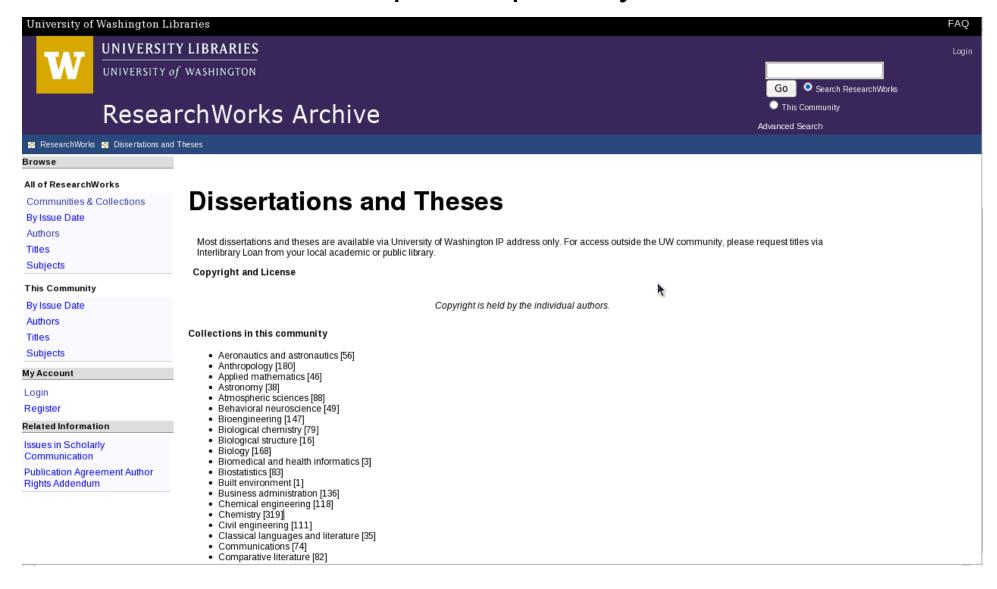


Jeff Sherwood, Programmer. Anjanette Young, Systems Librarian. University of Washington, Libraries.

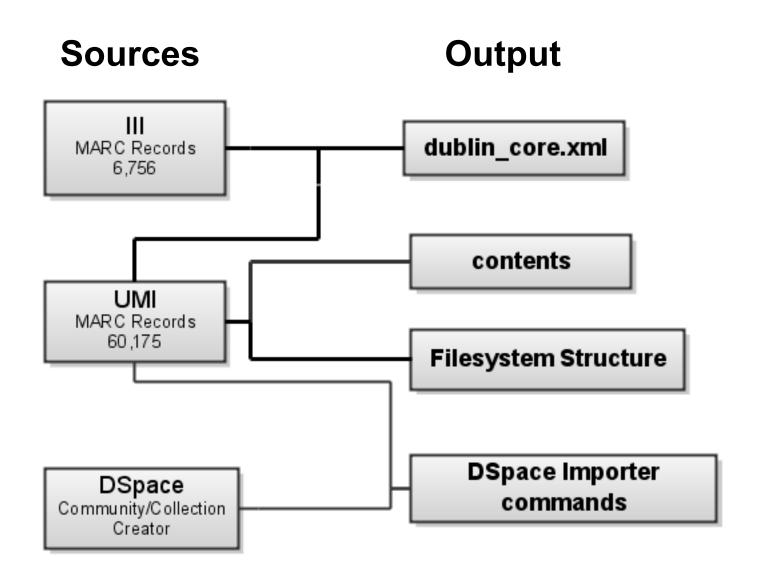
DSpace Repository

Goal

Ingest Metadata and PDF's for ETD's received from UMI into a DSpace repository.



