.7276	0.4802	0.5915		
	mDC	0.9820	0.9807	0.9839	0.9244	0.9226	0.9759	0.6668	0.5217	0.3961		- 0.7
	mESC	0.9871	0.9868	0.9846	0.9619	0.9057	0.9781	0.6829	0.4947	0.6378		
	mHSC-E	0.9807	0.9809	0.9790	0.9619	0.9039	0.9665	0.6530	0.4674	0.4771		
	mHSC-GM	0.9808	0.9809	0.9769	0.9619	0.9314	0.9651	0.7234	0.5231	0.4661		
	mHSC-L	0.9512	0.9500	0.9422	0.9317	0.9121	0.9338	0.7364	0.5315	0.5804		
Non-specifi ChIP-seq	hESC	0.9293	0.9269	0.9254	0.7908	0.7843	0.8547	0.7012	0.5470	0.5223		- 0.6
	hHEP	0.9362	0.9356	0.9282	0.8359	0.8621	0.8312	0.7225	0.5555	0.5558		
	mDC	0.9412	0.9396	0.9350	0.8645	0.8881	0.8193	0.7061	0.6037	0.5216		
	fic mESC	0.9458	0.9427	0.9381	0.8659	0.8918	0.6906	0.7200	0.5721	0.5517		
	mHSC-E	0.9266	0.9291	0.9078	0.8332	0.8371	0.8225	0.6112	0.6163	0.6308		- 0.5
	mHSC-GM	0.9268	0.9197	0.9168	0.8330	0.8006	0.8685	0.7240	0.6578	0.6706		
	mHSC-L	0.8041	0.8050	0.8032	0.6831	0.5895	0.8225	0.5421	0.5903	0.5811		
												- 0.4
LOF/GOF	mESC	0.9896						0.7898	0.5060	0.6469		
		GAN GEORGE	iel cletol	GRING GERE	GRANLINK	CMMTLUX	<b>GENELIN</b> *	GNE	& <sub>CC</sub>	'u		
		GRIT CEPS	CKW.	O. Go	Q,	~	G.					