Supplementary Materials for

"Predicting Human Microbe-Drug Associations via Graph Convolution Network with Conditional Random Field"

1. Supplementary Figures

In this paper, we compared our method with some diverse types of state-of-the-art computational methods developed for addressing other biomedical problems, such as microbe-disease prediction (i.e., KATZHMDA, NTSHMDA and WMGHMDA), miRNA-disease prediction (i.e., IMCMDA), miRNA-drug resistance prediction (i.e., GCMDR) and drug-target prediction (i.e., WNN-GIP and GCNMDA). We evaluated the performance for all methods using 2-fold CV, 5-fold CV and 10-fold CV in terms of AUC and AUPR scores. The results for AUC and AUPR have been shown in Table S1 and Table S2, respectively. In addition, for more intuitive comparison, Figure S1 and Figure S2 display corresponding ROC and PR curves, respectively.

Table S1. Performance comparison between baseline methods and our method in terms of AUC. The best results are marked in bold. Underline denotes the second best method.

Method	2-fold CV	5-fold CV	10-fold CV
KATZHMDA	0.7865±0.0254	0.8723 ± 0.0032	0.8929 ± 0.0265
NTSHMDA	0.7766±0.0126	0.8302 ± 0.0089	0.8547 ± 0.0165
WMGHMDA	0.8519±0.0024	0.8654 ± 0.0122	0.8729 ± 0.0130
IMCMDA	0.7285±0.0065	0.7466 ± 0.0102	0.7466 ± 0.0221
GCMDR	0.8432±0.0016	0.8485 ± 0.0062	0.8590 ± 0.0241
BLMNII	0.8583 ± 0.0239	0.9231 ± 0.0170	0.8644 ± 0.0227
WNNGIP	0.8256±0.0516	0.8721 ± 0.0162	0.8711 ± 0.0130
GCNMDA	0.9384±0.0028	0.9423±0.0105	0.9420±0.0078

Table S2. Performance comparison between baseline methods and our method in terms of AUPR. The best results are marked in bold. Underline denotes the second best method.

Method	2-fold CV	5-fold CV	10-fold CV
KATZHMDA	0.6559±0.0034	0.8081 ± 0.0209	0.8403 ± 0.0182
NTSHMDA	0.5674±0.0057	0.6735 ± 0.0092	0.7118 ± 0.0442
WMGHMDA	0.8201 ± 0.0086	0.8381 ± 0.0083	0.8452 ± 0.0117
IMCMDA	0.7618±0.0038	0.7773 ± 0.0113	0.7727 ± 0.0136
GCMDR	0.8423 ± 0.0025	0.8509 ± 0.0040	0.8634 ± 0.0210
BLMNII	0.7805±0.0318	0.9263 ± 0.0152	0.8792 ± 0.0221
WNNGIP	0.8299±0.0603	0.8922 ± 0.0137	0.8863 ± 0.0232
GCNMDA	0.9316±0.0017	0.9376±0.0114	0.9359±0.0087

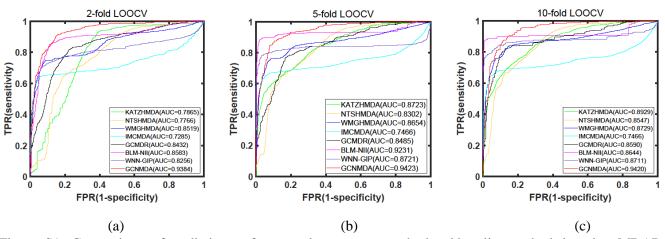


Figure. S1. Comparisons of prediction performance between our method and baseline methods based on MDAD under the frameworks of 2-fold CV, 5-fold CV and 10-fold CV in terms of AUC, respectively. (a) 2-fold CV; (b) 5-fold CV; (c) 10-fold CV.

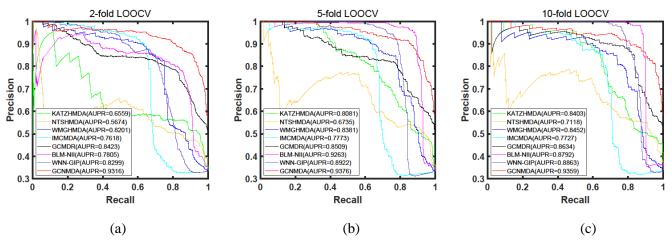


Figure. S2. Comparisons of prediction performance between our method and baseline methods based on MDAD under the frameworks of 2-fold CV, 5-fold CV and 10-fold CV in terms of AUPR, respectively. (a) 2-fold CV; (b) 5-fold CV; (c) 10-fold CV.

