

Supplementary

(Fig.S1-S4, Table S1-S9)

Figure S1-S4

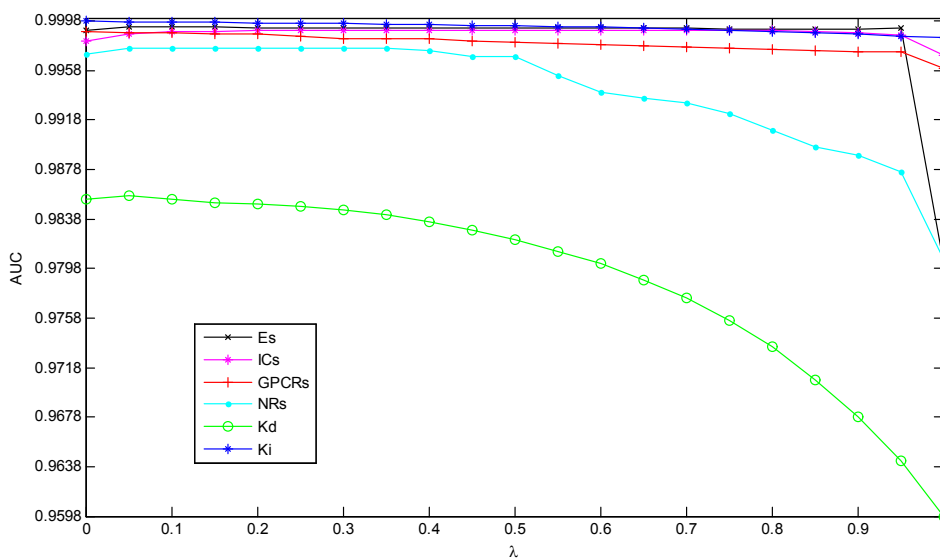


Figure S1 The relationship between the parameter λ and AUC for LPMIHN algorithm on the six binary drug-target datasets

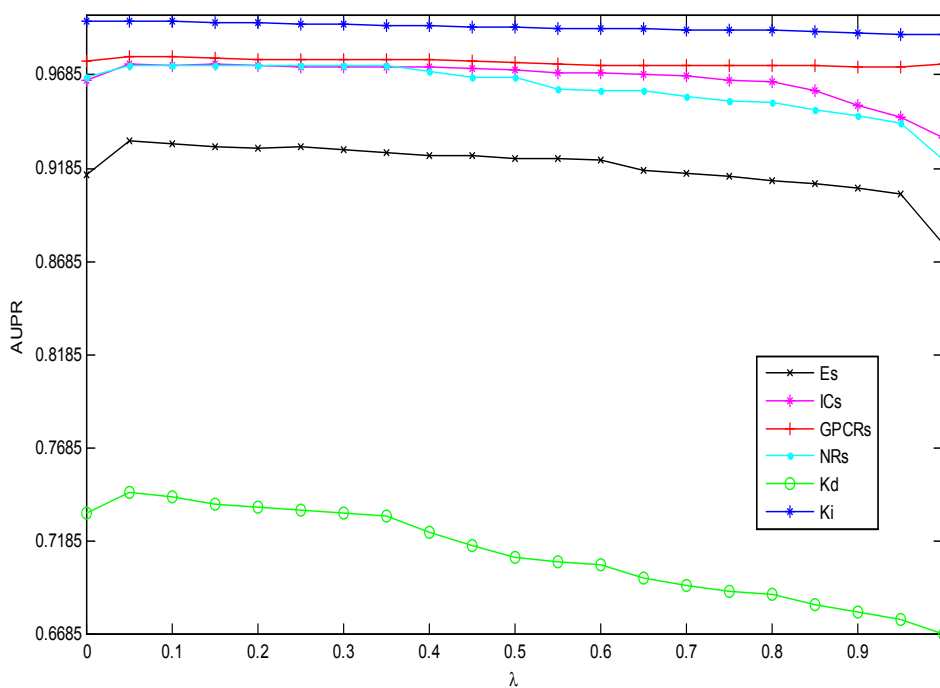


Figure S2 The relationship between the parameter λ and AUPR for LPMIHN algorithm on the six

