

前言

2017年11月29日 下午 03:11

INPUT

最主要的檔案:

resource-mentions.tsv 紀錄那些resource(rid)出現在哪些論文(mentionid or mentionid_int)內

resource-mentions-relationships.tsv 紀錄那些resource曾共同出現在那些論文內

resource-metadata.tsv 紀錄每個resource的meta information(如全名 相關網址)

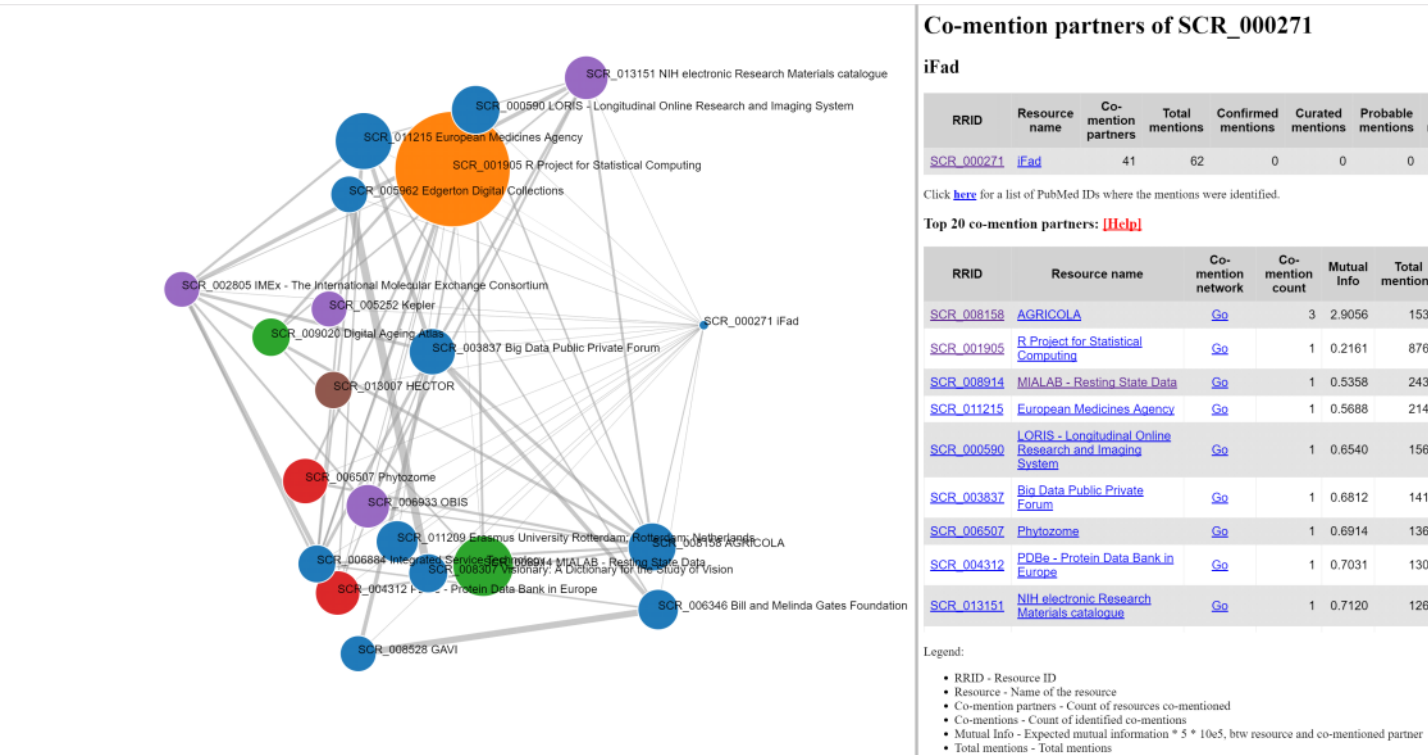
資料更正(除錯)檔:

exclusion.tsv 紀錄需要移除的resource id (要從上述的mention & mention-relationship檔刪除相關資料)

