Analysis

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1 摘要

1.1 主要生信部分

- Sjogren syndrome (SS), a chronic autoimmune disease
- 复方, Xuanmai Ganju (XMGJ)

原文以 GEO 做 SS 差异表达基因筛选,获取 XMGJ 的成分和靶点信息,构建 PPI 网络,筛选治疗疾病靶点。

1.2 修正内容

- 原文使用的是 TCMSP database https://tcmspw.com/tcmsp.php 检索复方成分数据和靶点数据,由于订单中不包含该部分原数据,所以该内容已全部重新分析。数据库修改为 HERB http://herb.ac.cn/¹。
- GEO 数据的分析全部重修。原文未包含具体的数据库来源(GEO 编号),也未提供原始数据,因此重修。
- 以上部分对应的图片全部重修。
- 2 前言
- 3 研究设计流程图
- 4 材料和方法

数据来源,见7。

方法:

- $HERB^1$
- \bullet clusterProfiler²
- $STRINGdb^3$
- $limma^4$
- 5 分析结果
- 6 结论
- 7 附:分析流程
- 7.1 复方成分和靶点
- 7.1.1 复方成分

Table 1为表格 Herbs information 概览。

(对应文件为 Figure+Table/Herbs-information.xlsx)

注:表格共有4行18列,以下预览的表格可能省略部分数据;表格含有4个唯一'Herb_'。

Table 1: Herbs information

Herb_	Herb	Herb	Herb	Herb	Prope	Merid	UsePart	Function	Indic	Toxicity
HERB0	GAN CAO	甘草	Root	Radix	Mild;	Lung;	root	To re	1. It	NA
HERB0	JIE GENG	桔梗	Platy	Radix	Mild;	Lung	root	To re	Cough	NA
${\rm HERB0}$	MAI DONG	麦冬	Lirio	Ophio	Minor	Lung;	tuberoid	To no	Angin	NA
HERB0	XUAN	玄参	Figwo	Radix	$\operatorname{Cold};$	Lung;	root	To re	Heat	NA

Figure 1为图 intersection of all compounds 概览。

(对应文件为 Figure+Table/intersection-of-all-compounds.pdf)

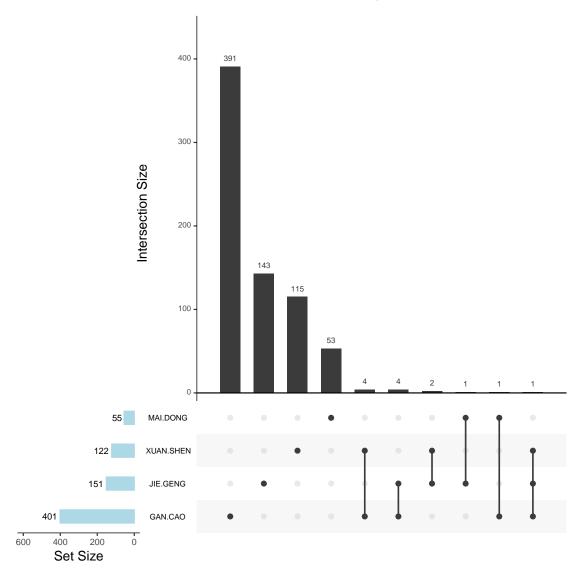


Figure 1: Intersection of all compounds

7.1.2 成分靶点

Table 2为表格 Components of the herbs and their targets 概览。

(对应文件为 Figure+Table/Components-of-the-herbs-and-their-targets.xlsx)

注:表格共有 6181 行 9 列,以下预览的表格可能省略部分数据;表格含有 715 个唯一'Ingredient.id'。

Table 2: Components of the herbs and their targets

Ingre1	Herb	Ingre3	Ingre4	Targe5	Targe6	Datab	Paper.id	
HBIN0	GAN CAO	11-de	11-De	HBTAR	NR3C1	NA	NA	
HBIN0	GAN CAO	11-de	11-De	HBTAR	AR	NA	NA	
HBIN0	GAN CAO	12-me	12-me	NA	NA	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	AR	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	CDK2	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	ESR1	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	GSK3B	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	MAPK14	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	CHEK1	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	PPARG	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	CCNA2	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	ESR2	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	CCNA2	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	CDK2	NA	NA	
HBIN0	GAN CAO	1,3-d	1,3-d	HBTAR	ESR1	NA	NA	
		•••						

