

Evaluating utility of subject headings in a data repository: A preliminary finding from a data search log and record classification

Presented by:

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Contributors:

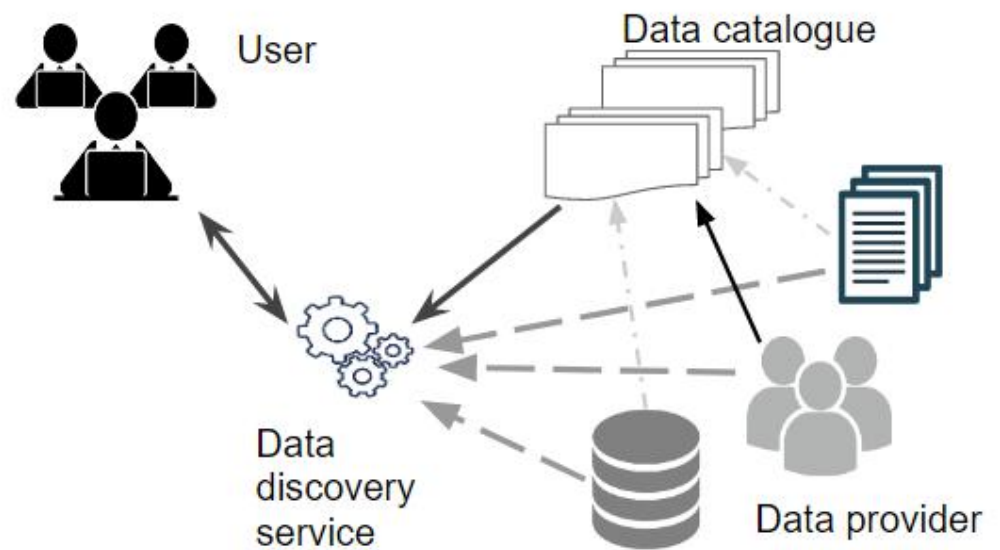
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Ying-Hsang Liu, University of Southern Denmark
Jenny Xiuzhen Zhang, RMIT University, Australia

NKOS, 10 Sept. 2020



Outlines

- A background about the studied data catalogue: Research Data Australia
- Log analysis: the usage of subject headings
- Experiments on data record classification
- Future work



Research Data Australia - A National Data Catalogue

Research Data Australia

Find data for research

Browse By Subjects

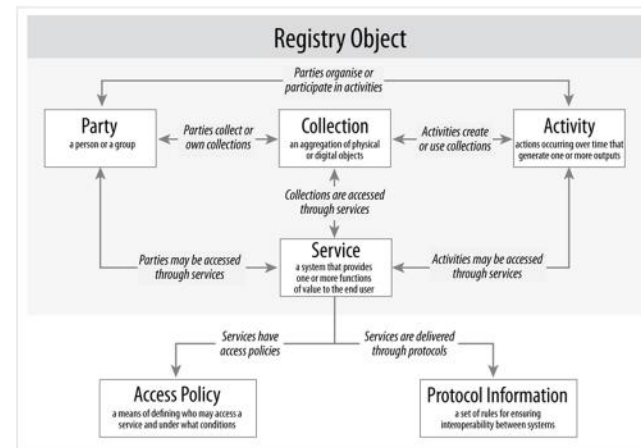
144K+ metadata records of dataset
60K+ research grants

99 Contributors

Griffith University, Deakin University, RMIT University, NCRIS, ARDC, AARDAC



Schema: The Registry Interchange Format - Collections and Services (RIF-CS, ISO 2146:2010)



