

Data Cleaning for Analysis and Publication Using the OpenRefine Software Package

Arctic Data Center, CUAHSI, DataONE, Environmental Data Initiative, GBIF, **iDigBio**, NEON, Neotoma

Jeanette Clark, **Deborah Paul**

#datahelpdesk

Ecological Society of America 2020

Career Central

August

<http://bit.ly/datahelpesa2020>





Data Archive



Portal for data discovery



Tools for data and metadata submission



Support services



Training and Outreach



Data Rescue



5,800
DATA SETS



1,500
CREATORS



725K
DATA FILES



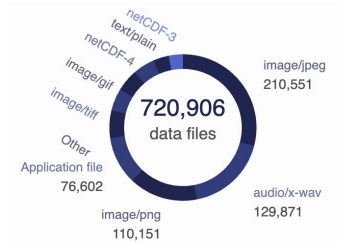
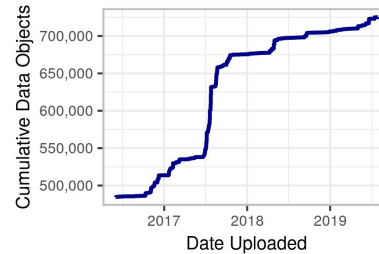
14,000
USERS



34 TB
DATA STORAGE



990K+
FILE DOWNLOADS



 @arcticdatactr



Advancing the Digitization of Biological Collections

iDigBio Hub and Thematic (Museum) Collection Networks

total: 121,428,342

Digitization

Workflows & Protocols
Dissemination

Research Use

Cyberinfrastructure
Tool collaboration
Portal development
ENM workshop

Research focus

Data quality
APIs

Training

Biodiversity informatics
Data skills and literacy
Collections software
Imaging
Project Management

www.idigbio.org

Search Records

search all fields

Must have media Must have map point

Filters Mapping Sorting Download

Add a field Clear

State/Province: kentucky

Present Missing

Top 6 Taxa

- Crinoidea
- Empoasca empoasca
- Agallia constricta
- Paraphlepsius irroratus
- Fungipata
- other

(Meta)data Aggregator

Community Building

Collections Data

PEOPLE IN THE LOOP

people graphic by Dorothy Allard

Education Outreach

Citizen Science
K-12 materials
Undergraduate
Fossil Clubs
Mentor teachers

Methods

Workshops
Webinars
Symposia
Conferences
Working Groups
Short Courses
Adobe Connect
Listservs
Publications
Social Media @idigbio



What do we mean by “Clean” Data?

More Data from More Sources =

- Structural Issues
- Inconsistent/unclear missing values
- Mixed data in single columns
- Mixed data types in single columns
- Ambiguous data values
- Data you can't use

JESLIB 2013; 2(2): 3-16
doi:10.7191/jeslib.2013.1024



Journal of eScience Librarianship

putting the pieces together: theory and practice

Common Errors in Ecological Data Sharing

Karina E. Kervin,¹ William K. Michener,² Robert B. Cook³

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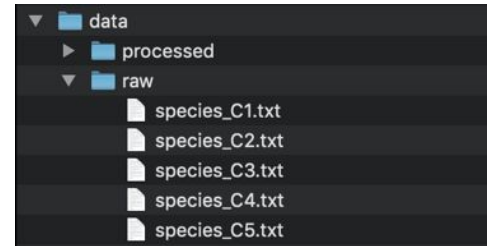
³ Oak Ridge National Laboratory, Oak Ridge, TN, USA

Simple guidelines for data management

- Use a scripted program
- Nonproprietary formats
- Keep a raw version of data
- Descriptive names
- Header line
- Plain ASCII text



.csv, .txt



```
1 Date,Time,Station,Latitude,Longitude,Target_Depth,CTD_Depth,CTD_Salinity,CTD_Temperature
2 3/21/08,1899-12-31 21:56:46,"73N,140W",73.02083333,-139.8885,20,15.127,26.0658,-1.423
3 3/21/08,1899-12-31 21:56:46,"73N,140W",73.02083333,-139.8885,60,60.5559,29.1798,-0.93431
4 3/21/08,1899-12-31 21:56:46,"73N,140W",73.02083333,-139.8885,85,85.7471,31.4023,-0.14583
5 3/21/08,1899-12-31 21:56:46,"73N,140W",73.02083333,-139.8885,190,191.4073,33.1268,-1.4775
6 3/21/08,1899-12-31 21:56:46,"73N,140W",73.02083333,-139.8885,310,309.2524,34.6233,0.25782
7 3/22/08,1899-12-31 21:45:27,"72N,140W",72.0505,-140.1118333,20,20.9588,26.1788,-1.4007
```

Borer, E. T. et al, (2009), Some Simple Guidelines for Effective Data Management. The Bulletin of the Ecological Society of America

Simple Guidelines for Data Management

- Design to add rows, not columns
- Each column should contain only one type of information
- Record a single piece of data only once; separate information collected at different scales into different tables. In other words, create a relational database.