binary drug-target datasets

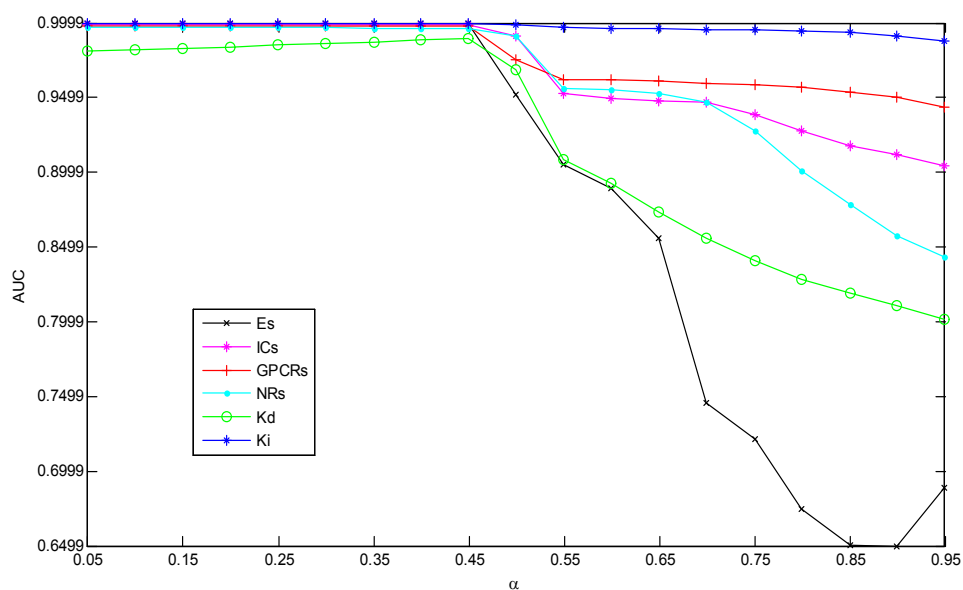


Figure S3 The relationship between the parameter α and AUC for LPMIHN algorithm on the six binary drug-target datasets

