	hESC	0.7132	0.7065	0.6973	0.6624	0.6653	0.2100	0.1978	0.0659	0.0362	
STRING	hHEP	0.7760	0.7602	0.7438	0.6028	0.5682	0.1583	0.1588	0.0483	0.0362	
	mDC	0.8377	0.8272	0.8235	0.6935	0.6627	0.3127	0.1960	0.0545	0.0391	
	mESC	0.7679	0.7145	0.7451	0.5111	0.5457	0.1158	0.1772	0.0379	0.0291	
	mHSC-E	0.6609	0.6707	0.6328	0.7297	0.6776	0.3328	0.0429	0.0801	0.0463	- 0.8
	mHSC-GM	0.6984	0.7100	0.6360	0.7803	0.7376	0.4548	0.2051	0.2445	0.1726	
	mHSC-L	0.3318	0.3501	0.3184	0.5464	0.5395	0.0988	0.1252	0.2071	0.2174	
Cell-type- specific ChIP-seq	hESC	0.8853	0.8868	0.8792	0.8038	0.7848	0.8238	0.3218	0.2019	0.2063	
	hHEP	0.9643	0.9627	0.9635	0.9486	0.9210	0.9575	0.5165	0.4027	0.4470	- 0.6
	mDC	0.7412	0.7695	0.7507	0.6163	0.6105	0.7520	0.1473	0.0803	0.0644	0.0
	mESC	0.9609	0.9602	0.9561	0.8880	0.8282	0.9355	0.5427	0.3910	0.4618	
	mHSC-E	0.9789	0.9780	0.9747	0.9474	0.8951	0.9406	0.6922	0.5766	0.5696	
	mHSC-GM	0.9678	0.9686	0.9607	0.9381	0.9047	0.9106	0.7190	0.5375	0.4901	
	mHSC-L	0.9364	0.9378	0.9207	0.9015	0.8808	0.9141	0.7366	0.5620	0.5636	
											- 0.4
Non-specific ChIP-seq	hESC	0.2629	0.2533	0.2368	0.1801	0.1971	0.0547	0.0390	0.0202	0.0172	
	hHEP	0.2642	0.2728	0.2402	0.2065	0.2249	0.0916	0.0699	0.0222	0.0179	
	mDC	0.4489	0.4446	0.4303	0.3731	0.3616	0.1364	0.0795	0.0254	0.0203	
	ic <sub>mESC</sub>	0.3587	0.3352	0.3048	0.2545	0.2778	0.0183	0.0631	0.0236	0.0184	
	mHSC-E	0.4613	0.4551	0.4090	0.3944	0.3615	0.1903	0.1188	0.0421	0.0332	- 0.2
	mHSC-GM	0.4744	0.4838	0.4836	0.4264	0.4026	0.2197	0.1054	0.0950	0.0701	
	mHSC-L	0.4617	0.4347	0.3322	0.4152	0.3159	0.3123	0.1066	0.1537	0.1309	
LOF/GOF	mESC	0.9385								0.3148	
		Gringer Co	CRANCTLE PO	GRING GERE	GRAMINA	Charling	GENELINY	CHE	& <sub>CC</sub>	'n	