# Towards Capturing Provenance of the Data Curation Process at Domain-specific Repositories

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#### 20,000 Leagues Under the Sea by Jules Verne

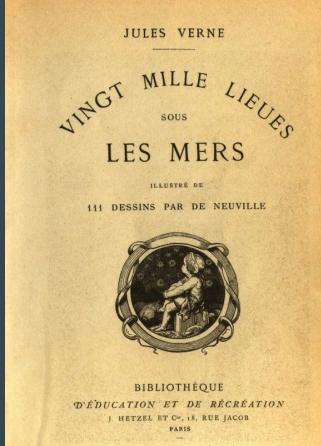


#### 20,000 Leagues Under the Sea

#### Plural 'Seas' became the singular 'Sea'

thus...

# 20,000 leagues was misinterpreted to mean **depth** instead of **distance**



Droits de traduction et de reproduction réservés.

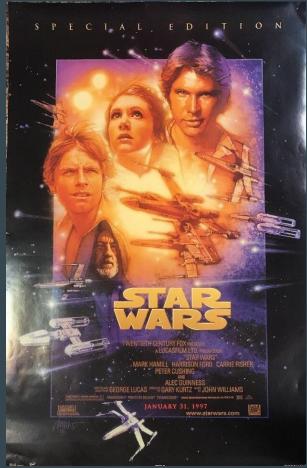
#### Lost in Translation

When variations are **perceived** to have changed the meaning of a work, there is potential for a **loss of trust**.



#### **1977** Star Wars





https://cdn.shopify.com/s/files/1/0371/9417/products/Original\_Star\_Wars080618PF\_eca1292e-66e9-417e-a811-43f2728f5542\_1024x1024.jpg?v=1542461664 https://d1w8cc2yygc27j.cloudfront.net/-1472768621735153225/1072936435962636846.jpg

#### **1997: A Disturbance in the Force**













# A single change in a split second altered the nature of a character.

#### **Did Han Shoot First?**





Biological & Chemical Oceanography Data Management Office

### **Submitted Columns Names**

A

st1\_50m

37:2 alkenone

## A Case of 'st1\_50m'

#### An observation made at the location of

station '1' at a depth of 50 meters

best hit annotation	best bit tayon id	st1_050m	st1 090m	ct1 120m	st1 200m	st1 300m	st1 400m	st1 600m	st3 040m	st3 060m	st3 120m	st3 1
nitrate reductase alpha subunit	247490		0 03011 0	3t1_120111 0	3t1_200m	122 sti	121	116	3t3_040111 0	313_000m 0	313_12011 17	313_1
nxrB1; putative nitrate oxidoreductase subunit beta (E	330214	0	0	0	1	136	173	153	0	0	18	
groEL; chaperonin GroEL (EC:3.6.4.9); K04077 chapero	167546	80	91	59	35	2	2	155	60	44	24	
ccmK; carboxysome shell protein CsoS1	167555	155	162	94	38	0	0		54	44	39	
putative UreA ABC transporter; substrate binding prot	167546	202	203	169	158	26	30	19	100	40 86	27	
	859653	202	205	8	158	20 51		74	3	1	19	
ligand-binding protein; OpuAC family		17	-	-			69	36	_			
ABC transporter	314261		20	19	9	30	35		12	15	22	
ABC transporter; substrate-binding protein; family 5	89187	0	0	0	0	50	50	65	0	0	2	
gInA; glutamine synthetase; glutamateammonia liga:	146891	62	60	54	58	3	4	2	60	53	4	
amino acid ABC transporter substrate-binding protein	913324	3	2	2	4	33	78	43	0	0	6	
F0F1 ATP synthase subunit beta	93058	57	63	76	34	5	4	3	39	40	47	
peptide ABC transporter; periplasmic substrate-bindin	375451	0	2	3	0	41	48	44	0	0	6	
glutamate/glutamine/aspartate/asparagine ABC trans	488538	2	7	11	2	31	52	44	2	3	5	
hypothetical protein	1090946	1	1	7	0	51	47	37	0	0	5	
ABC transporter binding protein	859653	88	68	33	29	1	4	3	38	32	10	
rbcL; ribulose bisophosphate carboxylase; K01601 ribu	146891	46	57	40	36	1	3	0	37	41	15	
nd	1073573	20	10	2	1	26	44	29	10	8	2	
formate dehydrogenase subunit alpha (EC:1.2.1.2); KO	639282	0	0	0	0	37	41	36	0	0	3	
amino acid ABC transporter substrate-binding protein,	644966	0	0	0	0	29	38	31	0	0	1	
ligand-binding protein; OpuAC family	859653	25	16	15	9	50	55	41	12	14	19	
glutamate/glutamine/aspartate/asparagine ABC trans	488538	0	14	18	0	36	45	33	0	0	22	
nd	1073573	4	5	1	2	35	27	35	0	0	3	
amino acid ABC transporter substrate-binding protein	859653	35	31	34	16	25	24	14	25	22	26	
TonB-dependent receptor plug	518766	0	0	0	0	38	55	31	0	0	2	
hypothetical protein	926566	0	0	0	0	19	64	21	0	0	0	
chlorophyll a/b binding light harvesting protein PcbD;	146891	8	69	91	8	3	3	0	32	25	0	
Phycobilisome protein	221359	2	26	10	4	0	0	0	3	2	0	
nd	4577	22	22	16	9	9	15	9	4	5	6	
ABC transporter	439493	31	29	21	10	7	4	3	28	24	17	
TIGR00065: ftsZ: cell division protein EtsZ PE00091 20:	35677	28	27	28	17	28	20	9	13	11	15	