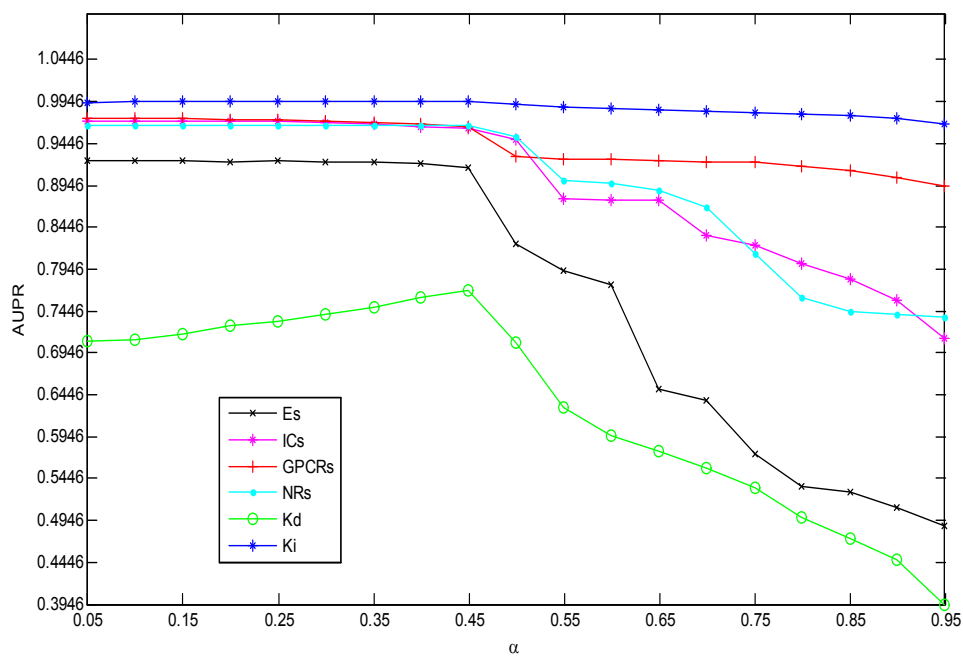


Figure S4 The relationship between the parameter α and AUPR for LPMIHN algorithm on the six binary drug-target datasets

Tables S1-S6

Table S1 Average results of LPMIHN-SS, LPMIHN-GO, LPMIHN-PPI and LPMIHN-IP on the four benchmark datasets in 10 CV test repeating 30 times

Dataset	AUC				AUPR				FPR (%)			
	SS	GO	PPI	IP	SS	GO	PPI	IP	SS	GO	PPI	IP
Es	0.950±0.001	0.941±0.002	0.955±0.003	0.988±0.001	0.698±0.003	0.565±0.002	0.588±0.010	0.695±0.003	0.45±0.006	0.50±0.004	0.50±0.01	0.45±0.002
ICs	0.982±0.006	0.972±0.007	0.982±0.003	0.995±0.004	0.815±0.006	0.767±0.004	0.884±0.002	0.86±0.005	0.99±0.03	1.13±0.02	0.75±0.01	0.86±0.03
GPCRs	0.967±0.002	0.965±0.002	0.975±0.003	0.987±0.002	0.915±0.005	0.912±0.005	0.932±0.004	0.934±0.004	0.37±0.02	0.38±0.02	0.38±0.03	0.34±0.02
NRs	0.962±0.01	0.961±0.01	0.94±0.01	0.964±0.007	0.913±0.02	0.907±0.02	0.886±0.014	0.913±0.02	0.93±0.2	1.04±0.17	1.03±0.14	0.99±0.2

Notes: SS, GO, PPI and IP represent the LPMIHN-SS, LPMIHN-GO, LPMIHN-PPI and LPMIHN-IP, respectively.

TableS2 Results of DBSI, TBSI, NBI, NRWRH and LPMIHN methods on the four benchmark datasets in terms of average AUC values with 10CV test repeating 30 times

	Es	ICs	GPCRs	NRs
DBSI	0.781±0.018	0.709±0.025	0.755±0.033	0.789±0.084
TBSI	0.903±0.016	0.905±0.018	0.745±0.044	0.528±0.139
NBI	0.975±0.006	0.976±0.007	0.946±0.019	0.838±0.087
NRWRH	0.953±0.003	0.971±0.005	0.942±0.003	0.867±0.013
LPMIHN	0.994±0.002	0.997±0.002	0.986±0.002	0.965±0.006

Table S3 Results of BLM-NII, NRWRH and LPMIHN methods on the four benchmark datasets in terms of average AUC values with 10CV test repeating 10 times

	Es	ICs	GPCRs	NRs
BLM-NII	0.984±0.001	0.987±0.001	0.978±0.001	0.959±0.006
NRWRH	0.952±0.004	0.972±0.008	0.941±0.003	0.871±0.008
LPMIHN	0.995±0.002	0.996±0.0002	0.986±0.003	0.964±0.008

Table S4 Results of NetlabRLS, NRWRH and LPMIHN methods on the four benchmark datasets in terms of average AUC values with 10CV test repeating 5 times

	Es	ICs	GPCRs	NRs
NetlabRLS	0.956±0.003	0.947±0.003	0.931±0.003	0.856±0.018
NRWRH	0.954±0.004	0.971±0.001	0.943±0.002	0.868±0.009
LPMIHN	0.992±0.002	0.997±0.0002	0.987±0.001	0.966±0.005

Table S5 Results of KBMF2K, NetCBP, NRWRH and LPMIHN methods on the four benchmark datasets in terms of average AUC values with 5CV test repeating 5 times