Figure 2为图 intersection of all targets 概览。

(对应文件为 Figure+Table/intersection-of-all-targets.pdf)

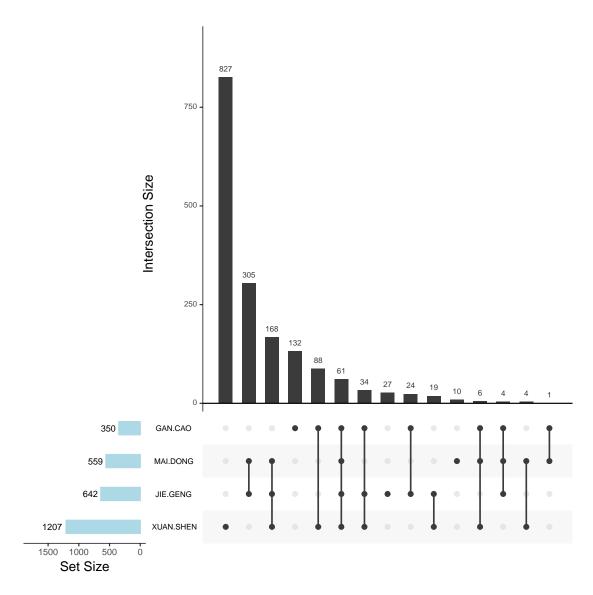


Figure 2: Intersection of all targets

7.2 GEO 数据分析

7.2.1 GSE7451: Gene profilling from 10 saliva samples from patients with primary Sojgren's syndrome and 10 saliva samples from control subjects using Affymetrix HGu133+2 microarray.

Table 3为表格 metadata of used samples of GSE7451 概览。

(对应文件为 Figure+Table/metadata-of-used-samples-of-GSE7451.xlsx)

注: 表格共有 20 行 7 列,以下预览的表格可能省略部分数据;表格含有 2 个唯一'group'。

Table 3: Metadata of used samples of GSE7451

rownames	group	lib.size	norm	sample	title	gende
GSM17	control	23247	1.101	GSM17	whole	femal
GSM18	control	23164	1.117	GSM18	whole	${\rm femal}$
GSM18	control	23523	1.213	GSM18	whole	femal
GSM18	control	23553	1.177	GSM18	whole	${\rm femal}$
GSM18	control	23114	1.049	GSM18	whole	femal
GSM18	control	23397	1.266	GSM18	whole	${\rm femal}$
GSM18	control	22724	0.996	GSM18	whole	${\rm femal}$
GSM18	control	23013	1.051	GSM18	whole	${\rm femal}$
GSM18	control	21367	0.721	GSM18	whole	${\rm femal}$
GSM18	control	20958	0.595	GSM18	whole	${\rm femal}$
GSM18	disease	23050	1.120	GSM18	whole	${\rm femal}$
GSM18	disease	22818	0.917	GSM18	whole	${\rm femal}$
GSM18	disease	21152	0.763	GSM18	whole	${\rm femal}$
GSM18	disease	22518	0.957	GSM18	whole	${\rm femal}$
GSM18	disease	22748	1.142	GSM18	whole	${\rm femal}$

Figure 3为图 DEGs of GSE7451 概览。

(对应文件为 Figure+Table/DEGs-of-GSE7451.pdf)

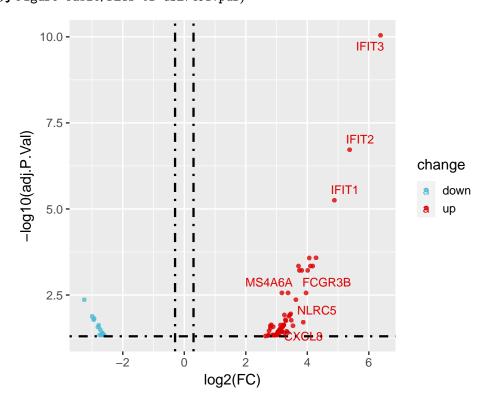


Table 4为表格 tables of DEGs of GSE7451 概览。

(对应文件为 Figure+Table/tables-of-DEGs-of-GSE7451.xlsx)