Types of subject vocabularies

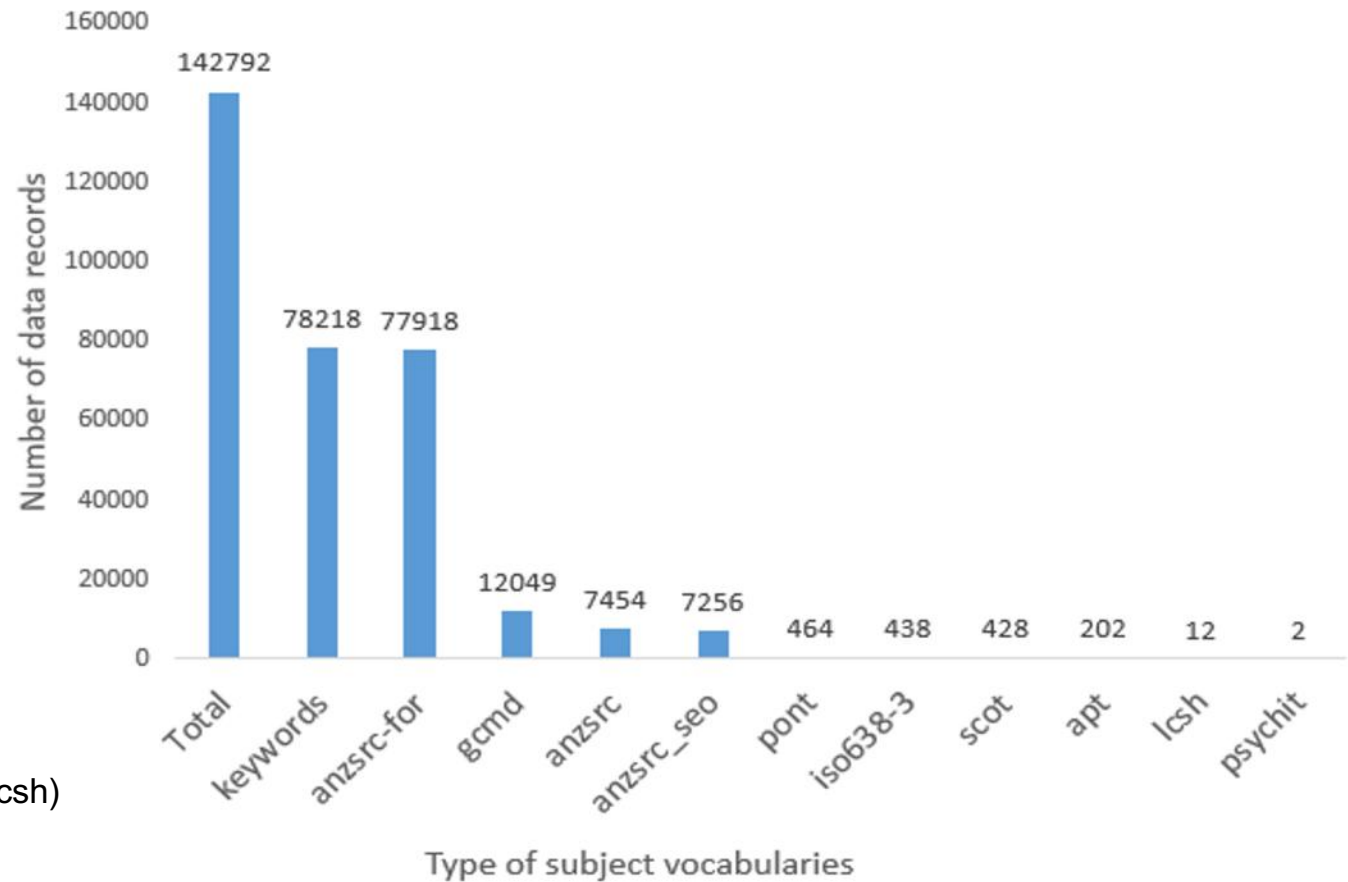
Anzsrc-for: The Australian and New Zealand Standard Research Classification (ANZSRC, fields of research)

Global change master directory (GCMD) keywords

Australian Pictorial Thesaurus (apt)

Thesaurus of Psychological Index Terms (psychit)