#### **Archive Quality**

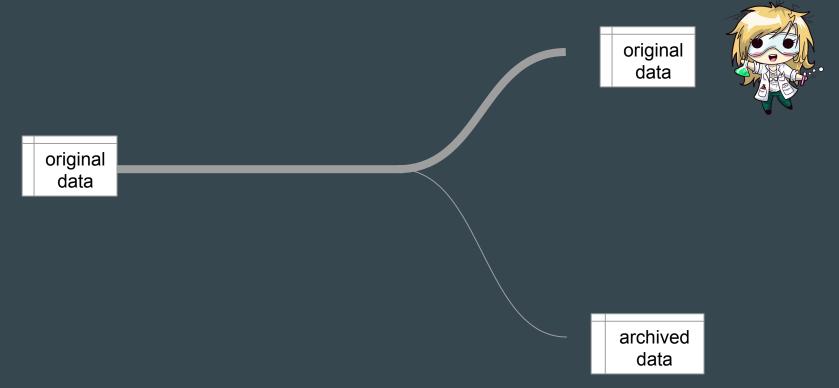
Why split these out into explicit columns if the science community prefers the original format?

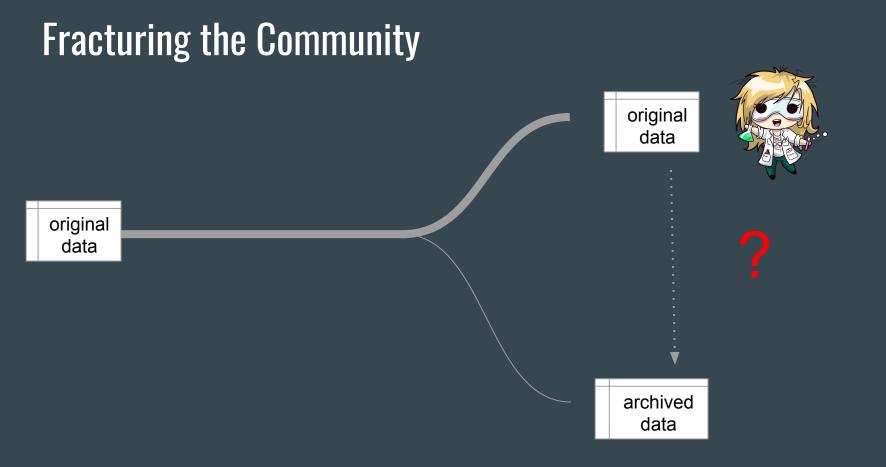
#### Putting the FIR in FAIR

**F**indable - get all datasets that measured 'depth' **I**nteroperable - linked to community vocabs **R**eusable - (meta)data with **provenance** 

	station	depth	spectral_count
5	8	200	0
5	9	40	12
5	9	70	4
5	9	380	0
L	12	40	0
L	12	120	0
L	12	300	0
ł	1	50	0
ł	1	90	0
ł	1	120	0
ł	1	200	0
ł	1	300	0
ł	1	400	0
ł	1	600	0
6	3	40	9
5	2	60	0

### Fracturing the Community





How do we capture provenance of the data curation process that links original & archived versions in a reusable way ?