注: 表格共有 75 行 23 列,以下预览的表格可能省略部分数据;表格含有 75 个唯一'rownames'。

Table 4: Tables of DEGs of GSE7451

rownames	hgnc	ID	GB_ACC	SPOT_ID	Speci	Annot	Seque8	Seque9	Targe	Re
22945	IFIT3	22945	AI075407		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AI	AI
22675	IFIT2	22675	AA131041		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AA	AA
20315	IFIT1	20315	NM_00		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	NN
22670	CMPK2	22670	AI742057		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AI	AI
21445	IFI44	21445	NM_00		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:NM	NN
20566	OASL	20566	NM_00		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	NN
15654	EGFR	15654	AF277897		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AF	AF
22861	XAF1	22861	AA142842		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AA	AA
21899	TMEM140	21899	NM_01		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	NN
22760	EPSTI1	22760	AA633203		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AA	AA
24262	RSAD2	24262	AW189843		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AW	AV
15584		15584	AW974642		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AW	AV
21966	MS4A6A	21966	NM_02		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	NN
20400	FCGR3B	20400	J04162		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:J0	$J0^2$
23839		23839	AW301504		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AW	AV

• saliva samples

7.2.2 GSE40611: Parotid gland tissues were harvested from 17 pSS and 14 non-pSS sicca patients and 18 controls.

• Parotid gland

Table 5为表格 metadata of used samples of GSE40611 概览。

(对应文件为 Figure+Table/metadata-of-used-samples-of-GSE40611.csv)

注:表格共有 35 行 9 列,以下预览的表格可能省略部分数据;表格含有 2 个唯一'group'。

Table 5: Metadata of used samples of GSE40611

rownames	group	lib.size	norm	sample	title	batch	disea	batch
GSM99	control	78182	1.949	GSM99	parot	1	Control	1
GSM99	control	65241	1.158	GSM99	parot	1	Control	1
GSM99	control	60270	0.969	GSM99	parot	1	Control	1
GSM99	control	71411	1.592	GSM99	parot	1	Control	1
GSM99	control	64687	1.171	GSM99	parot	1	Control	1
GSM99	control	57764	0.998	GSM99	parot	1	Control	1
GSM99	control	77992	1.919	GSM99	parot	1	Control	1
GSM99	control	61315	1.050	GSM99	parot	1	Control	1
GSM99	control	62902	1.120	GSM99	parot	1	Control	1
GSM99	control	44310	0.841	GSM99	parot	2	Control	2
GSM99	control	47089	0.891	GSM99	parot	2	Control	2
GSM99	control	48607	0.937	GSM99	parot	2	Control	2
GSM99	control	49006	0.958	GSM99	parot	2	Control	2
GSM99	control	41798	0.790	GSM99	parot	3	Control	3
GSM99	control	48423	0.829	GSM99	parot	3	Control	3

Figure 4为图 DEGs of GSE40611 概览。

(对应文件为 Figure+Table/DEGs-of-GSE40611.pdf)

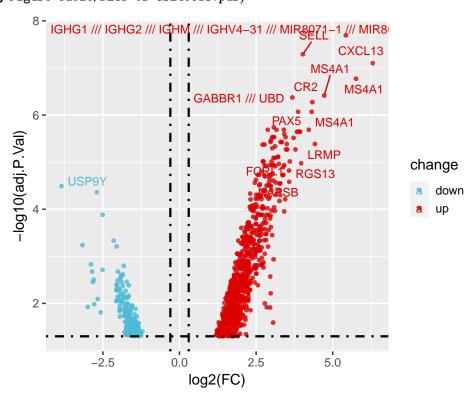


Table 6为表格 tables of DEGs of GSE40611 概览。

(对应文件为 Figure+Table/tables-of-DEGs-of-GSE40611.xlsx)