2. Supplementary Tables

For further verify the effectiveness of our method, we adopt two popular drugs (i.e., Ciprofloxacin and Moxifloxacin) and two popular microbes (i.e., Pseudomonas aeruginosa and Escherichia coli.) for case study. We validated the predicted candidate microbes using previous reports. The top 50 predicted candidate microbes have been listed in Table S3 and Table S4 for Ciprofloxacin and Moxifloxacin, respectively. The top 50 predicted candidate drugs have been listed in Table S5 and Table S6 for *Pseudomonas aeruginosa* and *Escherichia coli.*, respectively.

Table S3. Prediction results of the top 50 Ciprofloxacin-associated microbes. The first column records top 1-25 associated microbes. The third column records top 26-50 associated microbes.

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Microbe	Evidence	Microbe	Evidence
Candida albicans	PMID:31471074	Vibrio anguillarum	PMID:19146525
Streptococcus mutans	PMID:30468214	Proteus mirabilis	PMID:26953206
Staphylococcus epidermis	PMID:10632381	Eikenella corrodens	PMID:16875802
Staphylococcus epidermidis	PMID:28481197	Micrococcus luteus	PMID:26954218
Enterococcus faecalis	PMID:27790716	Bacillus cereus	PMID:26358183
Vibrio harveyi	PMID:27247095	Plasmodium falciparum	PMID:17214980
Salmonella enterica	PMID:26933017	Human immunodeficiency virus	PMID:26195465
Human immunodeficiency virus 1	PMID:9566552	Porphyromonas gingivalis	PMID: 15231772
Actinomyces oris	Unconfirmed	Enterococcus faecium	PMID:28193670
Streptococcus sanguis	PMID: 11347679	Streptococcus gordonii	PMID:22887906
Enteric bacteria and other eubacteria	PMID:27436461	Campylobacter jejuni	PMID:27900889
Listeria monocytogenes	PMID:28355096	Vibrio campbellii	Unconfirmed
Burkholderia cenocepacia	PMID:27799222	Klebsiella planticola	PMID:25465871
Streptococcus pneumoniae	PMID:26100702	Human herpesvirus 1	Unconfirmed
Burkholderia pseudomallei	PMID:24502667	Pichia anomala	Unconfirmed
Burkholderia multivorans	PMID: 19633000	Candida tropicalis	Unconfirmed
Clostridium perfringens	PMID:29978055	Candida glabrata	Unconfirmed
Serratia marcescens	PMID:23751969	Cryptococcus neoformans	Unconfirmed
Streptococcus epidermidis	Unconfirmed	Aeromonas hydrophila	PMID:23084650
Klebsiella pneumoniae	PMID:27257956	Streptococcus pneumoniae serotype 4	PMID: 12543698
Vibrio cholerae	PMID:28270803	Influenza A virus	Unconfirmed
Acinetobacter baumannii	PMID:29998878	Aeromonas sobria	PMID:28484157
Vibrio vulnificus	PMID:28971862	Hafnia alvei	PMID:28537065
Propionibacterium acnes	PMID:27801379	Enterococcus gallinarum	PMID:29030312
Streptococcus sanguinis	PMID:21507381	Mycobacterium avium	PMID:30012773

Table S4. Prediction results of the top 50 Moxifloxacin-associated microbes. The first column records top 1-25 associated microbes. The third column records top 26-50 associated microbes.

Microbe	Evidence	Microbe	Evidence
Pseudomonas aeruginosa	PMID:31691651	Mycobacterium tuberculosis	PMID:31713607
Staphylococcus aureus	PMID:31689174	Vibrio vulnificus	PMID: 12384368
Escherichia coli	PMID:31542319	Propionibacterium acnes	PMID:31538294
Streptococcus mutans	PMID:29160117	Streptococcus sanguinis	Unconfirmed
Staphylococcus epidermis	PMID: 11249827	Vibrio anguillarum	Unconfirmed
Staphylococcus epidermidis	PMID:31516359	Proteus mirabilis	PMID:27351708
Enterococcus faecalis	PMID:31763048	Eikenella corrodens	PMID:14614671
Bacillus subtilis	PMID:30036828	Micrococcus luteus	PMID:24231380
Vibrio harveyi	Unconfirmed	Bacillus cereus	PMID:21834669