Recognizing untidy data

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q
species	tree	main trunks kg	reiterated trunks kg	limbs kg	branches kg	leaves kg		type	species	main trunk	reiteration	dry masses (kg)			TOTAL	% total
												limb	branch	leaf		
SESE	Atlas	255144.9	46020.6	5477.7	13433.2	1101.2		tree	SESE	3569312	213247	53714	230945	17192	4084409	95.3491
SESE	Ballantine	221966.4	7651.6	5922.9	11210.0	1084.8		tree	PSME	135815	0	0	8338	961	145114	3.3876
SESE	Bell	253246.4	5454.3	5792.6	48500.7	1043.4		tree	THSE	31799	0	0	6343	864	39006	0.9105
SESE	Broken Top	130928.9	4805.2	1608.1	5137.4	729.9		tree	ACMA	4444	0	0	925	264	5634	0.1315
SESE	Buena Vista	128833.0	3486.5	0.0	8552.1	518.4		tree	UMCA	2921	0	0	937	273	4131	0.0964
SESE	Demeter	155896.0	11085.6	3204.3	10054.1	768.7		shrub	RUSP	0	0	0	1974	686	2660	0.0620
SESE	Epimetheus	226987.0	12915.7	1797.2	13585.2	1029.4		fern	POMU	0	0	0	0	1271	1271	0.0296
SESE	Iluvatar	349586.6	65003.9	12315.6	13987.0	1461.8		shrub	VAOV	0	0	0	526	26	552	0.0129
SESE	Kronos	134154.1	12204.4	7232.7	5036.1	597.3		shrub	COCO	0	0	0	284	6	289	0.0067
SESE	Pleiades I	182385.2	3735.0	1935.2	10846.6	762.2		fern	POSC	0	0	0	107	89	196	0.0045
SESE	Pleiades II	235838.8	11183.4	4306.0	11306.5	877.7		tree	RHPU	100	0	0	44	18	162	0.0037
SESE	Prometheus	239414.0	25228.9	1612.6	12458.2	1086.0		herb	OXOR	0	0	0	0	112	112	0.0026
SESE	Rhea	143710.4	487.8	730.1	5524.2	691.2		shrub	VAPA	0	0	0	94	4	99	0.0023
SESE	Zeus	243365.7	2885.5	1620.4	19104.7	954.3		tree	PISI	0	0	0	1	0	1	0.0000
SESE	3	1761.3	0.0	0.0	87.6	41.4		tree	CHLA	0	0	0	1	0	1	0.0000
SESE	4	6312.0	356.0	73.5	214.1	43.8		shrub	GASH	0	0	0	0	0	0	0.0000
SESE	5	206.0	0.0	0.0	8.7	2.5		shrub	SACA	0	0	0	0	0	0	0.0000
SESE	6E	18697.4	0.0	0.0	1055.2	66.3				3744390	213247	53714	250519	21767	4283636	
SESE	6W	14651.5	7.7	0.0	626.3	49.6										proportion
SESE	11	614.4	0.0	0.0	28.1	17.0				main trunk	reiteration	limb	branch	leaf	total	geophytic
SESE	12	232.1	0.0	0.0	11.2	10.3		SESE	geo	3569312	213247	53714	230945	17192	4084409	1.00
SESE	18	15632.0	0.0	0.0	946.3	106.8		SESE	epi	0	0	0	0	0	0	
SESE	19	11805.5	0.0	0.0	770.1	80.3		PSME	geo	135815	0	0	8338	961	145114	1.00
SESE	20	309.5	0.0	0.0	12.5	5.9		PSME	epi	0	0	0	0	0	0	
SESE	22	25618.3	0.0	0.0	1504.0	120.2		TSHE	geo	31740	0	0	6332	860	38932	0.99
SESE	23	482.7	0.0	0.0	18.8	4.6		TSHE	epi	0	0	0	13	4	74	

Characteristics of Tidy Data

Observations

- **Separate tables for each entity measured**

Recognizing untidy data

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q
species	tree	main trunks kg	reiterated trunks kg	limbs kg	branches kg	leaves kg		type	species	main trunk	reiteration	limb	branch	leaf	TOTAL	% total
SESE	Atlas	255144.9	46020.6	5477.7	13433.2	1101.2		tree	SESE	3569312	213247	53714	230945	17192	4084409	95.3491
SESE	Ballantine	221966.4	7651.6	5922.9	11210.0	1084.8		tree	PSME	135815	0	0	8338	961	145114	3.3876
SESE	Bell	253246.4	5454.3	5792.6	48500.7	1043.4		tree	THSE	31799	0	0	6343	864	39006	0.9105
SESE	Broken Top	130928.9	4805.2	1608.1	5137.4	729.9		tree	ACMA	4444	0	0	925	264	5634	0.1315
SESE	Buena Vista	128833.0	3486.5	0.0	8552.1	518.4		tree	UMCA	2921	0	0	937	273	4131	0.0964
SESE	Demeter	155896.0	11085.6	3204.3	10054.1	768.7		shrub	RUSP	0	0	0	1974	686	2660	0.0620
SESE	Epimetheus	226987.0	12915.7	1797.2	13585.2	1029.4		fem	POMU	0	0	0	0	1271	1271	0.0296
SESE	Iluvatar	349586.6	65003.9	12315.6	13987.0	1461.8		shrub	VAOV	0	0	0	0	26	552	0.0129
SESE	Kronos	134154.1	12204.4	7232.7	5036.1	597.3		shrub	COCO	0	0	0	0	6	289	0.0067
SESE	Pleiades I	182385.2	3735.0	1935.2	10846.6	762.2		fem	POSC	0	0	0	107	89	196	0.0045
SESE	Pleiades II	235838.8	11183.4	4306.0	11306.5	877.7		tree	RHPU	100	0	0	44	18	162	0.0037
SESE	Prometheus	239414.0	25228.9	1612.6	12458.2	1086.0		herb	OXOR	0	0	0	0	112	112	0.0026
SESE	Rhea	243997.4	487.7	730.1	5524.2	691.2		shrub	VAPA	0	0	0	94	4	99	0.0023
SESE	Zeus	243997.4	3885.5	1620.4	19104.7	954.3		tree	PISI	0	0	0	1	0	1	0.0000
SESE	3	1761.3	0.0	0.0	87.6	41.4		tree	CHLA	0	0	0	1	0	1	0.0000
SESE	4	6312.0	356.0	73.5	214.1	43.8		shrub	GASH	0	0	0	0	0	0	0.0000
SESE	5	206.0	0.0	0.0	8.7	2.5		shrub	SACA	0	0	0	0	0	0	0.0000
SESE	6E	18697.4	0.0	0.0	1055.2	66.3										
SESE	6W	14651.5	7.7	0.0	626.3	49.8										
SESE	11	614.4	0.0	0.0	28.1	17.0										
SESE	12	232.1	0.0	0.0	11.2	10.3										
SESE	18	15632.0	0.0	0.0	946.3	106.8										
SESE	19	11805.5	0.0	0.0	770.1	80.3										
SESE	20	309.5	0.0	0.0	12.5	5.9										
SESE	22	25618.3	0.0	0.0	1504.0	120.2										
SESE	23	483.7	0.0	0.0	18.8	4.6										
										3744390	213247	53714	250519	21767	4283636	
																proportion
										main trunk	reiteration	limb	branch	leaf	total	geophytic
									SESE geo	3569312	213247	53714	230945	17192	4084409	1.00
									SESE epi	0	0	0	0	0	0	
									PSME geo	135815	0	0	8338	961	145114	1.00
									PSME epi	0	0	0	0	0	0	
									TSHE geo	31740	0	0	632	860	38932	0.99
									TSHE epi	0	0	0	0	0	0	