注: 表格共有 1161 行 23 列,以下预览的表格可能省略部分数据;表格含有 1161 个唯一'rownames'。

Table 6: Tables of DEGs of GSE40611

rownames	hgnc	ID	GB_ACC	SPOT_ID	Speci	Annot	Seque8	Seque9	Targe	R
21143	IGHG1	21143	M87789		Homo	Oct 6	Exemp	GenBank	gb:M8	N
20456	SELL	20456	NM_00		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	N
20524	CXCL13	20524	NM_00		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	N
22859	MS4A1	22859	AW474852		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AW	A
21035	MS4A1	21035	BC002807		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:BC	В
20589	GABBR	20589	NM_00		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	N
20554	CR2	20554	NM_00		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	N
22196	PAX5	22196	BF510692		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:BF	В
22859	MS4A1	22859	AI862674		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AI	A
22853	SAMD9	22853	AA741307		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:AA	A
22660	SAMD9L	22660	BE966604		Homo	Oct 6	Conse	$\operatorname{GenBank}$	gb:BE	В
20329	HLA-DQA1	20329	NM_00		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	N
20423	FLI1	20423	NM_00		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	N
20411	CD48	20411	NM_00		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	N
20391	CXCL9	20391	NM_00		Homo	Oct 6	Exemp	$\operatorname{GenBank}$	gb:NM	N
					•••			•••		

7.2.3 GSE154926: Total mRNA was extracted from minor salivary glands of 43 SS patients and 7 healthy volunteers.

• minor salivary glands

Table 7为表格 metadata of used samples of GSE154926 概览。

(对应文件为 Figure+Table/metadata-of-used-samples-of-GSE154926.csv)

注:表格共有50行9列,以下预览的表格可能省略部分数据;表格含有2个唯一'group'。

Table 7: Metadata of used samples of GSE154926

rownames	group	lib.size	norm	sample	age.ch1	diagn	gende	tissu
BRA2	control	18452	0.860	BRA2	21	Healt	Female	minor

rownames	group	lib.size	norm	sample	age.ch1	diagn	gende	tissu
BRA3	control	35415	1.221	BRA3	59	Healt	Female	minor
BRA4	control	11889	0.970	BRA4	71	Healt	Female	minor
BRA31	control	15461	0.758	BRA31	24	Healt	Female	minor
BRA33	control	26419	1.448	BRA33	61	Healt	Female	minor
BRA36	control	18885	0.976	BRA36	14	Healt	Female	minor
BRA37	control	55874	1.782	BRA37	52	Healt	Female	minor
BRA41	disease	30095	1.303	BRA41	53	prima	Female	minor
BRA64	disease	14484	0.902	BRA64	55	prima	Female	minor
BRA11	disease	22380	1.116	BRA11	51	prima	Female	minor
BRA13	disease	21075	1.134	BRA13	51	prima	Female	minor
BRA14	disease	22754	1.497	BRA14	48	prima	Female	minor
BRA15	disease	16689	1.227	BRA15	58	prima	Female	minor
BRA16	disease	55177	0.367	BRA16	65	prima	Female	minor
BRA18	disease	13631	0.815	BRA18	69	prima	Female	minor
<u></u>	•••	•••	•••	•••	•••		•••	

Figure 5为图 DEGs of GSE154926 概览。

(对应文件为 Figure+Table/DEGs-of-GSE154926.pdf)

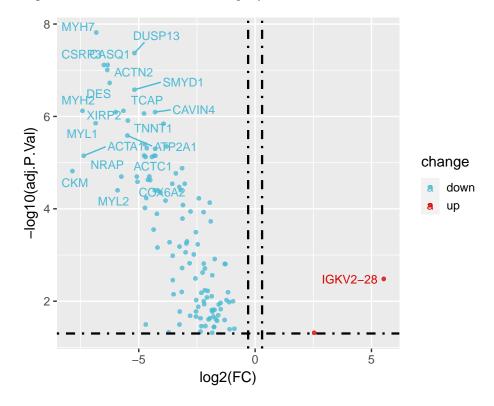


Figure 5: DEGs of GSE154926

Table 8为表格 tables of DEGs of GSE154926 概览。

(对应文件为 Figure+Table/tables-of-DEGs-of-GSE154926.xlsx)