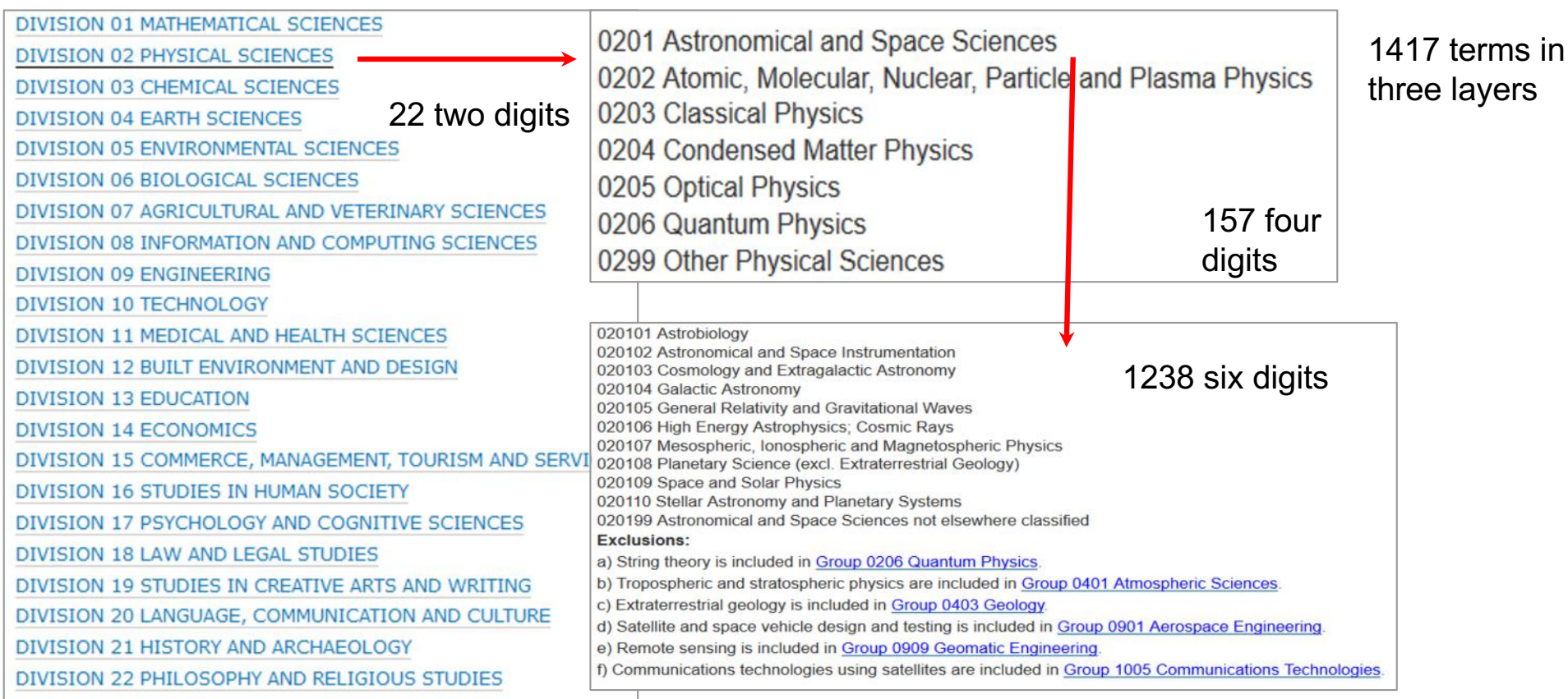
Library of Congress Subject Headings (lcsh)