Table 1

Table 2

Table 3

Characteristics of Tidy Data

Observations

- Separate tables for each entity measured
- **Each row represents a single observed entity**

Recognizing untidy data

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q
species	tree	main trunks kg	reiterated trunks kg	limbs kg	branches kg	leaves kg		type	species	main trunk	reiteration	limb	branch	leaf	TOTAL	% total
SESE	Atlas	255144.9	46020.6	5477.7	13433.2	1101.2		tree	SESE	3569312	213247	53714	230945	17192	4084409	95.3491
SESE	Ballantine	221966.4	7651.6	5922.9	11210.0	1084.8		tree	PSME	135815	0	0	8338	961	145114	3.3876
SESE	Bell	253246.4	5454.3	5792.6	48500.7	1043.4		tree	THSE	31799	0	0	6343	864	39006	0.9105
SESE	Broken Top	130928.9	4805.2	1608.1	5137.4	729.9		tree	ACMA	4444	0	0	925	264	5634	0.1315
SESE	Buena Vista	128833.0	3486.5	0.0	8552.1	518.4		tree	UMCA	2921	0	0	937	273	4131	0.0964
SESE	Demeter	155896.0	11000.0	3204.3	10054.1	768.7		shrub	RUSP	0	0	0	1974	686	2660	0.0620
SESE	Epimetheus	226987.0	12915.7	1797.2	13585.2						0	0	0	1271	1271	0.0296
SESE	Iluvatar	349586.6	65003.9	10015.6	13987.0						0	0	526	26	552	0.0129
SESE	Kronos	134154.1	12204.4	7200.7	5036.1						0	0	284	6	289	0.0067
SESE	Pleiades I	182385.2	3735.0	1935.2	10846.6						0	0	107	89	196	0.0045
SESE	Pleiades II	235838.8	11183.4	4306.0	14306.5						0	0	44	18	162	0.0037
SESE	Prometheus	239414.0	25228.9	1612.6	12700.2						0	0	0	112	112	0.0026
SESE	Rhea	143710.4	487.8	730.1	5524.2						0	0	94	4	99	0.0023
SESE	Zeus	243365.7	2885.5	1620.4	19104.7						0	0	1	0	1	0.0000
SESE	3	1761.3	0.0	0.0	87.6						0	0	1	0	1	0.0000
SESE	4	6312.0	356.0	73.5	214.1						0	0	0	0	0	0.0000
SESE	5	206.0	0.0	0.0	8.7						0	0	0	0	0	0.0000
SESE	6E	18697.4	0.0	0.0	1055.2						247	53714	250519	21767	4283636	
SESE	6W	14651.5	7.7	0.0	626.3											proportion
SESE	11	614.4	0.0	0.0	28.1	17.0				main trunk	reiteration	limb	branch	leaf	total	geophytic
SESE	12	232.1	0.0	0.0	11.2	10.3		SESE geo		3569312	213247	53714	230945	17192	4084409	1.00
SESE	18	15632.0	0.0	0.0	946.3	106.8		SESE epi		0	0	0	0	0	0	
SESE	19	11805.5	0.0	0.0	770.1	80.3		PSME geo		135815	0	0	8338	961	145114	1.00
SESE	20	309.5	0.0	0.0	12.5	5.9		PSME epi		0	0	0	0	0	0	
SESE	22	25618.3	0.0	0.0	1504.0	120.2		TSHE geo		31740	0	0	6332	860	38932	0.99
SESE	23	463.7	0.0	0.0	18.0	4.6		TSHE epi		0	0	0	0	0	0	

All the same observation?
No.

Characteristics of Tidy Data

Observations

- Separate tables for each entity measured
- Each row represents a single observed entity