resource-duplicates.tsv 紀錄一些相同resource卻不同(多餘)id的資料集合

OUTPUT

為每個resource產生 resource co-mention network (graph)還有相關的co-mention table



resource-mentions.tsv

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紀錄每個resource出現在那些論文內

	A	B	C	D	E	F	G	H	I	J	K	L
1	id	uid	rid	mentionid	rating	timestamp	mentionid_int	input_source	confidence	vote_sum	snippet	
2		1 NULL		2 PMID:9866185	none	1444789212	9866185	rdw	0.2	0		
3		2 NULL		2 PMID:9860986	none	1444789212	9860986	rdw	0.2	0		
4		3 NULL		2 PMID:979569	none	1444789212	979569	rdw	0.2	0		
5		4 NULL		2 PMID:9658713	none	1444789212	9658713	rdw	0.2	0		
6		5 NULL		2 PMID:9615423	none	1444789212	9615423	rdw	0.2	0		

主要欄位:

- rid: resource的id
- mentoid: 論文id
- mentionid_int: integer格式的論文id
- confidence: 介於0~1, 越大代表該resource出現在論文的可信度越高 (resource mention多半是透過information extraction (IE) 的程式擷取, 所以有些可能有錯, 而confidence = 1是人工判斷過的, 可信度最高)

主要產出資訊:

- 統計出每個resource的mention count(出現在多少篇論文內), Resource co-mention graph與table會使用到該值調整resource node的大小與table欄位 (如右例圖)

因為UCSD會不斷update這個檔案(修正IE程式擷取資料的錯誤), 請先了解秋中設計的db格式, 並設計Python程式來處理tsv檔以產生相關資料.



iFad

RRID	Resource name	Co-mention partners	Total mentions	Confirmed mentions
SCR_0000271	iFad	44	62	0

Click [here](#) for a list of PubMed IDs where the mentions were identified.

需要注意事項:

- exclusion.tsv紀錄了一些需要移除的resources, 要確定最後的db沒有這些resources
- resource-duplicates.tsv紀錄了一些多餘的resource ids, 要確定將這些多餘id的mentions (co-mention)整合到單一resource id下

紀錄圖中resource pair曾出現在那些論文內

#	A	B	r1	r2	count	comentions	count_hc	comentions_hc
1	1	1	8673	1	PMID:21505475	1	PMID:21505475	1
2	1	1	4	1	PMID:27990286	1	PMID:27990286	1
3	1	1	6917	1	PMID:22438826	0		0
4	1	1	4455	1	PMID:22859986	0		0
5	1	1	7817	1	PMID:22438826	1	PMID:22438826	1
6	1	1	3145	1	PMID:22859986	1	PMID:22859986	1
7	1	1	8426	1	PMID:27990286	1	PMID:27990286	1
8	1	1	4519	1	PMID:22438826	0		0
9	1	1	1905	1	PMID:27119341	1	PMID:27119341	1
10	1	1	1554	1	PMID:22438826	1	PMID:22438826	1
11	1	1	11840	1	PMID:22438826	0		0
12	1	1	8117	1	PMID:27990286	1	PMID:27990286	1
13	1	1	8982	1	PMID:22438826	0		0
14	1	1	3033	1	PMID:22438826	1	PMID:22438826	1
15	1	1	11417	1	PMID:22438826	1	PMID:22438826	1
16	1	1	4761	1	PMID:25847540	1	PMID:25847540	1
17	1	1	2110	1	PMID:22438826	1	PMID:22438826	1
18	1	1	9211	1	PMID:22438826	0		0
19	1	1	4727	1	PMID:22438826	1	PMID:22438826	1
20	1	1	4286	3	PMID:25847540,PMID:24843691,PMID:22859986	3	PMID:25847540,PMID:24843691,PMID:22859986	3
21	1	1	8639	1	PMID:24647409	0		0
22	2	2	10241	1	PMID:23431087	0		0

主要欄位:

- r1, r2: resource pair的resource id
- count: 共同出現的論文篇數
- comentions: 共同出現的論文ids

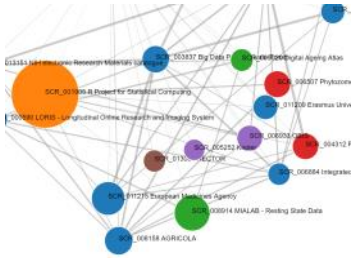
主要產出資訊:

- 結合resource-mentions.tsv中resource出現的次數與共同出現的次數來算“**預期**”resources的關聯性，關聯性數值與**expected mutual information (MI)**，該值用來決定resource graph上link的粗細(見右圖)
- Resource table會列出與當前resource關係最強的resources，且秀出其co-mention count和MI值

因為UCSD會不斷update這個檔案(修正因爬取資料的錯誤，
請先了解表中設計的db格式，
並設計Python程式來處理tsv檔以產生相關資料。

需要注意事項:

- exclusion.tsv記錄了一些需要移除的resources，要確定最後的db沒有這些resources的co-mentions
- resource-duplicates.tsv記錄了一些多餘的resource ids，要確定將這些多餘ids的mentions整合到單一resource id下，且相對應的co-mention count與comention PMID要調整



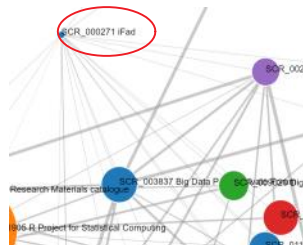
Top 20 co-mention partners: [JHclh]

RRID	Resource name	Co-mention network	Co-mention count	Mutual Info.	Total mentions
SCR_008158	AGRICOLA	Go	3	2.9056	1531
SCR_001905	R Project for Statistical Computing	Go	1	0.2161	8769
SCR_008914	MALAB - Rastling State Data	Go	1	0.5358	2430
SCR_011215	European Medicines Agency	Go	1	0.5688	2148
SCR_000569	LOUIS - Longitudinal Online Research and Imaging System	Go	1	0.6540	1567
SCR_003837	Bio Data Public Private Forum	Go	1	0.6612	1418
SCR_006507	Physiozone	Go	1	0.6914	1366
SCR_004312	PDBe - Protein Data Bank in Europe	Go	1	0.7031	1309
SCR_013351	NIH electronic Research Materials catalogue	Go	1	0.7125	1267