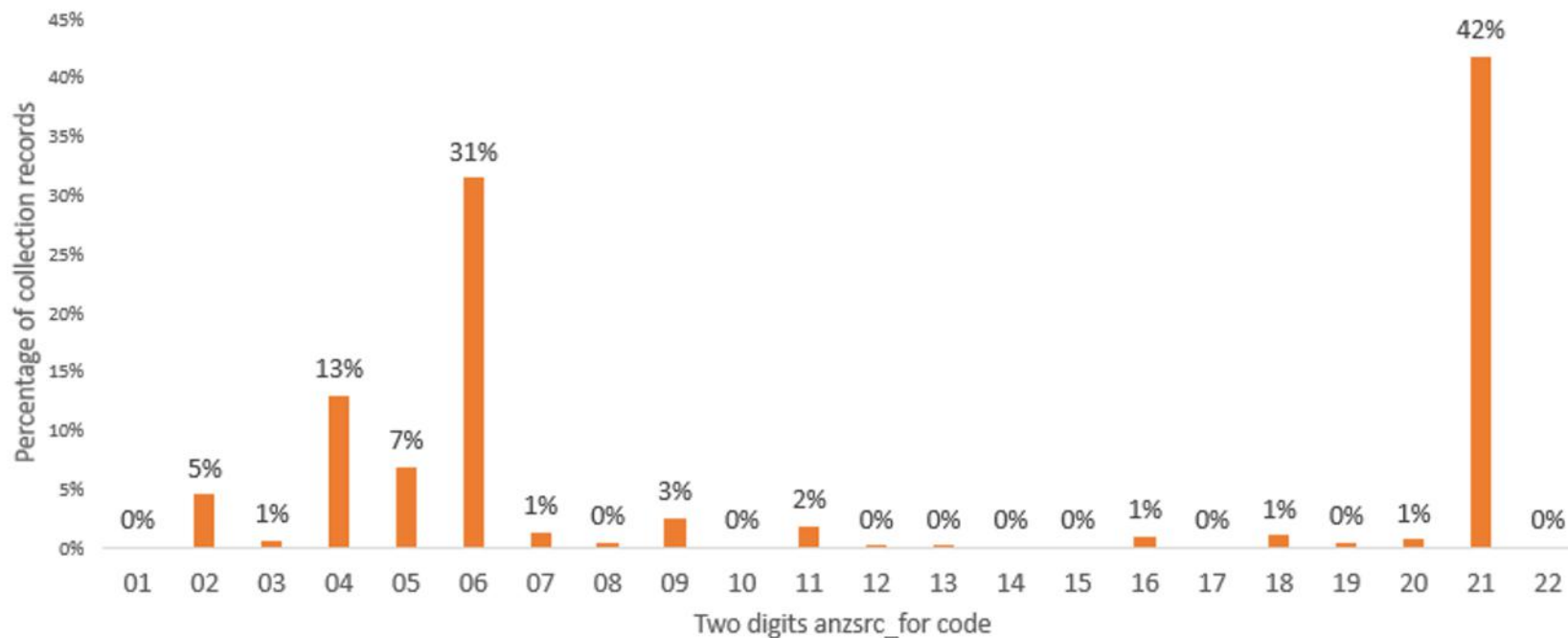
Anzsrc-for: The Australian and New Zealand Standard Research Classification - Fields of Research

- ANZSRC ensures that R&D statistics collected are useful to governments, educational institutions, international organisations, scientific, professional or business organisations, business enterprises, community groups and private individuals in Australia and New Zealand.
- ANZSRC-FoR include major fields and related sub-fields of research and emerging areas of study investigated by businesses, universities, tertiary institutions, national research institutions and other organisations.

Anzsrc-for: The Australian and New Zealand Standard Research Classification - Fields of Research)



Number of records per anzsrc-for two digits



04: Earth Sciences

06: Biological Sciences

21: History and Archaeology

Search interface

All text strings (including subject headings) are indexed.

The screenshot displays a search interface with a search bar at the top containing the text 'gene'. Below the search bar, it indicates '766 results (48 milliseconds)' and 'Records selected: 0'. The interface is divided into several sections:

- Current Search:** Shows the search term 'gene' and options to 'Save Search' or 'Clear Search'.
- Refine search results:** A sidebar with filters for 'Type', 'Subject', and 'Data Provider'.
 - Type:** Data (762), Software (4).
 - Subject:** Biological Sciences (330), Medical And Health Sciences (240), Agricultural And Veterinary... (26), Environmental Sciences (21), Information And Computing S... (13).
 - Data Provider:** Monash University (235), Australian Ocean Data Network (86).
- Search Results:** A list of search results, each with a checkbox and a title:
 - Gene Sherman Collection** (Museum Metadata Exchange). Description: A collection of Japanese fashion owned and worn by Gene Sherman from the late 1980s to 2007.
 - Disease gene prediction database** (Deakin University). Description: This database includes gene predictions for disease phenotypes based on published Genome-Wide Association Data.
 - Play to Cure: Genes in Space** (Atlas of Living Australia). Description: We know that faults in our genes can lead to cancer cells forming.
 - Lactation related gene expression data** (Deakin University). Description: RNA sequencing and gene expression data related to lactation.
 - Antibiotic resistance gene cassettes** (University of New South Wales). Description: Gene cassettes and cassette arrays.