注: 表格共有 123 行 14 列,以下预览的表格可能省略部分数据;表格含有 123 个唯一'ensembl_gene_id'。

Table 8: Tables of DEGs of GSE154926

ensem	hgnc	entre	refse	chrom	start	end_p	descr	logFC	AveExpr	t	P.Va
ENSG0	MYH7	4625	NM_00	14	23412740	23435660	myosi	-6.82	-2.41	-9.37	9.110
ENSG0	DUSP13	51207	NM_01	10	75094432	75109221	dual	-5.18	-3.45	-8.88	5.071
ENSG0	CASQ1	844	NM_00	1	16019	16020	calse	-6.33	-1.73	-8.57	1.549
ENSG0	CSRP3	8048		11	19182030	19210571	cyste	-6.49	-1.64	-8.53	1.838
ENSG0	ACTN2	88		1	23666	23676	actin	-6.35	-1.67	-8.40	2.948
ENSG0	DES	1674	NM_00	2	21941	21942	desmi	-6.25	-1.35	-8.17	6.739
ENSG0	SMYD1	150572	NM_19	2	88067825	88113384	SET a	-5.18	-3.00	-8.03	1.102
ENSG0	MYH2	4620		17	10521148	10549700	myosi	-7.41	-0.01	-7.68	3.967
ENSG0	TCAP	8557	NM_00	17	39665349	39666554	titin	-5.65	-2.06	-7.67	4.063
ENSG0	XIRP2	129446	NM_00	2	16688	16725	xin a	-5.98	-2.27	-7.61	5.165
ENSG0	NEB	4703	NM_00	2	15148	15173	nebul	-4.77	1.733	-7.56	6.197
ENSG0	CAVIN4	347273	NM_00	9	10057	10058	caveo	-4.29	-2.27	-7.60	5.282
ENSG0	TNNT1	7138	NM_00	19	55132698	55149206	tropo	-5.46	-0.98	-7.44	9.538
ENSG0	MYL1	4632	NM_07	2	21029	21031	myosi	-6.85	-0.75	-7.39	1.171
ENSG0	TNNI2	7136	NM_00	11	1838981	1841680	tropo	-3.93	0.268	-7.36	1.285
•••	•••	•••	•••	•••				•••			

7.2.4 GSE135635: RNA-sequencing in primary circulating plasmacytoid dendritic cells from patients with pSS, nSS, and HC. Two cohorts of n=31 each are included (discovery and replication), no duplicates samples

Table 9为表格 metadata of used samples of GSE135635 概览。

(对应文件为 Figure+Table/metadata-of-used-samples-of-GSE135635.csv)

注:表格共有 22 行 5 列,以下预览的表格可能省略部分数据;表格含有 2 个唯一'group'。

Table 9: Metadata of used samples of GSE135635

rownames	group	lib.size	norm	sample
Disc.HC1	control	56201	0.986	Disc.HC1
${\bf Disc. HC2}$	control	54656	0.984	${\bf Disc. HC2}$
${\bf Disc. HC3}$	control	55765	0.959	${\bf Disc. HC3}$
Disc.HC4	control	59405	1.037	Disc.HC4

rownames	group	lib.size	norm	sample
Disc.HC5	control	55793	0.957	Disc.HC5
Disc.HC6	control	57688	0.997	Disc.HC6
${\rm Disc. HC7}$	control	55350	0.961	Disc.HC7
Disc.HC8	control	56148	0.995	Disc.HC8
Disc	disease	56631	1.016	Disc
Disc	disease	59579	1.007	Disc
Disc	disease	59215	1.038	Disc
Disc	disease	58367	1.033	Disc
Disc	disease	56710	1.002	Disc
Disc	disease	58039	1.024	Disc
Disc	disease	56652	1.008	Disc
	•••			

Figure 6为图 DEGs of GSE135635 概览。

(对应文件为 Figure+Table/DEGs-of-GSE135635.pdf)

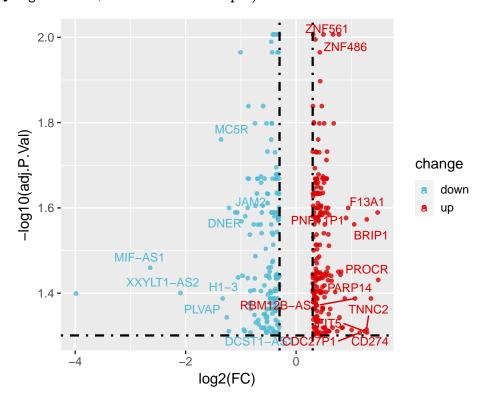


Figure 6: DEGs of GSE135635

Table 10为表格 tables of DEGs of GSE135635 概览。

(对应文件为 Figure+Table/tables-of-DEGs-of-GSE135635.xlsx)