Salmonella enterica	PMID:22151215	Plasmodium falciparum	PMID:15125930
Human immunodeficiency virus 1	Unconfirmed	Human immunodeficiency virus	PMID:31388545
Actinomyces oris	PMID: 26538502	Porphyromonas gingivalis	PMID:30048853
Streptococcus sanguis	PMID:10629010	Enterococcus faecium	PMID:23524466
Enteric bacteria and other eubacteria	Unconfirmed	Streptococcus gordonii	PMID:29160117
Listeria monocytogenes	PMID:28739228	Campylobacter jejuni	PMID:16027651
Burkholderia cenocepacia	Unconfirmed	Vibrio campbellii	Unconfirmed
Streptococcus pneumoniae	PMID:31542319	Klebsiella planticola	Unconfirmed
Burkholderia pseudomallei	PMID:15731198	Proteus vulgaris	PMID: 19692210
Burkholderia multivorans	Unconfirmed	Human herpesvirus 1	Unconfirmed
Clostridium perfringens	PMID:29486533	Pichia anomala	Unconfirmed
Serratia marcescens	PMID:17592324	Candida tropicalis	PMID:20455400
Streptococcus epidermidis	Unconfirmed	Candida glabrata	PMID:20455400
Klebsiella pneumoniae	PMID:30058553	Cryptococcus neoformans	Unconfirmed
Vibrio cholerae	PMID:16341343	Aeromonas hydrophila	PMID:12821471
Acinetobacter baumannii	PMID:31261589	Streptococcus pneumoniae serotype	Unconfirmed
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Table S5. Prediction results of the top 50 *Pseudomonas aeruginosa*-associated drugs. The first column records top 1-25 associated drugs. The third column records top 26-50 associated drugs.

Microbe	Evidence	Microbe	Evidence
Biphenyl	PMID: 26519802	Isohydnocarpin	Unknown
L-Phenylalanine-L-Lysine	Unknown	Quercetin Dihydrate	Unknown
Ticarcillin	PMID: 26248364	Sage oil	PMID: 27585258
Ethyl acetate	PMID: 26887538	Silver sulphate	Unknown
2,3-dihydroxybenzoic acid	PMID: 24449781	Phylloseptin-1	Unknown
Ionic silver	PMID: 25001302	Resorcinol	PMID: 28989246
Phloridzin	Unknown	D-Aspartate	PMID: 25656450
3-tetra- decanoyltetronic	Unknown	Oregano oil	PMID: 30344513
Dichloromethane	PMID: 29375509	Quercetin-3-O-glucuronide	Unknown
3-tetradecanoylthiotetronic	PMID: 27999062	3-Hydroxytyrosine	Unknown
(5Z)-2-(1,1-dichloro-7-methyloctyl)-5-{[4-(trifluoromethyl)phenyl]methylidene}-1,3-oxazolidin-4-one	Unknown	(5Z)-2-(1,1-dichloropropyl)-5-{[4-(trifluoromethyl)phenyl]methyliden e}-1,3-oxazolidin-4-one	Unknown
(5Z)-2-(1,1-dichlorobut-3-yn-1-yl)-5-{[4-(trifluoromethyl)phenyl]methylidene}-1,3-oxazolidin-4-one	Unknown	PTP-7	Unknown
5- hydroxyethyl-3-tetradecanoyltetramic acid	PMID: 27999062	Silver dihydrogen citrate	Unknown
P10	PMID: 3141560	Pine needle oil	Unknown
CAMA	PMID: 30849936	Na-Propionate	PMID: 10515415
Teichoplanin	Unknown	Tangerine oil	Unknown
cis-Stillbene	Unknown	Diphenyl methane	Unknown
Myricitrin	PMID: 25993634	Clary sage oil	Unknown
Cecropin (1-7)-melittin A (2-9) amide	Unknown	alpha-tocopherol	PMID: 25458794
Myrrh oil	PMID: 22288378	Protamine sulphate	PMID: 7574516
HE4	Unknown	Silybin	PMID: 18758739
Tea tree oil	PMID: 23581401	(2Z)-N-hexyl-2-hydroxy-3-[4- (trifluoromethyl)phenyl]prop-2- enamide	Unknown
Myricetin	PMID: 30213633	Hesperetin	Unknown
trans-Resveratrol	PMID: 29530588	(5Z)-2-(1,1-dichloro-3,3-dimethylbutyl)-5-	Unknown

		(phenylmethylidene)-1,3- oxazolidin-4-one	
Methanol	PMID: 28637123	(10R,11R)-Hydnocarpin D	Unknown

Table S6. Prediction results of the top 50 *Escherichia coli*.-associated drugs. The first column records top 1-25 associated drugs. The third column records top 26-50 associated drugs.