Subject headings


The screenshot shows a search interface with the following components:

- Search Bar:** Contains the text "All Fields" and "gene". A search button with a magnifying glass icon is present.
- Filters:** A checkbox labeled "Publicly accessible online" is checked.
- Results Summary:** "766 results (48 milliseconds)", "Records selected: 0", and buttons for "Save Records" and "Export".
- Current Search:** A sidebar on the left showing "All Fields" with "gene" entered. It includes "Save Search" and "Clear Search" buttons.
- Refine search results:** A section with "Add more keywords" and a "Go" button. It lists "Type" filters: "Data" (762) and "Software" (4).
- Subject Facet Filter:** A table listing subject categories with their respective counts, highlighted with a red box and labeled "2. Facet filter".

Subject	Count
Biological Sciences	330
Medical And Health Sciences	240
Agricultural And Veterinary...	26
Environmental Sciences	21
Information And Computing S...	13
- Advanced Search Panel:** A sidebar on the right with a "Subject" filter selected. It lists various subject categories, with "Vocabulary ANZSRC FOR -" circled in red and labeled "1. Advanced search".


Subject	Count
Agricultural And Veterinary Sciences	26
Biological Sciences	330
Built Environment And Design	1
Chemical Sciences	2
Commerce, Management, Tourism And Services	1
Earth Sciences	1
Economics	1
Education	0
Engineering	2
Environmental Sciences	21
History And Archaeology	1
Information And Computing Sciences	13
Language, Communication And Culture	1
Law And Legal Studies	0
Mathematical Sciences	1
- Search Results:** A list of search results including "Gene Sherman Collection", "Disease gene prediction database", "Play to Cure: Genes in Space", "Lactation related gene expression d...", and "Antibiotic resistance gene cassettes".


Record view



Disease gene prediction database

Deakin University
Dr Merridee Wouters (Aggregated by) Mr Martin Oti (Aggregated by)

 Dataset

 Viewed: 946 Accessed: 15

[Access the data](#)

[Cite](#) [Save to MyRDA](#)

Licence & Rights:
Other [view details](#)

Access:
Other [view details](#)

Contact Information
Postal Address:
School of Life and Environmental Sciences,
Deakin University, 75 Pigdons Road, Waurn
Ponds, Victoria 3216 Australia

Full description

This database includes gene predictions for disease phenotypes based on published Genome-Wide Association Data. May be used to choose primers for phenotype-specific resquencing of patient DNA.

For each prediction for following data is listed: phenotype, predicted gene, significant SNP, datasource, datasource reference.

Notes

The data was generated by a computer from clinical data, and some data from HuGE (<http://hugenavigator.net/HuGENavigator/home.do>) was used. The data is organised within a searchable