注:表格共有 394 行 14 列,以下预览的表格可能省略部分数据;表格含有 394 个唯一 'ensembl_gene_id'。

Table 10: Tables of DEGs of GSE135635

ensem hgnc entre	refse chrom.	. start end_pdescr	$\log FC$	AveExprt	P.Value
ENSG0.ZNF56193134	19	96046809621236z inc	0.668	4.194 6.524	1.120
ENSG0.TCF3 6929	19	 16092911652615 trans	-	7.500	1.935
			0.36	6.29	
ENSG0.UBE4A 9354	NM_001.1	11835 11839 ubiqu	0.309	5.755 6.074	3.266
ENSG0.ATP7A 538	X	7791069 7 805039 A TPas.	0.494	3.524 6.094	3.112
${\bf ENSG0.TMIGD226259}$	NM_0019	$42922274302431\mathrm{trans}$	-	5.504	4.615
			0.39	5.93	
ENSG0.TRAF4 9618	NM_0017	$2874401 2875095 \mathbf{GNF}$	-	8.484	5.132
		r	0.42	5.88	
${\rm ENSG0.FAM3A60343}$	NM_00X	15450 15451 FAM3	-	4.948	5.324
			0.30	5.87	
ENSG0.RELT 84957	NM_151.1	7337639 9 339747 4 RELT	-	6.709	5.780
			0.33	5.83	
ENSG0.NUP58 9818	13	2530152 2 536539@ucle	0.349	5.040 5.791	6.476
ENSG0.IRF2BP 6 4207	NM_0214	7702454 3 702870 & nter	_	6.989	7.496
			0.44	5.73	
ENSG0.ZNF512 B 7473	NM 0220	6395670 \$ 396993 Ø inc	_	5.529	8.722
	_		0.33	5.66	
ENSG0.ZNF48690649	NM 0519	2016721 2 020048 8 inc			2.586
ENSG0.EPS8L264787	11	694438 727727 EPS8	_	6.161	9.891
21.5 00.21 502201.01				5.61	0.001
ENSG0.CSNK1E454	NM 1522	3829069 \$ 831808 ¢ asei			1.082
LIVEGU.CEIVITIE	1111_1022	3023003 9 031000 4 asci		5.58	1.002
ENSG0.CORO1NA	16	3018350 3 018495 C ORO			0.851
EINDGU.DORUINA	10	9010990 9 01049910ORO		5.61	J.001
			•••		

7.2.5 Gather

Figure 7为图 all DEGs of GSE datasets 概览。

(对应文件为 Figure+Table/all-DEGs-of-GSE-datasets.pdf)

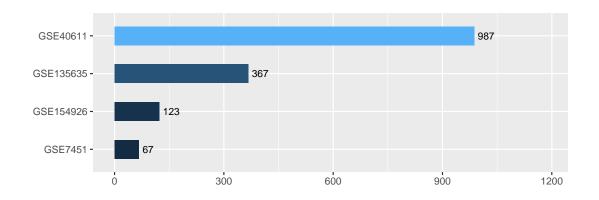


Figure 7: All DEGs of GSE datasets

7.3 PPI STRINGdb

Figure 8为图 intersection of SS DEGs with XMGJ targets 概览。

(对应文件为 Figure+Table/intersection-of-SS-DEGs-with-XMGJ-targets.pdf)

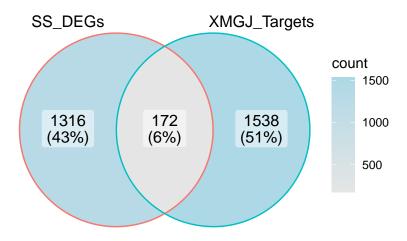


Figure 8: Intersection of SS DEGs with XMGJ targets

Figure 9为图 PPI networks 概览。

(对应文件为 Figure+Table/PPI-networks.pdf)