Variables

- **All values in a column are of the same type**

Recognizing untidy data

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q
species	tree	main trunks kg	reiterated trunks kg	limbs kg	branches kg	leaves kg		type	species	main trunk	reiteration	limb	branch	leaf	TOTAL	% total
SESE	Atlas	255144.9	46020.6	5477.7	13433.2	1101.2		tree	SESE	3569312	213247	53714	230945	17192	4084409	95.3491
SESE	Ballantine	221966.4	7651.6	5922.9	11210.0	1084.8		tree	PSME	135815	0	0	8338	961	145114	3.3876
SESE	Bell	253246.4	5454.3	5792.6	48500.7	1043.4		tree	THSE	31799	0	0	6343	864	39006	0.9105
SESE	Broken Top	130928.9	4805.2	1608.1	5137.4	729.9		tree	ACMA	4444	0	0	925	264	5634	0.1315
SESE	Buena Vista	128833.0	3486.5	0.0	8552.1	518.4		shrub	UMCA	2921	0	0	937	273	4131	0.0964
SESE	Demeter	155896.0	11085.6	3204.3	10054.1	768.7		shrub	RUSP	0	0	0	1974	686	2660	0.0620
SESE	Epimetheus	226987.0	12915.7	1797.2	13585.2	1029.4		fem	POMU	0	0	0	0	1271	1271	0.0296
SESE	Iluvatar	349586.6	65003.9	12315.6	13987.0	1481.8		shrub	VAOV	0	0	0	526	26	552	0.0129
SESE	Kronos	134154.1	12204.4	7232.7	5036.0	503.6				0	0	0	284	6	289	0.0067
SESE	Pleiades I	182385.2	3735.0	1935.2	10846.0	1084.6				0	0	0	107	89	196	0.0045
SESE	Pleiades II	235838.8	11183.4	4306.0	11306.0	1130.6				0	0	0	44	18	162	0.0037
SESE	Prometheus	239414.0	25228.9	1612.6	12456.0	1245.6				0	0	0	0	112	112	0.0026
SESE	Rhea	143710.4	487.8	730.1	5524.0	552.4				0	0	0	94	4	99	0.0023
SESE	Zeus	243365.7	2885.5	1620.4	19104.0	1910.4				0	0	0	1	0	1	0.0000
SESE	3	1761.3	0.0	0.0	87.0	8.7				0	0	0	1	0	1	0.0000
SESE	4	6312.0	356.0	73.5	214.0	21.4				0	0	0	0	0	0	0.0000
SESE	5	206.0	0.0	0.0	6.0	0.6				0	0	0	0	0	0	0.0000
SESE	6E	18697.4	0.0	0.0	1055.0	105.5				213247	53714	250519	21767	4283636		
SESE	6W	14651.5	7.7	0.0	626.0	62.6										
SESE	11	614.4	0.0	0.0	24.0	2.4										
SESE	12	232.1	0.0	0.0	11.2	1.1		SESE geo		3569312	213247	53714	230945	17192	4084409	1.00
SESE	18	15632.0	0.0	0.0	946.3	94.6		SESE epi		0	0	0	0	0	0	
SESE	19	11805.5	0.0	0.0	770.1	77.0		PSME geo		135815	0	0	8338	961	145114	1.00
SESE	20	309.5	0.0	0.0	12.5	1.2		PSME epi		0	0	0	0	0	0	
SESE	22	25618.3	0.0	0.0	1504.0	150.4		TSHE geo		31740	0	0	6332	860	38932	0.99
SESE	23	463.7	0.0	0.0	18.0	1.8		TSHE epi		0	0	0	0	0	0	

All the same variable?
No.



Characteristics of Tidy Data

Observations

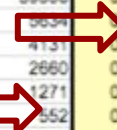
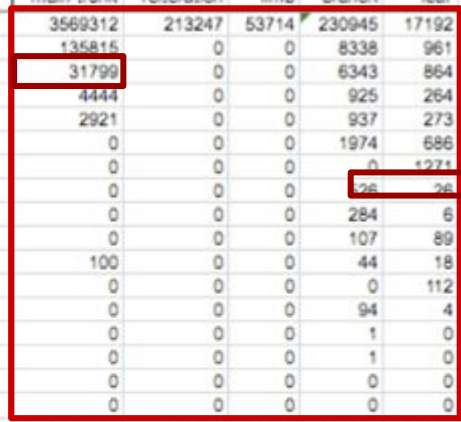
- Separate tables for each entity measured
- Each row represents a single observed entity
- **Observations (rows) are all unique**

Variables

- All values in a column are of the same type
- **All columns pertain to the same observation (row)**
- **Each column represents either an identifying or measured variable**

A not-so-reproducible workflow

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q
species	tree	main trunks kg	reiterated trunks kg	limbs kg	branches kg	leaves kg		type	species	main trunk	reiteration	dry masses (kg)		leaf	TOTAL	% total
												limb	branch			
SESE	Atlas	255144.9	46020.6	5477.7	13433.2	1101.2		tree	SESE	3569312	213247	53714	230945	17192	4084409	95.3491
SESE	Baillantine	221966.4	7651.6	5922.9	11210.0	1084.8		tree	PSME	135815	0	0	8338	961	145114	3.3876
SESE	Bell	253246.4	5454.3	5792.6	48500.7	1043.4		tree	THSE	31799	0	0	6343	864	39006	0.9105
SESE	Broken Top	130958.9	4805.2	1608.1	5137.4	729.9		tree	ACMA	4444	0	0	925	264	5633	0.1315
SESE	Buena Vista	35833.0	3486.5	0.0	8552.1	518.4		tree	UMCA	2921	0	0	937	273	4131	0.0964
SESE	Demeter	55896.0	11085.6	3204.3	10054.1	768.7		shrub	RUSP	0	0	0	1974	686	2660	0.0620
SESE	Epimetheus	226987.0	12915.7	1797.2	13585.2	1029.4		ferm	POMU	0	0	0	0	1271	1271	0.0296
SESE	Iliuvatar	349586.6	7003.9	12315.6	13987.0	1481.8		shrub	VAOV	0	0	0	28	28	52	0.0129
SESE	Kronos	134154.1	12204.4	7232.7	5036.1	597.3		shrub	COCO	0	0	0	284	6	289	0.0067
SESE	Pleiades I	182385.2	1735.0	1935.2	10846.6	762.2		ferm	POSC	0	0	0	107	89	196	0.0045
SESE	Pleiades II	235838.8	1183.4	4306.0	11306.5	877.7		tree	RHPU	100	0	0	44	18	162	0.0037
SESE	Prometheus	239414.0	25228.9	1612.6	12458.2	1086.0		herb	OXOR	0	0	0	0	112	112	0.0026
SESE	Rhea	13710.4	487.8	730.1	5524.2	691.2		shrub	VAPA	0	0	0	94	4	99	0.0023
SESE	Zeus	2005.7	2885.5	1620.4	19104.7	954.3		tree	PISI	0	0	0	1	0	1	0.0000
SESE	3	1761.3	0.0	0.0	87.6	41.4		tree	CHLA	0	0	0	1	0	1	0.0000
SESE	4	6312.0	356.0	73.5	214.1	43.8		shrub	GASH	0	0	0	0	0	0	0.0000
SESE	5	206.0	0.0	0.0	8.7	2.5		shrub	SACA	0	0	0	0	0	0	0.0000
SESE	6E	18697.4	0.0	0.0	1055.2	66.3				3744390	213247	53714	250519	21767	4283636	
SESE	6W	14651.5	7.7	0.0	626.3	49.6										
SESE	11	614.4	0.0	0.0	28.1	17.0										
SESE	12	232.1	0.0	0.0	11.2	10.3		SESE geo		3569312	213247	53714	230945	17192	4084409	1.00
SESE	18	15632.0	0.0	0.0	946.3	106.8		SESE epi		0	0	0	0	0	0	0
SESE	19	11805.5	0.0	0.0	770.1	80.3		PSME geo		135815	0	0	8338	961	145114	1.00
SESE	20	309.5	0.0	0.0	12.5	5.9		PSME epi		0	0	0	0	0	0	0
SESE	22	25618.3	0.0	0.0	1504.0	120.2		THSE geo		31740	0	0	6332	860	38932	0.99
SESE	23	463.7	0.0	0.0	18.0	4.6		THSE epi		0	0	0	0	0	0	0