	Es	ICs	GPCRs	NRs
KBMF2K	0.832	0.799	0.857	0.824
NetCBP	0.983	0.986	0.971	0.888
NRWRH	0.950±0.001	0.969±0.002	0.935±0.005	0.866±0.02
LPMIHN	0.976±0.002	0.991±0.001	0.964±0.002	0.94±0.007

Table S6 Top 50 predicted potential drug-target interactions for Enzymes (Es). Some are verified manually in the latest online versions of ChEMBL(Marked as 'C'), KEGG DRUG(Marked as 'K'), Drugbank(Marked as 'D') and SuperTarget (Marked as 'S') datasets

Drug ID	Gene ID	Verified
D00097	hsa:5743	CD
D00449	hsa:5742	C
D00947	hsa:4129	C
D05458	hsa:4128	D
D00005	hsa:4128	D
D00733	hsa:43	
D01118	hsa:43	C
D00437	hsa:1559	CD
D00567	hsa:1557	
D00126	hsa:1557	D
D00537	hsa:759	
D00703	hsa:762	
D01977	hsa:22954	
D03218	hsa:22954	
D03252	hsa:22954	
D03350	hsa:22954	
D04023	hsa:22954	
D04024	hsa:22954	
D04025	hsa:22954	
D00593	hsa:1557	C
D00283	hsa:1559	D
D00542	hsa:1571	CDK
D03670	hsa:1610	
D00139	hsa:1543	DK
D00293	hsa:1559	D
D00410	hsa:1543	
D00217	hsa:1576	K
D00533	hsa:1559	C
D00536	hsa:1559	C
D00454	hsa:1576	
D01441	hsa:22954	
D00487	hsa:590	D
D00994	hsa:590	D
D00995	hsa:590	C
D00998	hsa:590	C
D01001	hsa:590	
D01228	hsa:590	
D02068	hsa:590	D
D02173	hsa:590	D
D02418	hsa:590	C

D02558	hsa:590	K
D03751	hsa:590	C
D03823	hsa:590	
D03826	hsa:590	
D04292	hsa:590	D
D00300	hsa:1576	
D01164	hsa:1576	D
D02342	hsa:1576	
D02671	hsa:1576	
D00217	hsa:1610	

Table S7 Top 50 predicted potential drug-target interactions for Ion Channels. Some are verified manually in the latest online versions of ChEMBL(Marked as 'C'), KEGG DRUG(Marked as 'K'), Drugbank(Marked as 'D') and SuperTarget(Marked as 'S') datasets.

Drug ID	Gene ID	Verified
D05453	hsa:3738	
D02207	hsa:3782	
D02356	hsa:6323	
D00616	hsa:9312	
D03830	hsa:9312	
D00619	hsa:9312	
D02356	hsa:3776	
D00775	hsa:3778	
D00799	hsa:3782	
D01448	hsa:3782	
D02234	hsa:3782	
D02356	hsa:3767	D
D02356	hsa:1080	
D03742	hsa:6530	D
D00294	hsa:3767	K
D00480	hsa:3781	
D00791	hsa:3781	
D00812	hsa:3781	
D00816	hsa:3781	
D01242	hsa:3781	
D02163	hsa:3781	
D00228	hsa:6531	D
D00394	hsa:6531	D
D00640	hsa:6336	
D00332	hsa:6323	
D00512	hsa:6323	DK
D02356	hsa:2554	
D00294	hsa:2554	
D00542	hsa:3736	
D00538	hsa:6331	
D00648	hsa:3758	
D02360	hsa:6530	C
D02362	hsa:6530	D
D02356	hsa:6326	
D00438	hsa:779	
D01828	hsa:3758	
D01599	hsa:3758	
D01799	hsa:3758	
D01854	hsa:3758	
D00294	hsa:3776	

D00607	hsa:6531	
D01603	hsa:6531	
D02237	hsa:6531	
D02566	hsa:6531	C
D00294	hsa:2741	
D02356	hsa:2741	
D00552	hsa:6331	
D00542	hsa:6323	
D00649	hsa:8911	
D00437	hsa:6323	

Table S8 Top 50 predicted potential drug-target interactions for G-protein coupled receptors. Some are verified manually in the latest online versions of ChEMBL(Marked as 'C'), KEGG DRUG(Marked as 'K'), Drugbank(Marked as 'D') and SuperTarget(Marked as 'S') datasets.