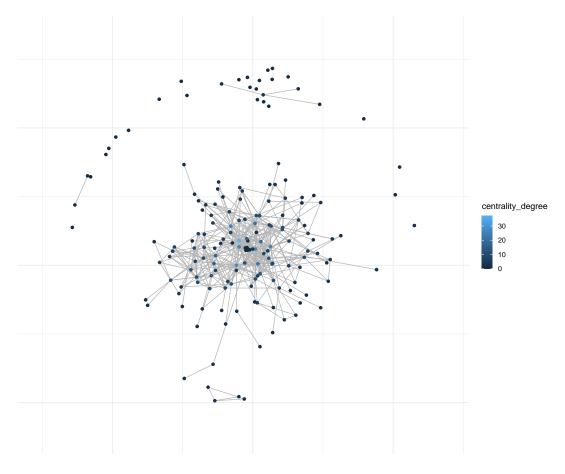


Figure 9: PPI networks

Figure 10为图 MCC top30 概览。

(对应文件为 Figure+Table/MCC-top30.pdf)

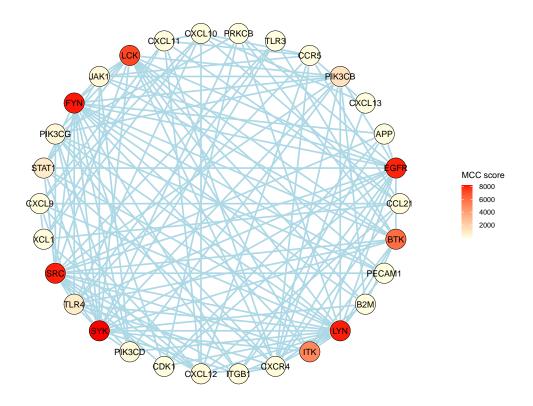


Figure 10: MCC top30

7.4 富集分析

Figure 11为图 KEGG Enrichment 概览。

(对应文件为 Figure+Table/KEGG-Enrichment.pdf)

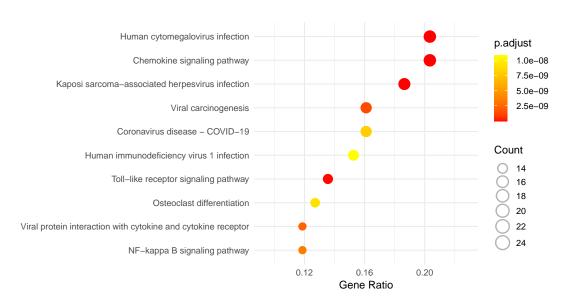


Figure 11: KEGG Enrichment

Table 11为表格 tables of KEGG Enrichment 概览。

(对应文件为 Figure+Table/tables-of-KEGG-Enrichment.xlsx)

注:表格共有 233 行 9 列,以下预览的表格可能省略部分数据;表格含有 233 个唯一'ID'。

Table 11: Tables of KEGG Enrichment

ID	Descr	GeneR	BgRatio	pvalue	p.adjust	qvalue	geneID	Count
hsa04062	Chemo	24/118	192/8622	5.925	1.380	5.613	5291/	24
$\rm hsa05163$	Human	24/118	225/8622	2.350	2.738	1.113	5291/	24
hsa05167	Kapos	22/118	194/8622	1.091	8.475	3.446	5291/	22
hsa04620	Toll	16/118	108/8622	9.575	5.577	2.267	5291/	16
hsa05203	Viral	19/118	204/8622	2.918	1.360	5.530	5291/	19
hsa04061	Viral	14/118	100/8622	6.090	2.365	9.616	6387/	14
hsa04064	NF-ka	14/118	104/8622	1.047	3.485	1.417	695/3	14
hsa05171	Coron	19/118	232/8622	2.736	7.971	3.241	5291/	19
hsa04380	Osteo	15/118	135/8622	3.498	9.056	3.682	695/2	15
hsa05170	Human	18/118	212/8622	4.708	1.082	4.400	5291/	18
hsa05161	Hepat	16/118	162/8622	5.109	1.082	4.400	5291/	16
hsa05417	Lipid	18/118	215/8622	5.924	1.150	4.677	5291/	18
hsa05164	Influ	16/118	171/8622	1.142	2.047	8.325	5291/	16
hsa05169	Epste	17/118	202/8622	1.719	2.861	1.163	695/5	17
hsa05134	Legio	10/118	56/8622	3.289	5.110	2.077	841/4	10