Microbe	Evidence	Microbe	Evidence
Indol AHL	PMID: 22101045	Seg6D	Unknown
7-bromoindole	PMID: 27921270	Zinc Dichloride	PMID: 16955743
(2S,3R,4S,5S,6R)-6-(Acetoxymethyl)-3-(3-oxohexanamido)tetrahydro-2H-pyran-2,4,5-triyl triacetate	PMID: 28049617	D, L- K6L9	Unknown
LF11-322	PMID: 21515687	2-(5-Bromo-2-(2- chloroacetamido)phenyl)-2-oxo- N-(prop-2-yn-1-yl)acetamide	Unknown
N-((2R,3R,4S,5S,6R)-2,4,5-Trihydroxy-6-(hydroxymethyl)tetrahydro-2H-pyran-3-yl)hexanamide	PMID: 28049617	Gallium cation	Unknown
Cobalt Ion	PMID: 16158232	Silver Ion	PMID: 28458540
ZnO nanoparticles	PMID: 27614583	3beta-Hydroxyolean-12-en-28-al 3-p-coumarate	Unknown
Indirubin	PMID: 24500796	Furanone	PMID: 24118802
4-tertiary butyl benzyl decyl beta-maltoside	Unknown	3-fluoroaniline	Turan <i>et al</i> . (2012)
14,15,29,30-Tetrahydro-5H,20H-dibenzo[h,s]bis([1,2,3]triazolo)[5,1-c:5_,1n][1,4,7,12,15,18]hexaazacyclodocosine-6,12,13,21,27,28(7H,22H)-hexaone 14,15,29,30-Tetrahydro-5H,20H-	Unknown	D-galactose cluster	PMID: 15382922
dibenzo[h,s]bis([1,2,3]triazolo)[5,1-c:5_,1n][1,4,7,12,15,18]hexaazacyclodocosine-6,12,13,21,27,28(7H,22H)-hexaone	Unknown	14-Glycinyl andrographolide hydrochloride	PMID: 19152987
2-((3,4-dichlorophenyl)sulfonyl)-4-(1H-tetrazol-1-yl)pyrimidine	Unknown	7-fluoroindole	PMID: 22251040
Tellurite	PMID: 25447814	7-methylindole	Unknown
7-chloroindole	PMID: 27921270	14-dehydroxy-11,12- didehydroandrographolide- 3,19- bis(succinic acid) potasium sodium salt	Unknown
2-oxindole	PMID: 25664629	4-tertiary butyl benzyl dodecyl beta-maltoside	Unknown
7-Bicyclic brominated furanones	Unknown	Norpinguisone	Jorge et al. 2013
N-(5-Methyl-1,2-oxazol-3-yl)butanamide	Unknown	Delphinidin	PMID: 29765781
Tricaine	PMID: 20019794	5-Chloro-1-(2-chloroacetyl)indoline-2,3-dione	PMID: 28654117
N-Decanoyl cyclopentylamide	PMID: 17369333	Decyl beta-maltoside	Unknown
O-LF11-215	Unknown	para-ethylaniline	Unknown
Pelargonidin	PMID: 29554804	6-MO-LF11-215	Unknown
Benzocaine	PMID: 22556388	2-(2-(2-Azidoacetamido)-5- fluorophenyl)-2-oxo-N-(prop-2- yn-1-yl)acetamide	PMID: 28654117
(2S,3R,4S,5S,6R)-6-(Acetoxymethyl)-3-(3-oxooctanamido)tetrahydro-2H-pyran-2,4,5-triyl triacetate	PMID: 28049617	N-((2R,3R,4S,5S,6R)-2,4,5- Trihydroxy-6-	PMID: 28049617

		(hydroxymethyl)tetrahydro-2H- pyran-3-yl)butyramide	
2-(4-Bromophenyl)-N-((2R,3R,4S,5S,6R)-2,4,5-trihydroxy-6-(hydroxymethyl)tetrahydro-2H-pyran-3-yl)acetamide	PMID: 28049617	4-methanesulfonylaniline	Unknown
para-ethoxyaniline	Unknown	8-fluoroquinoline	PMID: 22251040

Reference

Turan, N. *et al.* (2012) Synthesis, Characterization of Poly(E)-3-amino-4-((3-bromophenyl)diazenyl)1H-pyrazol-5-ol: Investigation of Antibacterial Activity, Fluorescence, and Optical Properties. *Fibers and Polymers*, 13, 415-424.

Jorge, C.S. et al. (2013) Bioprospecting, a tool to conserve Chilean bryophytes. Gayana Bot, 70, 16-25.