#### **Declarative Workflows**

```
lat_lon_DDM_to_DD:
title: lat_lon_DDM_to_DD
description: "Add lat and lon columns in decimal degrees (DD) given one column with\
    \ lat and lon in format degrees decimal minutes (DDM) (e.g. \"77\xB0 51.3'S 166\xB0\
    \ 40.1'E\"). "
pipeline:
    - run: add_resource
    parameters:
        name: mcmurdo_epifauna,
        url: 'http://datadocs.bco-dmo.org/docs/TestProject/data_docs/latlon_DDM_to_DD/McMurdoEpifauna.xlsx',
        format: xlsx,
        sheet: animals,
        headers: 1,
```

#### **Declarative Workflows - A set of steps to execute**

```
lat_lon_DDM_to_DD:
title: lat_lon_DDM_to_DD
description: "Add lat and lon columns in decimal degrees (DD) given one column with\
  \ lat and lon in format degrees decimal minutes (DDM) (e.g. \"77\xB0 51.3'S 166\xB0\
  \ 40.1'E\"). "
pipeline:
  - run: add_resource
parameters:
    name: mcmurdo_epifauna,
    url: 'http://datadocs.bco-dmo.org/docs/TestProject/data_docs/latlon_DDM_to_DD/McMurdoEpifauna.xlsx',
    format: xlsx,
    sheet: animals,
    headers: 1,
```

#### Declarative Workflows - Each step is "named"

```
lat_lon_DDM_to_DD:
title: lat_lon_DDM_to_DD
description: "Add lat and lon columns in decimal degrees (DD) given one column with\
  \ lat and lon in format degrees decimal minutes (DDM) (e.g. \"77\xB0 51.3'S 166\xB0\
  \ 40.1'E\"). "
pipeline:
  - run: add_resource
parameters:
    name: mcmurdo_epifauna,
    url: 'http://datadocs.bco-dmo.org/docs/TestProject/data_docs/latlon_DDM_to_DD/McMurdoEpifauna.xlsx',
    format: xlsx,
    sheet: animals,
    headers: 1,
```

#### **Declarative Workflows - Each step has inputs**

```
lat_lon_DDM_to_DD:
title: lat_lon_DDM_to_DD
description: "Add lat and lon columns in decimal degrees (DD) given one column with\
  \ lat and lon in format degrees decimal minutes (DDM) (e.g. \"77\xB0 51.3'S 166\xB0\
  \ 40.1'E\"). "
pipeline:
  - run: add_resource
parameters:
  name: mcmurdo_epifauna,
  url: 'http://datadocs.bco-dmo.org/docs/TestProject/data_docs/latlon_DDM_to_DD/McMurdoEpifauna.xlsx',
  format: xlsx,
    sheet: animals,
    headers: 1,
```

#### **Declarative Workflows - More steps**

<pre>lat_lon_DDM_to_DD: title: lat_lon_DDM_to_DD description: "Add lat and lon columns in decimal degrees (DD) \ lat and lon in format degrees decimal minutes (DDM) (e.g. \ 40.1'E\"). " pipeline:</pre>	The second							
<pre>pipeline: - run: ad - run: bcodmo_pipeline_processors.convert_to_dec</pre>	imal degrees							
paramet cache: True								
nam parameters:								
url resources: [mcmurdo_epifauna]								
for fields:								
<pre>she - {input_field: lat_long, format: degrees-de</pre>								
<pre>hea pattern: "(?P<degrees>.*)\xB0 (?P<decimal_< pre=""></decimal_<></degrees></pre>	pattern: "(?P <degrees>.*)\xB0 (?P<decimal_minutes>.*)'(?P<directional>.)\\ .*\xB0 .*'."}</directional></decimal_minutes></degrees>							
<pre>- run: bcodmo_pipeline_processors.convert_to_dec</pre>	- run: bcodmo_pipeline_processors.convert_to_decimal_degrees							
cache: true								
parameters:								
resources: [mcmurdo_epifauna]								
fields:								
<pre>- {input_field: lat_long, format: degrees-de</pre>	- {input_field: lat_long, format: degrees-decimal_minutes, output_field: long_converted, directional: '',							
<pre>pattern: ".*\xB0 .*'. (?P<degrees>.*)\xB0</degrees></pre>	pattern: ".*\xB0 .*'. (?P <degrees>.*)\xB0 (?P<decimal_minutes>.*)'(?P<directional>.)"}</directional></decimal_minutes></degrees>							