Building a reproducible workflow

id	date	site	spcode	height
1	2017-10-10	1	DAPU	4.6
2	2017-09-05	2	DAMA	3.5
3	2017-10-10	1	DAMA	4.5
4	2017-09-05	2	DAPU	3.9

Raw, messy data



id	date	site	spcode	height
1	2017-10-10	1	DAPU	4.6
2	2017-09-05	2	DAMA	3.5
3	2017-10-10	1	DAMA	4.5
4	2017-09-05	2	DAPU	3.9

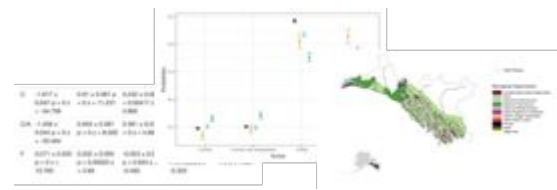
site	name	elev
1	Taku	3.7
2	Lituya	3.2

Clean, raw data



id	date	site	spcode	height	name	elev
1	2017-10-10	1	DAPU	4.6	Taku	3.7
2	2017-09-05	2	DAMA	3.5	Lituya	3.2
3	2017-10-10	1	DAMA	4.5	Taku	3.7
4	2017-09-05	2	DAPU	3.9	Lituya	3.2

Merged/summarized derived data



Figures, tables, maps

Where data come from matters! (a sample)

- **Excel** Issues that may arise from these data sources are our focus today
 - Automatic conversion of gene names to dates or floating point numbers*
 - Date values can be converted when transferring data between operating systems and applications
- **Text (e.g. CSV) & Excel**
 - Free-form structure - lack of enforcement of column-row structure, type consistency
- **Text (e.g. CSV)**
 - Inconsistent structure - quotes, commas, missing values, spaces
- **Database**
 - Enforced structure - tables, column typing
 - Specialized methods for interaction (pros and cons to this)

The ESAUSSEE Data Help Desk

who we are and how to find us

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Messy data?

Repetitive data tasks?

Increase Reproducibility and Productivity using tools like [Open Refine](#)



OpenRefine

A power tool for working with messy data.



Magic is here.
Ask for it, plan for it.



GBIF



iDigBio
Integrated Digitized Biocollections



neon
Operated by Battelle



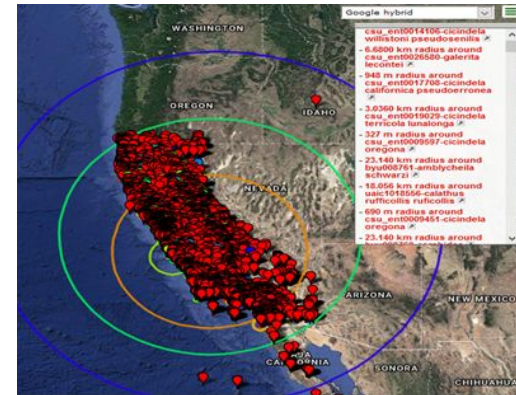
NEOTOMA PALEOECOLOGY DATABASE



Data lessons compiled - inspired by workshop

Georeferencing for Research Use of Museum Collections Data

- **Data mapped to standards**
 - supports use and re-use (e.g. Darwin Core DwC, Ecological Metadata Language EML)
 - standards help with data validation and cleaning
- **Data have issues**
 - what are some you have experienced
 - need to be addressed before applying research methods
 - keep raw data raw
 - track your changes
- **Data visualization is key**
 - QGIS lessons
 - Open Refine
 - R, etc.