Subjects

**3. Facet search
(vocabulary + keyword)**

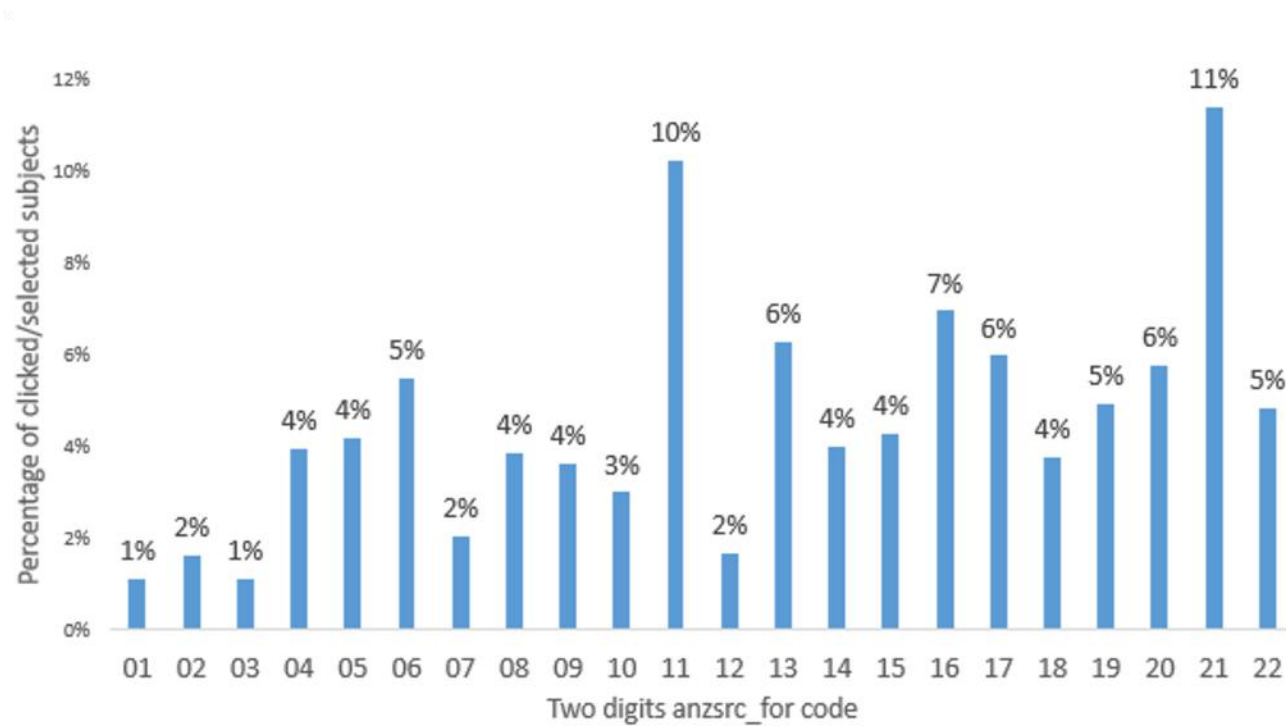
[Biological Sciences](#) | [Clinical Health \(Organs, Diseases and Abnormal Conditions\)](#) | [Genetics](#) | [Genetics Not Elsewhere Classified](#) | [Health](#) | [Inherited Diseases \(Incl. Gene Therapy\)](#) | [database](#) | [genetic databases](#) | [genome-wide association study](#) | [humans](#) | [polymorphism](#) | [protein disease/genetics](#) | [single nucleotide](#) | [software](#) |