Electronic Theses & Dissertations



MARC Fields

UMI Records

=001 (Filename) =520 (Abstract)

III Records

=001 (OCLC number)

=100 (Author)

=245 (Title)

=260 (Date published)

=502 (type and date)

=695 (Department)

=941 (Local identifier)

dublin_core.xml

```
<dublin core>
  <dcvalue element="identifier" qualifier="other">
     iii[941]</dcvalue>
  <dcvalue element="title" qualifier="none">
     iii[245][a][b]</dcvalue>
  <dcvalue element="contributor" qualifier="author">
     iii[100][a][b][c]</dcvalue>
  <dcvalue element="description" qualifier="abstract">
     umi[520][a]</dcvalue>
  <dcvalue element="subject" qualifier="other">
     iii[655][a][x]</dcvalue>
</dublin core>
```

MARC Loader . . . No.

```
|||0|0| | |0|n|G|0|@ov action="o"
|||0|0| | |0|n|G|0
@ov protect="b=V0123456789d(690,695:d)
                    hn(590:d)y(099,249,852,856:d)y(910,925,
                    980,981)F26"
035|001 |+|0|0|b|o|0|y|N|0|%001(start="1-9",char="!-~")
245||+|0|0|b|t|0|y|N|0|%bracket="h"
500-599||+|0|0|b|n|0|y|N|0|
600-651||-w|0|0|b|d|0|y|N|0|
653-657||+|0|0|b|d|0|y|N|0|
690-699||-w|0|0|b|d|0|y|N|0|
700-715||-w|0|0|b|b|0|y|N|0|
730-740||-w|0|0|b|f|0|y|N|0|
```

Matching overview

Ham-fisted Method

- 1. Exact Title + Exact Author
- 2. Exact Title + Shortened Author

Cool Math Method

Calculate Similarity of Title Calculate Similarity of Author

- 1. Exact Title + Fuzzy Author
- 2. Fuzzy Title + Fuzzy Author
- 3. Fuzzy Title or Fuzzy Author

Pymarc - the MARC Hammer

```
umi dict = {
 Alaskan Bootlegger: {author: Leon Kania, umi count = 1},
 title2 value: {author: author2 value, umi count = index2},
iii dict = {
 Alaskan Bootlegger: {author: Leon W. Kania, iii count = 9},
 title2_value: {author: author2 value, iii count = index2},
```

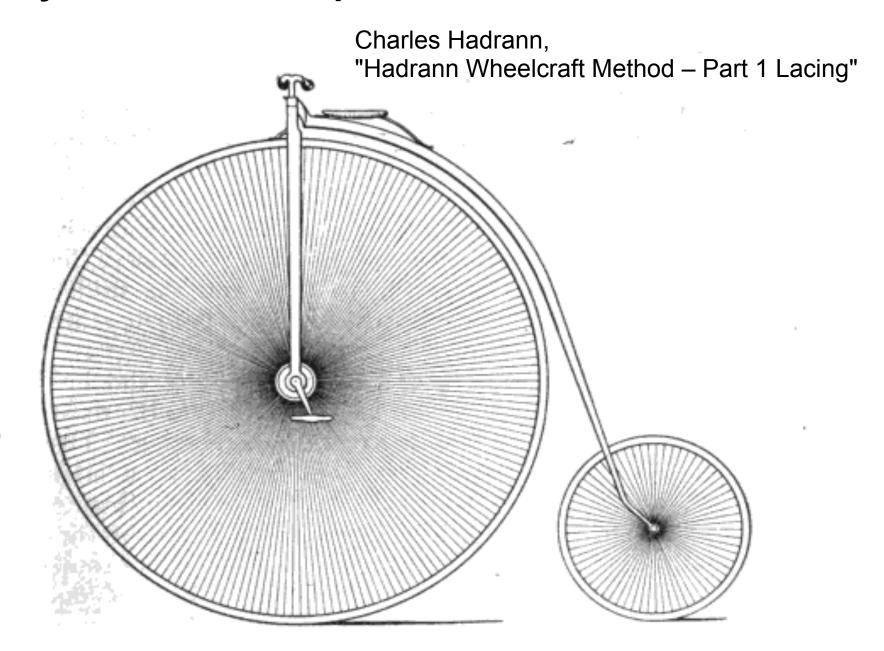
Exact title + exact author

```
# Exact Title
# Create sets out of the dictionary keys
umi set = set(umi dict.iterkeys())
iii set = set(iii dict.iterkeys())
# Find the Intersection of sets.
title match = umi set & iii set
# Verify Intersection with Exact Author
for x in title match:
  if umi_dict[x][author] == iii dict[x][author]:
     ... do stuff.
```