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A	B	C	D	E	F	G	H	I	J	K	L
e_uid	resource_name	abbreviation	description	url	see_full_record_url	see_full_record	alternative_ids	original_id	canonical_id		
1	TransGenic		A commercial antib	https://www.transgenic.com/browse/resources/SCISCR_000001			nix_152482	SCR_000001			
2	monarch-ontologies	Monarch Ontologies	The set of ontologies	http://purl.obolibrary.org/ontology/scienc	https://www.bioscience.com/browse/resources/SCISCR_000002		nix_152901	SCR_000002			
3	Sarah Cannon Researc	SCRI	A global cancer insti	http://sarahcannonres	https://www.bioscience.com/browse/resources/SCISCR_000003		nix_158000	SCR_000003			
4	GE Healthcare		A commercial antib	http://www.gelifescier	https://www.bioscience.com/browse/resources/SCISCR_000004		nix_152368	SCR_000004			
5	Neuroshare - Open data specifications and s		Neuroshare aims to	http://neuroshare.sour	https://www.bioscience.com/browse/resources/SCISCR_000005		nif-0000-00023	SCR_000005			
6	University of Algarve; UAlg		A young state univer	https://www.ualg.pt/en	https://www.bioscience.com/browse/resources/SCISCR_000006		nix_157657	SCR_000006			
7	G Biosciences		A commercial antib	https://www.gbiosci	https://www.bioscience.com/browse/resources/SCISCR_000007		nix_152367	SCR_000007			
8	University at Albany SUNY Labs and Facilit		A facility that conda	https://www.albany.edu	https://www.bioscience.com/browse/resources/SCISCR_000008		SciEx_4303	SCR_000008			
9	ncdfFlow		Software package th	https://www.ncdfcondu	https://www.bioscience.com/browse/resources/SCISCR_000009		OMICS_05617	SCR_000009			
10	Computational Neuroscience on the Web		An annotated index	(http://theme.eartlink	https://www.bioscience.com/browse/resources/SCISCR_000010		nif-0000-01007	SCR_000010			
11	Leica DMRE Fluoresce	Leica DMRE micros	Microscope that enal	http://www.unilfaco	https://www.bioscience.com/browse/resources/SCISCR_000011		SciRes_000155	SCR_000011			
12	Offline Sorter	OPS	Offline spike sorting	http://www.plexon.co	https://www.bioscience.com/browse/resources/SCISCR_000012		nix_1558484	SCR_000012			
13	BSmodot-align		A statistics and align	https://github.com/Bie	https://www.bioscience.com/browse/resources/SCISCR_000013		OMICS_018486	SCR_000013			
14	University of Pittsburgh Pitt CCNMD, Conte C	The Conte Center fo	https://www.ccnmd.pit	https://www.bioscience.com/browse/resources/SCISCR_000014			nix_144496	SCR_000014			
15	4Peaks	4Peaks	Software application	https://nucleobytes	https://www.bioscience.com/browse/resources/SCISCR_000015		OMICS_01015	SCR_000015			
16	CSDeconv	CSDeconv	A software applicat	https://crab.rutgers	edu/https://www.bioscience.com/browse/resources/SCISCR_000016		OMICS_00436	SCR_000016			
17	Tablet	Tablet	A lightweight, high-y	https://bioinf.scri	ac.uk/https://www.bioscience.com/browse/resources/SCISCR_000017		OMICS_00896	SCR_000017			
18	Midwest Transplant MNTN		An organization that	https://www.mwtn	.org/https://www.bioscience.com/browse/resources/SCISCR_000018		nix_87553	SCR_000018			
19	NeuroTribes		Steve Silberman's pe	https://blogs.plos	.org/h/https://www.bioscience.com/browse/resources/SCISCR_000019		nix_91543	SCR_000019			
20	Ludwig Boltzmann Cl	Ludwig Boltzmann Cl	The projected cluster	https://oc.lbg.ac.at/	https://www.bioscience.com/browse/resources/SCISCR_000020		nix_143958	SCR_000020			

- e_uid, see_full_record: resource id
- resource_name: 資源名稱
- abbreviation: 資源名稱縮寫
- url: 該資源的官方網址
- see_full_record_url: 該資源於scicrunch的說明網址



- Resource graph與table會使用這些資訊來產生相關圖表 (見右圖)

因為UCSD會不斷update這個檔案(修正正程式擷取資料的錯誤),
請先了解秋中設計的db格式,
並設計Python程式來處理tsv檔以產生相關資料.

- 小心欄位會有missing value!! (如 1 TransGenic的abbreviation missing)

RRID	Resource name	Co-mention network
SCR_008158	AGRICOLA	Go
SCR_001905	R Project for Statistical Computing	Go
SCR_008914	MIALAB - Resting State Data	Go
SCR_011215	European Medicines Agency	Go
SCR_000590	LORIS - Longitudinal Online Research and Imaging System	Go
SCR_003837	Big Data Public Private Forum	Go
SCR_006507	Phylozome	Go
SCR_004319	PDRe - Protein Data Bank in Europe	Go