10

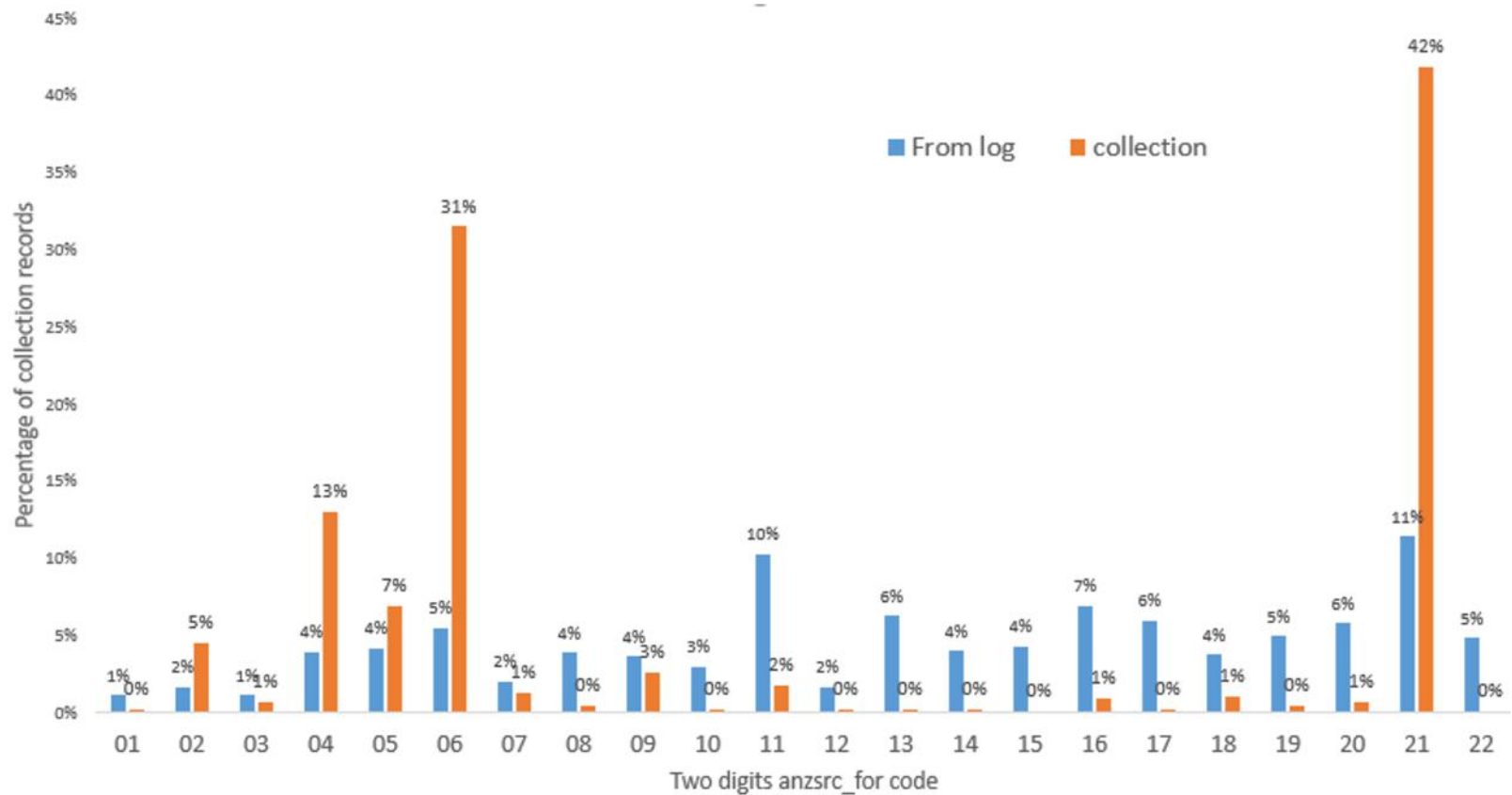
Log analysis: the usage of subject headings

- Transaction log: one year (2019) of activities recorded from the RDA catalogue
- About 2 million entries/activities, 63% from Australia
- About 496,739 sessions (with 30 minutes duration from the same IP address)
- 37,056 sessions have at least a search event (keyword search, advanced search, subject (facet) filter, subject search)
- 4668 (12.6%) of search sessions involved filters/search with the ansrc-for subjects, only 45 (0.1%) with gcmd subject

Subject usages per anzsrc-for two digits code



Subject distribution among clicks and the collection



Log analysis: the usage of subject headings

- There is less bias in user's behaviour of applying subject headings, compared to the content bias toward a few subject headings.
- However, this log shows low usage of subject headings
- Exploring causes
 - Further log analysis, e.g. correlation between subject usage and
 - query types
 - domain knowledge
 - search quality
 - Interface design
 - At the record level: only half of the indexed records have anzsrc-for codes

Machine learning for record classification

- Assign anzsrc-for code to unlabelled records automatically
 - Aim to improve search experience for both human and machine
 - Understand domain coverage of the collection
- Train models, three components are essential for the training:
 - Labels - anzsrc-for code
 - Classifier - four supervised machine learning methods:
 - multinomial logistic regression (MLR), multinomial naive bayes (MNB), K Nearest Neighbors (KNN), Support Vector Machine (SVM)
 - Data - (~78k) records with anzsrc-for code
 - Split into two sets: training set, test set
- Apply model(s)/best prediction to unlabelled records

Record classification with ansrc-for code

- Use 77918 records that have an ansrc-for code for training models
- Step by step: first the top two digits, then move down to four, six digits
- Four models: multinomial logistic regression (MLR), multinomial naive bayes (MNB), K Nearest Neighbors (KNN), Support Vector Machine (SVM)