#### Declarative Workflows - Names identify code to execute

```
lat lon DDM to DD:
 title: lat_lon_DDM_to_DD
 description: "Add lat and lon columns in decimal degrees (DD) given one column with\
   \ lat and lon in format degrees decimal minutes (DDM) (e.g. \"77\xB0 51.3'S 166\xB0\
   \ 40.1'E\"). "
 pipeline:
              - run: bcodmo_pipeline_processors.convert_to_decimal_degrees
 - run: ad
                cache: True
   paramet
                parameters:
       nam
                  resources: [mcmurdo_epifauna]
       url
                  fields:
       for
                  - {input_field: lat_long, format: degrees-decimal_minutes, output_field: lat_converted, directional: '',
       she
                     pattern: "(?P<degrees>.*)\xB0 (?P<decimal_minutes>.*)'(?P<directional>.)\\ .*\xB0 .*'."}
       hea
              - run: bcodmo pipeline processors.convert to decimal degrees
                cache: true
                parameters:
                  resources: [mcmurdo_epifauna]
                  fields:
                  - {input_field: lat_long, format: degrees-decimal_minutes, output_field: long_converted, directional: '',
                     pattern: ".*\xB0 .*'. (?P<degrees>.*)\xB0 (?P<decimal minutes>.*)'(?P<directional>.)"}
```

#### Declarative Workflows - Each step has its own inputs

```
lat lon DDM to DD:
 title: lat_lon_DDM_to_DD
 description: "Add lat and lon columns in decimal degrees (DD) given one column with\
   \ lat and lon in format degrees decimal minutes (DDM) (e.g. \"77\xB0 51.3'S 166\xB0\
   \ 40.1'E\"). "
 pipeline:
              - run: bcodmo_pipeline_processors.convert_to_decimal_degrees
 - run: ad
                cache: True
   paramet
                parameters:
       nam
                  resources: [mcmurdo_epifauna]
       url
                   fields:
       for
                  - {input_field: lat_long, format: degrees-decimal_minutes, output_field: lat_converted, directional: '',
       she
                     pattern: "(?P<degrees>.*)\xB0 (?P<decimal minutes>.*)'(?P<directional>.)\\ .*\xB0 .*'."}
       hea
              - run: bcodmo pipeline processors.convert to decimal degrees
                cache: true
                parameters:
                   resources: [mcmurdo_epifauna]
                   fields:
                   - {input_field: lat_long, format: degrees-decimal_minutes, output_field: long_converted, directional: '',
                     pattern: ".*\xB0 .*'. (?P<degrees>.*)\xB0 (?P<decimal minutes>.*)'(?P<directional>.)"}
```

## **Benefits of Declarative Workflows**

Pipeline Title	lat_lon_DDM_to_DD						
nue							
Pipeline Description	Add lat and lon columns in decimal degrees (DD) given one column with lat and lon in format degrees decimal minutes (DDM) (e.g. "77° 51.3'S 166° 40.1'E").						
1	▲ Add resource + ► ×	C					
Processor	Add resource						
Name r	nust match the $([-a-z0-9.])+$ regular expression.						
Name	mcmurdo_epifauna						
URL	http://datadocs.bco-dmo.org/docs/TestProject/data_docs						
Format	xlsx	J					
Header	1						
1010 #							

Because declarative workflows build a configuration (WHAT) instead of code (HOW)

#### Build tools to:

- construct these configurations,
- execute the workflow,
- convert the configuration into provenance

### **Benefits of Declarative Workflows**

Pipeline lat lon DDM to DD Title Pipeline Add lat and lon columns in decimal degrees (DD) Description given one column with lat and lon in format degrees decimal minutes (DDM) (e.g. "77° 51.3'S 166° 40.1'E"). 1 Add resource Processor Add resource Name must match the ([-a-z0-9, /])+ regular expression. mcmurdo epifauna Name http://datadocs.bco-dmo.org/docs/TestProject/data docs URL Format xlsx Header 1 row #

#### lat\_lon\_DDM\_to\_DD:

title: lat\_lon\_DDM\_to\_DD

- description: "Add lat and lon columns in decimal degrees (DD) given one column with
- \ lat and lon in format degrees decimal minutes (DDM) (e.g. \"77\xB0 51.3'S 166\xB0\ 40.1'E\"). "