Exact title + Truncated author

```
def shortenAuthorName(name):
    #Leon W. Kania   -> [Leon, W., Kania]
    namelist = str(name).split()
    if len(namelist) > 2:
        shortname = "%s %s" % (namelist[0], namelist[-1])
    else:
        shortname = name
    return shortname
```

"If you break three spokes, it is time for a rebuild"



Rogues Gallery

USE OF CROWN LENGTH TO DEFINE STEM FORM: SEGMENTED TAPER EQUATION (DOUGLAS FIR)

Use of crown length to define stem form :: segmented taper equation Towards an understanding of seismic performance of three-dimensional structures: Stability and reliability

Towards an understanding of seismic performance of 3D structures :: stability & reliability

Hoekstra, Hopi Danielle Elisabeth

Hoekstra, Danielle E

Arnason, Halldor

Halldór Árnason



Levenshtein Edit Distance

Edit distance is the number of operations required to transform one string of characters into the another.

How many steps to turn kitten into sitting?

kitten → **sitten** (k changes to s)

sitten → sittin (e changes to i)

sittin → sitting (insert g)

LD is Always...

- ≥ difference in string lengths
- ≤ length of the longer string
- = 0 if the strings are identical

Similarity Score

$$similarity = 1 - \frac{d_L}{\max(|s_1|, |s_2|)}$$

Optimizations

"A stochastic model of cyclical interaction processes"

All titles

Identify Stopwords

the: 24587

for: 7643

with: 3323

effects: 1958

evaluation: 1073

. . .

hypoxic: 1

reduplication: 1

picaresque: 1

emperador 1

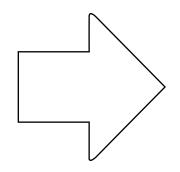
heteroduplex 1

Throw out common words in titles

Keep the rarer ones

Extract Significant Words

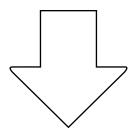
"Stochastic models for DNA sequence data"



stochastic dna sequence

```
rec = {'title': 'Stochastic models...',}
index['stochastic'].append(rec)
index['dna'].append(rec)
index['sequence'].append(rec)
```

index['stochastic']



```
{'title': "Stochastic models for DNA sequence data", ...}
{'title': "A stochastic model of clan systems", ...}
{'title': "A stochastic model of cyclical interaction processes", ...}
{'title': "Stochastic reliability models for maintained systems", ...}
{'title': "Uniform approximation and almost periodicity of doubly stochastic operators", ...}
```

Hoekstra, Hopi Danielle Elisabeth

Hoekstra, Danielle E



Hoekstra, H

Hoekstra, D

Arnason, Halldor

Halldór Árnason



Arnason, H

Árnason, H

mprovements

Jaro-Winkler Algorithm

$$d_j = \frac{1}{3} \left(\frac{m}{|s_1|} + \frac{m}{|s_2|} + \frac{m-t}{m} \right)$$

What's a "match"?

Two characters match if they are a reasonable distance from one another as defined by:

$$\left| \frac{max(|s_1|, |s_2|)}{2} \right| - 1$$

Example

$$dj = \frac{1}{3} \left(\frac{m}{|s_1|} + \frac{m}{|s_2|} + \frac{m-t}{m} \right)$$

s1 = Martha

s2 = Marhta

$$dj = \frac{1}{3} \left(\frac{6}{6} + \frac{6}{6} + \frac{6-1}{6} \right)$$

$$\approx 0.944$$

Example

$$similarity = 1 - \frac{d_L}{\max(|s_1|, |s_2|)}$$

s1 = Martha

s2 = Marhta

$$similarity = 1 - \frac{2}{6}$$

 ≈ 0.667

Jaro-Winkler works best for short strings

Resources

Levenshtein & Jaro-Winkler

Background

http://en.wikipedia.org/wiki/Levenshtein_distance http://en.wikipedia.org/wiki/Jaro-Winkler distance

Code

http://pypi.python.org/pypi/editdist/0.1 http://pypi.python.org/pypi/python-Levenshtein/0.10.1

Miscellaneous

String Comparison Tutorial

http://bit.ly/ZGSmF

SecondString - Java text analysis library

http://secondstring.sourceforge.net/

MarcXimiL - MARC de-duping package

http://marcximil.sourceforge.net/

http://snurl.com/uggtn