Drug ID	Gene ID	Verified
D02910	hsa:154	
D00454	hsa:152	D
D02358	hsa:154	CD
D04625	hsa:154	CK
D00283	hsa:152	D
D02614	hsa:154	
D00683	hsa:153	
D00684	hsa:153	D
D00687	hsa:153	S
D00688	hsa:153	S
D01386	hsa:153	K
D02147	hsa:153	D
D02359	hsa:153	
D05792	hsa:153	C
D00283	hsa:11255	
D00760	hsa:1128	
D00765	hsa:1128	
D00513	hsa:152	C
D00110	hsa:4988	
D00332	hsa:135	C
D00437	hsa:152	D
D00255	hsa:152	C
D00726	hsa:1129	
D01713	hsa:152	
D02356	hsa:152	
D02327	hsa:1129	
D00613	hsa:1813	

D00110	hsa:1813	
D00525	hsa:1129	K
D00715	hsa:1129	K
D01103	hsa:1129	K
D01118	hsa:1129	K
D01269	hsa:1129	D
D01297	hsa:1129	
D00604	hsa:148	C
D00604	hsa:147	
D00283	hsa:4543	
D00283	hsa:1814	CD
D00454	hsa:4543	
D00095	hsa:155	K
D00270	hsa:152	
D00136	hsa:152	
D00318	hsa:3269	
D00235	hsa:155	
D00598	hsa:155	
D00601	hsa:155	
D00632	hsa:155	
D00635	hsa:155	
D00645	hsa:155	
D02342	hsa:155	

Table S9 Top 50 predicted potential drug-target interactions for Nuclear receptors. Some are verified manually in the latest online versions of ChEMBL(Marked as 'C'), KEGG DRUG(Marked as 'K'), Drugbank(Marked as 'D') and SuperTarget(Marked as 'S') datasets.

Drug ID	Gene ID	Verified
D00585	hsa:2099	C
D00182	hsa:2099	C
D00951	hsa:2099	D
D00690	hsa:2099	
D01217	hsa:2099	
D00577	hsa:5241	
D00105	hsa:5241	
D01161	hsa:5241	
D02217	hsa:5241	
D04066	hsa:5241	
D00327	hsa:5241	
D00066	hsa:2100	
D00950	hsa:2100	
D00954	hsa:2100	

D01294	hsa:2100	
D02367	hsa:2100	
D00067	hsa:5241	
D00312	hsa:5241	
D00554	hsa:5241	
D00898	hsa:5241	
D00962	hsa:5241	
D00327	hsa:2100	
D00067	hsa:2100	K
D00312	hsa:2100	K
D00554	hsa:2100	CK
D00898	hsa:2100	CK
D00962	hsa:2100	K
D00075	hsa:2099	C
D00462	hsa:2099	
D00586	hsa:2099	C
D00956	hsa:2099	
D00961	hsa:2099	
D00965	hsa:2099	
D00066	hsa:367	K
D00950	hsa:367	D
D00954	hsa:367	D
D01294	hsa:367	C
D02367	hsa:367	
D00577	hsa:367	
D00316	hsa:6096	
D00088	hsa:5241	
D00246	hsa:5241	
D01387	hsa:5241	
D01689	hsa:5241	
D00066	hsa:2908	
D00950	hsa:2908	C
D00954	hsa:2908	C
D01294	hsa:2908	C
D02367	hsa:2908	
D00105	hsa:367	