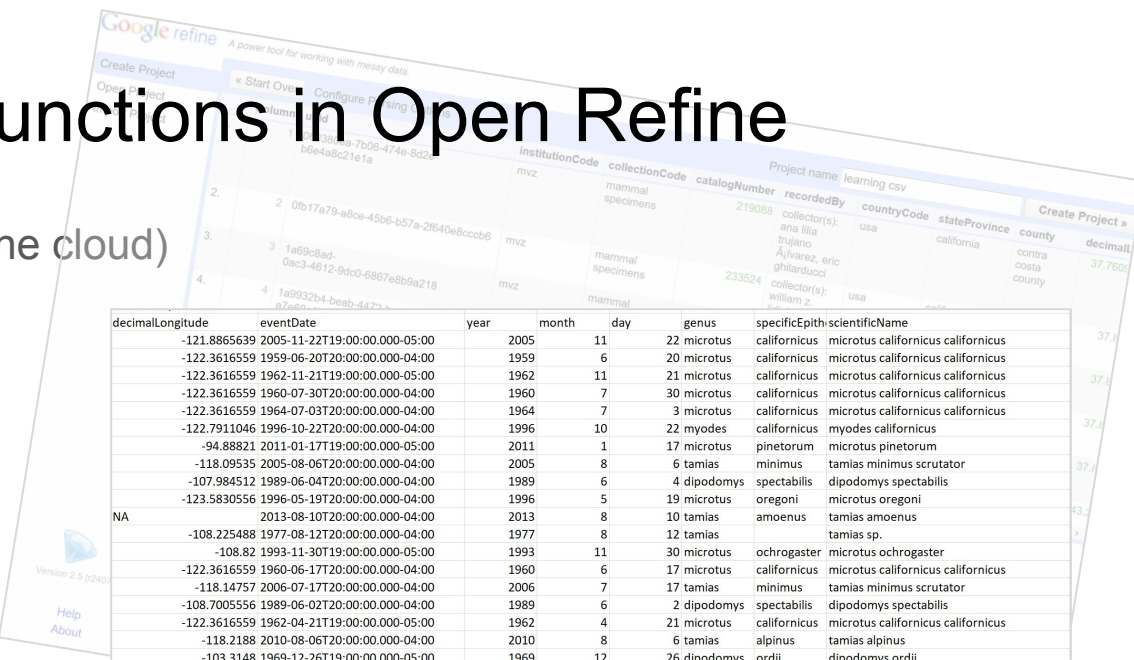


A	B	C	D	E
id	id gn barcode value	desc:basisOfRecord	desc:catNumber	desc:catalogNumber
1	515ab548-0c1a-4a53-9212-f68a90702a9	preservedspecimen	ouac	474848
2	52182933-ab95-442a-ba20-4d9933be21af	preservedspecimen	cuu_ent0017464	inse
4	533c7312-3970-4e12-8190-917752c94899	preservedspecimen	cuu_ent0009279	inse
5	5515a8f4-0d6a-425e-404e-172263ca3b84	preservedspecimen	cuu_ent0014106	inse
6	559e6e45-d60b-452d-871e-9f3131962a55	preservedspecimen	ouac	479715
7	5737e4dc-e600-4565-9676-93cf71913123	preservedspecimen	cuu_ent0026580	inse
8	1e87c060-c52d-4f05-b609-f8282a4f3869	preservedspecimen	ouac	471442
9	1a0f6460-1c6c-440b-b08c-96a3272f2a2a	preservedspecimen	asuac0045568	inse
10	1899a27f-0364-4b7c-8e5e-18993082815	preservedspecimen	ouac	474825
11	222770e9-2bb9-44e8-b477-13145632f512	preservedspecimen	ouac	454183
12	1393b0e2-82c3-42b1-bd74-90494e3c9a34	preservedspecimen	ouac	465205
13	158bc3d9-0c37-4c90-8c2c-2c8b948c8e28	preservedspecimen	cuu_ent0017708	inse
14	15c78b66-14bb-4e4a-8972-4921f82eabc3	preservedspecimen	ouac	475101

Carabidae (beetles) of California

(Fun!) features and functions in Open Refine

- runs on your computer (not in the cloud)
- data formats supported
- raw data
- column manipulation
- text facet
- routine cleaning (white space)
- clustering
- step-wise editable task script
- APIs
- regular expressions
- export
- share project files



The screenshot shows the Open Refine interface with a data table and a task script. The table has columns for decimalLongitude, eventDate, year, month, day, genus, specificEpith, and scientificName. The task script below the table shows a series of operations like 'clean', 'cluster', and 'facet' applied to the data.

decimalLongitude	eventDate	year	month	day	genus	specificEpith	scientificName
-121.8865639	2005-11-22T19:00:00.000-05:00	2005	11	22	microtus	californicus	microtus californicus californicus
-122.3616559	1959-06-20T20:00:00.000-04:00	1959	6	20	microtus	californicus	microtus californicus californicus
-122.3616559	1962-11-21T19:00:00.000-05:00	1962	11	21	microtus	californicus	microtus californicus californicus
-122.3616559	1960-07-30T20:00:00.000-04:00	1960	7	30	microtus	californicus	microtus californicus californicus
-122.3616559	1964-07-03T20:00:00.000-04:00	1964	7	3	microtus	californicus	microtus californicus californicus
-122.7911046	1996-10-22T20:00:00.000-04:00	1996	10	22	myodes	californicus	myodes californicus
-94.88821	2011-01-17T19:00:00.000-05:00	2011	1	17	microtus	pinetorum	microtus pinetorum
-118.09535	2005-08-06T20:00:00.000-04:00	2005	8	6	tamias	minimus	tamias minimus scrutator
-107.984512	1989-06-04T20:00:00.000-04:00	1989	6	4	dipodomys	spectabilis	dipodomys spectabilis
-123.5830556	1996-05-19T20:00:00.000-04:00	1996	5	19	microtus	oregoni	microtus oregoni
NA	2013-08-10T20:00:00.000-04:00	2013	8	10	tamias	amoenus	tamias amoenus
-108.225488	1977-08-12T20:00:00.000-04:00	1977	8	12	tamias		tamias sp.
-108.82	1993-11-30T19:00:00.000-05:00	1993	11	30	microtus	ochrogaster	microtus ochrogaster
-122.3616559	1960-06-17T20:00:00.000-04:00	1960	6	17	microtus	californicus	microtus californicus californicus
-118.14757	2006-07-17T20:00:00.000-04:00	2006	7	17	tamias	minimus	tamias minimus scrutator
-108.7005556	1989-06-02T20:00:00.000-04:00	1989	6	2	dipodomys	spectabilis	dipodomys spectabilis
-122.3616559	1962-04-21T19:00:00.000-05:00	1962	4	21	microtus	californicus	microtus californicus californicus
-118.2188	2010-08-06T20:00:00.000-04:00	2010	8	6	tamias	alpinus	tamias alpinus
-103.3148	1969-12-26T19:00:00.000-05:00	1969	12	26	dipodomys	ordii	dipodomys ordii
-107.464764	2012-10-26T20:00:00.000-04:00	2012	10	26	dipodomys	merriami	dipodomys merriami
NA	1968-01-17T19:00:00.000-05:00	1968	1	17	microtus	ochrogaster	microtus ochrogaster
-118.19192	2007-09-18T20:00:00.000-04:00	2007	9	18	tamias	minimus	tamias minimus scrutator
-110.78851	2011-07-20T20:00:00.000-04:00	2011	7	20	tamias	rufus	tamias rufus
-78.7133	1993-08-25T20:00:00.000-04:00	1993	8	25	myodes	gapperi	myodes gapperi
NA	1960-03-01T19:00:00.000-05:00	1960	3	1	microtus	californicus	microtus californicus californicus
-113.6134	2010-06-12T20:00:00.000-04:00	2010	6	12	tamias	amoenus	tamias amoenus
-120.1599	2006-08-18T20:00:00.000-04:00	2006	8	18	tamias	minimus	tamias minimus scrutator