exclusion.tsv

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不需產生資料的 resource ids

	A	B	C	D
	1	SCR_000001	TransGenic	Commercial antibody supplier
	4	SCR_000004	GE Healthcare	Commercial antibody supplier
	7	SCR_000007	G Biosciences	Commercial antibody supplier
	69	SCR_000069	GeneTex	Commercial antibody supplier
	70	SCR_000070	Genemed	Commercial antibody supplier
	215	SCR_000215	Full Moon BioSyste	Commercial antibody supplier
	314	SCR_000314	DB BioTech	Commercial antibody supplier
	382	SCR_000382	SunnyLab	Commercial antibody supplier
	1108	SCR_001108	Academy Biomedica	Commercial antibody supplier
0	1129	SCR_001129	NewEast Bioscience	Commercial antibody supplier
L	1130	SCR_001130	MitoScience	Commercial antibody supplier
2	1133	SCR_001133	Hytest	Commercial antibody supplier
3	1134	SCR_001134	BioLegend	Commercial antibody supplier
4	1136	SCR_001136	Aves Labs	Commercial antibody supplier
5	1137	SCR_001137	Atlas Antibodies	Commercial antibody supplier
5	1139	SCR_001139	Abazyme	Commercial antibody supplier
7	1141	SCR_001141	Phoenix Pharmaceut	Commercial antibody supplier
3	1220	SCR_001220	ChanTest	Commercial antibody supplier
9	1224	SCR_001224	Covance	Commercial antibody supplier
0	1287	SCR_001287	Merck	Commercial antibody supplier
L	SCR_001	Integrated Animals	Interated animal family	
2	SCR_001	Integrated Models	Interated animal family	
3	1932	SCR_001932	Immune Technology	Commercial antibody supplier
4	2087	SCR_002087	Icosagen AS	Commercial antibody supplier
5	SCR_002	Integrated	Interated animal family	
5	2891	SCR_002891	GenScript	Commercial antibody supplier
7	2930	SCR_002930	Genox Corpooration	Commercial antibody supplier
3	SCR_003	Integrated Grants	Interated animal family	
9	3145	SCR_003145	GeneCopoeia	Commercial antibody supplier
0	3200	SCR_003200	Gen Probe	Commercial antibody supplier

數字id missing!!

不需為這些resource產生table & graph,
也不能讓這些resource出現在其他resource的graph & table內
保險起見, 在db內把他們mention co-mention的紀錄移除,

因為UCSD會不斷update這個檔案(修正E程式擷取資料的錯誤),
請先了解秋中設計的db格式,
並設計Python程式來處理tsv檔以產生相關資料.

需要注意事項:

- 小心欄位會有missing value!!

resource-duplicates.tsv

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紀錄著本是同一resource, 卻因為一些緣故產生多組(餘)的resource id, 需依照該檔來將mention & co-mention整合(修正)

會影響資料一致性...需小心處理!!!

	A	B	C	D	E	F	G	H	I	J
1	id	uid	id1	id2	type1	type2	reltype_id	canon_id	timestamp	
2	21081		31497	SCR_008406	SCR_013567	res	res	1	1	1.45E+09
3	21440		31497	SCR_001915	SCR_005576	res	res	1	1	1.45E+09
4	21467		31497	SCR_010243	SCR_003614	res	res	1	1	1.45E+09
5	21585		31497	SCR_007058	SCR_007057	res	res	1	1	1.45E+09
6	21587		31497	SCR_013733	SCR_014203	res	res	1	1	1.45E+09
7	21588		31497	SCR_007394	SCR_000951	res	res	1	1	1.45E+09
8	21622		511	SCR_002823	SCR_007368	res	res	1	1	1.45E+09
9	21652		31537	SCR_011249	SCR_005074	res	res	1	1	1.46E+09
10	21659		31537	SCR_004513	SCR_010892	res	res	1	1	1.46E+09
11	21664		31537	SCR_008302	SCR_010795	res	res	1	1	1.46E+09
12	21666		31497	SCR_000325	SCR_014216	res	res	1	1	1.46E+09
13	21667		31537	SCR_013504	SCR_013506	res	res	1	1	1.46E+09
14	21668		31537	SCR_008249	SCR_011948	res	res	1	1	1.46E+09
15	21673		31497	SCR_007030	SCR_005244	res	res	1	1	1.46E+09

主要欄位:

- Id1, id2: 多餘的resource id

整合(修正)方式:

duplicate會造成一個resource的mentions與co-mentions的紀錄四散,

如一個resource如果有3個ids,

則應該把這三個ids的mentions, co-mentions整合

首先要確定這些duplicate ids有無串連,

如 SCR_1 跟 SCR_5 duplicate,

SCR_5又跟 SCR_10 duplicate,

要將所有相關(串聯)的ids進行整合,

即 SCR_1, SCR_5 & SCR_10 是代表同一個resource,

假設最後這三個ids都用SCR_1來統稱(代表) - (先前挑代表id的方式有點問題, 我目前正在等 UCSD答覆如何挑對的代表id...)