Model	Training Set Accuracy	Test Set Accuracy
Logistic Regression	0.769149	0.701299
SVM	0.696435	0.676324
Multinomial Naïve Bayes	0.702965	0.659341
KNN	0.906460	0.642358

Acknowledgement:

Adapted [the code](#) from Miguel Frenandez Zafra

Performance per category

Most correlated unigrams:

Code	Top 5	Bottom 5
04	earth airborne geophysical mount ignsn	al unit two australia region
15	study financial survey university dataset	given number received document expert

04: Earth Science

15: Commerce, Management, Tourism and Services

2 digitis Code	MLR	SVM	MNB	KNN	No. of records
01	0.29	0.00	0.43	0.33	111
02	0.97	1.00	0.95	1.00	*300
03	0.67	0.56	0.55	0.58	499
04	0.98	0.96	0.94	0.96	*600
05	0.68	0.71	0.53	0.54	*400
06	0.98	1.00	0.89	0.78	*600
07	0.63	0.55	0.52	0.79	*200
08	0.42	0.22	0.23	0.41	386
09	0.95	1.00	1.00	0.84	*200
10	0.33	0.00	0.00	0.19	128
11	0.82	0.81	0.83	0.66	*400
12	0.71	1.00	0.77	0.81	174
13	0.58	0.82	0.54	0.56	148
14	0.35	0.00	0.83	0.57	122
15	0.23	0.00	0.00	0.25	76
16	0.49	0.47	0.44	0.50	*300
17	0.47	0.00	0.57	0.50	112
18	1.00	1.00	1.00	1.00	*400
19	0.77	0.62	0.48	0.58	343
20	0.64	0.72	0.48	0.20	*300
21	0.96	0.94	0.88	0.98	*600
22	0.22	0.00	0.33	0.20	79
micro ave	0.70	0.68	0.66	0.64	
macro ave	0.65	0.56	0.60	0.61	

Examples of classification within two-digits code

Method: MLR

06: Biological Sciences (41505 records)

02: Physical Sciences (3533 records)

06: 17268 records (out of 41505) have both 0601 and 0604 labels

	precision	test data		precision	test data
0601	0.58	2859	0201	1.00	752
0602	0.99	652	0202	0.00	1
0603	0.15	22	0203	0.04	2
0604	0.49	2560	0204	0.32	13
0605	0.01	11	0205	0.00	0
0606	1	3	0206	0.00	0
0607	0.1	48	0299	1.00	116
0608	0.52	51			
0699		20			
micro avg	0.5	6226		0.58	884
macro ave	0.43			0.34	
weighted ave	0.58			0.99	

Confusion matrix - MLR (06 - Biological Sciences)

	601	602	603	604	605	606	607	608	699
Biochemistry and Cell Biology (601) -	560	0	6	2248	0	1	0	44	0
Ecology (602) -	4	325	21	0	0	57	0	230	15
Evolutionary Biology (603) -	2	0	6	0	0	3	0	11	0
Genetics (604) -	397	1	1	2130	0	15	0	16	0
Microbiology (605) -	1	1	2	0	0	4	0	3	0
Physiology (606) -	0	0	0	0	0	1	0	2	0
Plant Biology (607) -	3	0	2	0	0	15	10	18	0
Zoology (608) -	2	1	3	0	0	7	0	38	0
Other Biological Sciences (699) -	1	0	0	0	0	1	0	2	16
	601	602	603	604	605	606	607	608	699

Predicted

Actual

Discussion and future work

- User behaviour:
 - Evidence that subject headings are used
 - Why and why not
 - Low usage of subject headings from this log collection
 - Is this unique to this data catalogue and interface?

Log analysis + survey and interview
- Collection characteristics:
 - Large proportion of records from the catalogue without a “standard” vocabulary for the subject headings **a known issue**
 - Those with subject headings are biased toward a few categories
 - Encourage underrepresented subject areas to publish and share data
 - Record classification works for some categories
 - Explore correlation, improvement

Thanks!