Open Refine - getting started is quick and easy

- download and install
- launch
- import your data
- your raw data is NOT touched
- supported data formats
- subset data

Google refine A power tool for working with messy data.

Project name: learning.csv Create Project »

Column	uid	InstitutionCode	collectionCode	catalogNumber	recordedBy	countryCode	stateProvince	county	decimal
1.	060380ea-7b06-474e-8d2e-b6e4a8c21e1a	mvz	mammal specimens	219088	collector(s): ana lilia trujano Álvarez, eric ghilarducci	usa	california	contra costa county	37.7601
2.	0fb17a79-a8ce-45b6-b57a-2f640e8ccb6	mvz	mammal specimens	233524	collector(s): william z. lidicker jr.	usa	california	contra costa county	37.1
3.	1a69c8ad-0ac3-4612-9dc0-6867e8b9a218	mvz	mammal specimens	234346	collector(s): william z. lidicker jr.	usa	california	contra costa county	37.1
4.	1a9932b4-beab-4472-bec1-a7e68c4b9e6e	mvz	mammal specimens	233951	collector(s): william z. lidicker jr.	usa	california	contra costa county	37.1
5.	1f3b8aea-fbae-46d1-91c8-274924b40c9f	mvz	mammal specimens	235290	collector(s): william z. lidicker jr.	usa	california	contra costa county	37.1
6.	203f0531-9b46-403f-ac09-3acab5be977c	uam	mammal specimens	85106	collector(s): tom manning	usa	oregon	douglas county	43.1

Parse data as

Character encoding:

Update Preview

CSV / TSV / separator-based files

Columns are separated by

- commas (CSV)
- tabs (TSV)
- custom ,

Escape special characters with \

- Ignore first 0 line(s) at beginning of file
- Parse next 1 line(s) as column headers
- Discard initial 0 row(s) of data
- Load at most 0 row(s) of data

- Parse cell text into numbers, dates, ...
- Store blank cells as nulls
- Store file source (file names, URLs) in each row

Version 2.5 (2407)

Help About

Open Refine - managing columns

- reorganize columns easily

The screenshot shows the Open Refine interface with a data table and a modal dialog box for managing columns.

Table Data:

logNumber	recordedBy	countryCode	stateProvince
219088	collector(s): ana lilia trujano Álvarez, eric ghilarducci	usa	calif
233524	collector(s): william z. lidicker jr.	usa	calif
234346	collector(s): william z. lidicker jr.	usa	calif
233951	collector(s): william z. lidicker jr.	usa	calif
235290	collector(s): william z. lidicker jr.	usa	calif
85106	collector(s): tom manning; preparator(s): amber baxter	usa	oreg
50048	caidwell, j. p. and vitt, l. j.	usa	okla
216309	collector(s): james l. patton	usa	calif
294933	collector(s): troy l. best; preparator(s): troy l. best	usa	new
50255	collector(s): karl j. martin; preparator(s): paul ollig	usa	oreg

Re-order / Remove Columns Dialog:

Drag columns to re-order: uuid, institutionCode, collectionCode, catalogNumber, recordedBy, countryCode, stateProvince, county, decimalLatitude, decimalLongitude, eventDate, year, month, day, genus

Drop columns here to remove: (empty)

Buttons: OK, Cancel

Open refine - text facet

lists and counts the distinct values in a column

Facet / Filter Undo / Redo 1 **10767 rows**

Refresh Reset All Remove All Show as: rows record

scientificName change

162 choices Sort by: name count Cluster

cipodomys agilis	1
clethrionomys gapperi gapperi	3
dipodomis agilis	1
dipodomys agilis	1
dipodomys agilia	1
dipodomys agilis	5
dipodomys agilis perplexus	13
dipodomys agilis simulans	4
dipodomys agilus	1
dipodomys californicus	1
dipodomys californicus	
californicus	35

scientificName
microtus californicus californicus
microtus californicus californicus
microtus californicus californicus
microtus californicus californicus
microtus californicus californicus
microtus californicus californicus
myodes californicus

Open Refine - the magic of clustering algorithms

or how to find issues that abc sort won't and fix them all at once - no hunting

Method: key collision Keying Function: fingerprint 16 clusters found

Cluster Size	Row Count	Values in Cluster	Merge?	New Cell Value
2	59	<ul style="list-style-type: none"> dipodomys deserti deserti (54 rows) dipodomys deserti (5 rows) 	<input type="checkbox"/>	dipodomys deserti deserti
2	227	<ul style="list-style-type: none"> microtus pennsylvanicus (217 rows) microtus pennsylvanicus pennsylvanicus (10 rows) 	<input type="checkbox"/>	microtus pennsylvanicus
2	36	<ul style="list-style-type: none"> dipodomys californicus californicus (35 rows) dipodomys californicus (1 rows) 	<input type="checkbox"/>	dipodomys californicus califoi
2	23	<ul style="list-style-type: none"> tamias panamintinus panamintinus (17 rows) tamias panamintinus (6 rows) 	<input type="checkbox"/>	tamias panamintinus panami
2	2319	<ul style="list-style-type: none"> microtus californicus californicus (2295 rows) microtus californicus (24 rows) 	<input type="checkbox"/>	microtus californicus californi
2	46	<ul style="list-style-type: none"> dipodomys microps (38 rows) dipodomys microps microps (8 rows) 	<input type="checkbox"/>	dipodomys microps
2	6	<ul style="list-style-type: none"> dipodomys agilis (5 rows) dipodomys agilis (1 rows) 	<input type="checkbox"/>	dipodomys agilis