修正mention -

假如SCR_5曾出現在PMID:9000, 但SCR_1沒有, 則要修正成SCR_1有出現在PMID:9000, 且要更正SCR_1的total_mention次數 (加一)

但若SCR_1也有出現在PMID:9000, 則不修正, 且捨棄SCR_5出現在PMID:9000的資料

****若有修正, 記得要更正total_mention次數****

修正co-mention -

若SCR_5 與 SCR_2 曾一起出現在PMID:3,PMID:5, PMID:7, 而SCR_1 跟 SCR_2曾一起出現在PMID:1,PMID:5, PMID:4,

則要刪掉SCR_5與SCR_2的co-mention, 且修正SCR_1與SCR_2的co-mention為PMID:1, PMID:3, PMID:4, PMID:5, PMID:7,

**** 要同時修正SCR_1與SCR_2的co-mention count為 5(原本為3) ****

若SCR_5 與 SCR_4 曾一起出現在PMID:11,PMID:19, PMID:37, 而SCR_1 跟 SCR_4不曾一起出現過,

則要刪掉SCR_5與SCR_4的co-mention, 且修正SCR_1與SCR_4的co-mention為PMID:11, PMID:19, PMID:37

**** 要同時修正SCR_1與SCR_4的co-mention count為 3(原本沒這筆資料) ****

若發現SCR_5與SCR_1有一起出現的PMIDs, 則可以delete這些資料,

**** 要check資料的正確性, 如SCR_5與SCR_1一起出現在PMID:200, PMID:400,**

則需要看"修正後的mention"是否有記錄到SCR_1出現在PMID:200, PMID:400,

如果有漏...表示資料不一致!! **

因為UCSD會不斷update這個檔案(修正IE程式擷取資料的錯誤),
請先了解秋中設計的db格式,
並設計Python程式來處理tsv檔以產生相關資料.

需要注意事項:

- 目前還在釐清正確的id置換法則

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先前秋中已產生30個clusters of resources,
每個cluster存成一個文字檔 (0.txt, 1.txt, ..., 29.txt)

主要欄位:

- RRID: resource id
- Resource Name: resource名稱
- total mention count: resource出現的在論文篇數

- Resource graph內同cluster的resource會標示同個顏色 (見右圖)

用現行的package SLM

<http://www.ludowaltman.nl/slm/>

- 我們先把資料(tsv & db)整理好, 確定無誤後再來研究SLM的使用方式,



RRID	Resource name	Co-mention partners	Total mentions	Confirmed mentions	Curated mentions	Probable mentions
RSC 000071	Fad	41	62	0	0	0

Click [here](#) for a list of PubMed IDs where the mentions were identified.

Top 20 co-mention partners: [Help](#)

RRIID	Resource name	Co- mension code	Co- mension count	Manual Info	Total mensions
SCE_000008	AGRICOLA	Go	3	2,9256	153
SCE_000009	B.Project for Statistical Computation	Go	1	1,2161	879
SCE_000014	MOLAB - Reading Steps Data	Go	1	1,5358	245
SCE_000025	Program: Medication Appliance	Go	1	1,5688	214
SCE_000028	LORIS - Linguistical Online Research and Inquiry System	Go	1	1,0540	106
SCE_000037	Big Data Public: Protein 2045	Go	1	1,0682	1411
SCE_000052	Phylozone	Go	1	1,0594	130
SCE_000071	PDBE - Protein Data Bank in Europe	Go	1	1,0701	130
SCE_000113	NPL electronic Research Materials collection	Go	1	1,07120	126

Legend:

- RRID - Resource ID
- Resource - Name of the resource
- Co-mention partners - Count of resources co-mentioned
- Co-mentions - Count of identified co-mentions
- Mental info - Expected mental information * 5 * 10e5, how resource and co-mentioned partner
- Total mentions - Total mentions