Figure 12为图 GO enrichment 概览。

(对应文件为 Figure+Table/GO-enrichment.pdf)

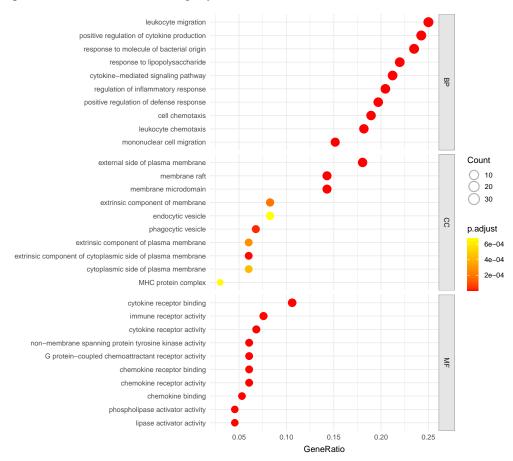


Figure 12: GO enrichment

'Tables of GO enrichment' 数据已全部提供。

(对应文件为 Figure+Table/Tables-of-GO-enrichment)

注:文件夹 Figure+Table/Tables-of-GO-enrichment 共包含 3 个文件。

- $1.\ 1_{\rm BP.csv}$
- 2. 2_CC.csv
- 3. 3_MF.csv

The Cellular Component (CC), the Molecular Function (MF) and the Biological Process (BP).

7.5 MCC top30 和 Chemokine signaling pathway

Figure 13为图 intersection of MCC top30 and the hitted genes of Chemokine pathway 概览。

(对应文件为Figure+Table/intersection-of-MCC-top30-and-the-hitted-genes-ofChemokine-pathway.pdf)

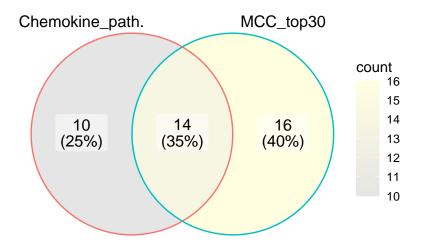


Figure 13: Intersection of MCC top30 and the hitted genes of Chemokine pathway

7.6 Xuanmai Ganju 靶向 Chemokine signaling pathway 治疗 Sjogren syndrome 的成分

Table 12为表格 tables of components of XMGJ targeting Chemoking signaling for treating SS 概览。

(对应文件为 Figure+Table/tables-of-components-of-XMGJ-targeting-Chemoking-signaling-for-treating-SS.xlsx)

注: 表格共有 68 行 9 列,以下预览的表格可能省略部分数据; 表格含有 20 个唯一'Ingredient.name'。

Table 12: Tables of components of XMGJ targeting Chemoking signaling for treating SS

Ingre1	Herb	Ingre3	Ingre4	Targe5	Targe6	Datab	Paper.id	
HBIN0	GAN CAO	18bet	AJ-72	HBTAR	SRC	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CCR1	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CCR5	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CCR6	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CCR7	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CXCL8	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CXCR2	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CXCL10	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CXCL9	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CXCL12	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CXCR4	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CXCR6	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	GNG2	NA	NA	
HBIN0	JIE GENG	adeni	NA	HBTAR	CCR2	NA	NA	
HBIN0	MAI DONG	adeni	NA	HBTAR	CCR1	NA	NA	

Ingre1	Herb	Ingre3	Ingre4	Targe5	Targe6	Datab	Paper.id	

Reference

- 1. Fang, S. et al. HERB: A high-throughput experiment- and reference-guided database of traditional chinese medicine. *Nucleic Acids Research* 49, D1197–D1206 (2021).
- 2. Wu, T. et al. ClusterProfiler 4.0: A universal enrichment tool for interpreting omics data. The Innovation 2, (2021).
- 3. Szklarczyk, D. et al. The string database in 2021: Customizable proteinprotein networks, and functional characterization of user-uploaded gene/measurement sets. Nucleic Acids Research 49, D605–D612 (2021).
- 4. Ritchie, M. E. et al. Limma powers differential expression analyses for rna-sequencing and microarray studies. Nucleic Acids Research 43, e47 (2015).