Rows in Cluster

0 — 2400

Average Length of Choices

15 — 31

Length Variance of Choices

0.5 — 7.5

Open Refine - manages pesky white spaces

/: 5 10 25 50 rows

specificEpithet	scientificName	weight	length	sex
californicus	Facet	30.5	165	male
californicus	Text filter			
californicus	Edit cells			
Trim leading and trailing whitespace	Transform...			
Collapse consecutive whitespace	Common transforms			
Unescape HTML entities	Fill down			
To titlecase	Blank down			
To uppercase	Split multi-valued cells...			
To lowercase	Join multi-valued cells...			
To number	Cluster and edit...	23.5	141	female
To date				
To text		24	121	
Blank out cells		27	176	female

Open Refine - add data to your data using APIs *that's application programming interface*

Add column by fetching URLs based on column scientificName

New column name Throttle delay milliseconds

On error set to blank store error

Formulate the URLs to fetch:

Expression Language

```
"http://webservice.catalogueoflife.org  
/col/webservice?scientificName="+escape(value, 'url')
```

No syntax error.

Preview History Starred Help

row	value	"http://webservice.catalogueoflife.org /col/webservice?scientificName="+escape(value,'url')
1.	microtus californicus californicus	http://webservice.catalogueoflife.org /col/webservice?scientificName=microtus+californicus+californicus
2.	microtus californicus californicus	http://webservice.catalogueoflife.org /col/webservice?scientificName=microtus+californicus+californicus
3.	microtus californicus californicus	http://webservice.catalogueoflife.org /col/webservice?scientificName=microtus+californicus+californicus
4.	microtus californicus californicus	http://webservice.catalogueoflife.org /col/webservice?scientificName=microtus+californicus+californicus

OK Cancel

Open Refine - saves your steps

☐ *supports
reproducibility*

☐ *tracks your work
for you*

☐ *easy to go back to
earlier steps with
confidence*

Extract Operation History

Extract and save parts of your operation history as JSON that you can apply to this or other projects in the future.

- Reorder columns
- Mass edit cells in column scientificName
- Mass edit cells in column scientificName
- Create column c at index 4 by fetching URLs based on column scientificName using expression
grel:"http://webservice.catalogueoflife.org/col/webservice?scientificName="+escape(value,'url')

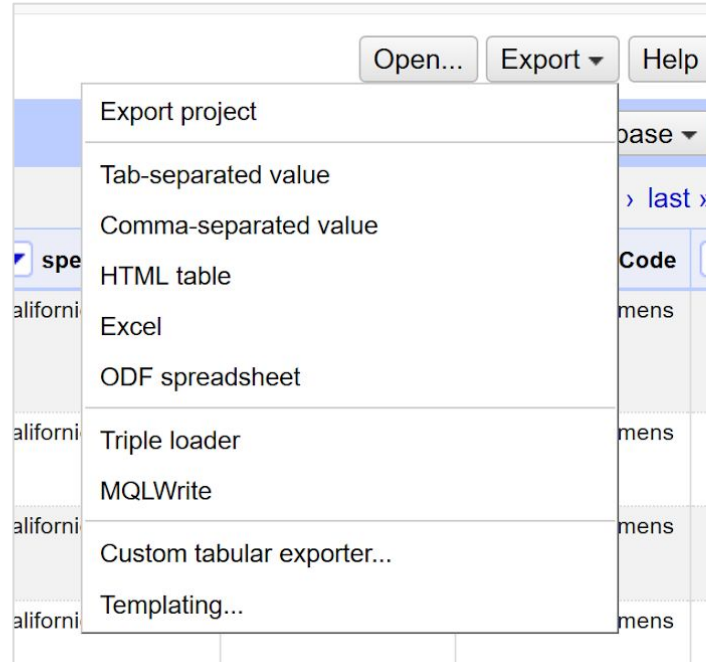
```
{
  "v": {
    "v": "dipodomys agilis",
    "l": "dipodomys agilis"
  },
  "v": {
    "v": "microtus californicus sc",
    "l": "microtus californicus sc"
  }
},
"selectError": false,
"invert": false,
"name": "scientificName",
"omitBlank": false,
"type": "list",
"columnName": "scientificName"
}
],
"newColumnName": "c",
"columnInsertIndex": 4,
"baseColumnName": "scientificName",
"urlExpression": "grel:\`http://webservice",
"onError": "set-to-blank",
"delay": 250
}
]
```

Select All Unselect All

Close

Open Refine - export your data, share project files

select the *format*
export *subsets* too
and *project files*



Open Refine - make some friends

- share this tool with students, friends, families, colleagues
- **imagine future tools, think beyond spreadsheets**

Increase Reproducibility and Productivity
using tools like Open Refine

Magic is here.
Ask for it, plan for it.



CUAHSI DataONE
Universities Allied for Water Research



GBIF
iDigBio
Integrated Digital Biodiversity Collections



neon
Operated by Battelle



NEOTOMA PALEOECOLOGY DATABASE

Looking for next steps now?

R, Open Refine, and Data Management resources

- The #datahelpdesk is ready to offer data assistance!
- #CareerCentral Q and A: Wednesday, August 5th, 9:30-10:30 PDT (12:30-1:00 EDT)
- Data Help Desk Wiki <https://bit.ly/datahelpesa2020>
- Data Carpentry lessons
- on Twitter #ESA2020 #datahelpdesk