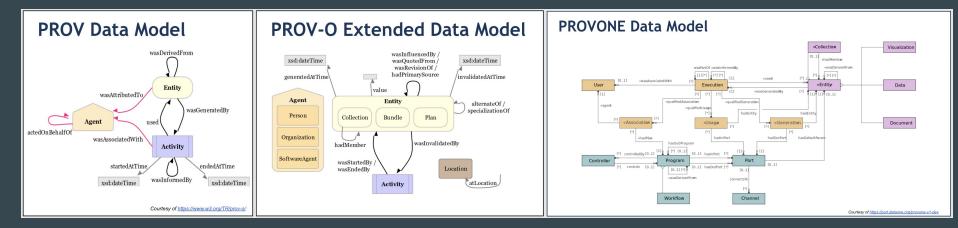
pipeline:

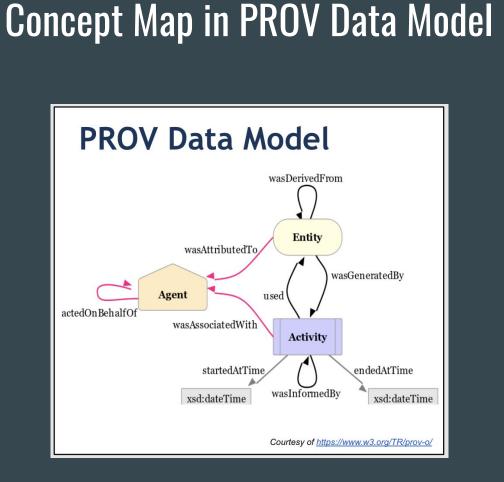
- run: add\_resource

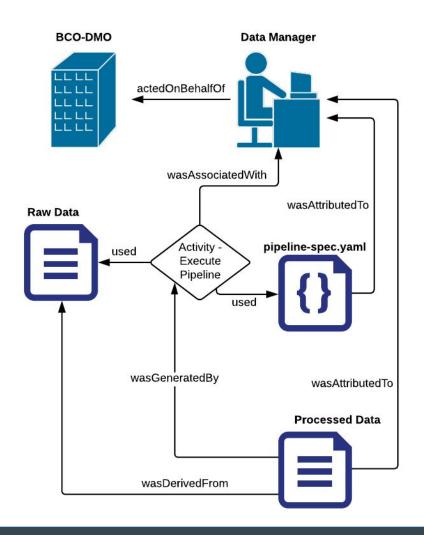
parameters:

- name: mcmurdo\_epifauna,
- url: 'http://datadocs.bco-dmo.org/docs/TestProject/data\_docs/latlon\_DDM\_to\_DD/McMurdoEpifauna.xlsx',
  format: xlsx,
- sheet: animals,
- headers: 1.
- run: stream\_remote\_resources
- cache: True
- parameters:
- resources: [mcmurdo\_epifauna]
- run: set\_types cache: True
- parameters:
- resources: [mcmurdo\_epifauna]
- types:
- year: {type: number}
- site: {type: string}
- lat\_long: {type: string}
- genus\_species: {type: number}
- run: bcodmo\_pipeline\_processors.add\_schema\_metadata
- cache: True
- parameters:
- resources: [mcmurdo\_epifauna]
- missingValues: ["nd"]
- run: bcodmo\_pipeline\_processors.convert\_to\_decimal\_degrees
- cache: True
- parameters:
- resources: [mcmurdo\_epifauna]
- fields:
- {input\_field: lat\_long, format: degrees-decimal\_minutes, output\_field: lat\_converted, directional: '', pattern: "(?P<degrees>.\*)\xB0 (?P<decimal\_minutes>.\*)'(?P<directional>.)\\. .\*\xB0 .\*',"}
- run: bcodmo\_pipeline\_processors.convert\_to\_decimal\_degrees
- cache: true
  - parameters:
  - resources: [mcmurdo\_epifauna]
  - fields:
  - {input\_field: lat\_long, format: degrees-decimal\_minutes, output\_field: long\_converted, directional: '',
    pattern: ".\*\xB0 .\*'. (?P<degrees-.\*)\xB0 (?P<decimal\_minutes>.\*)'(?P<directional>.)"}
- run: bcodmo\_pipeline\_processors.round\_fields
- cache: True
- parameters:
- resources: [mcmurdo\_epifauna]
- fields:
- {digits: 5, name: lat\_converted}
- run: bcodmo\_pipeline\_processors.round\_fields

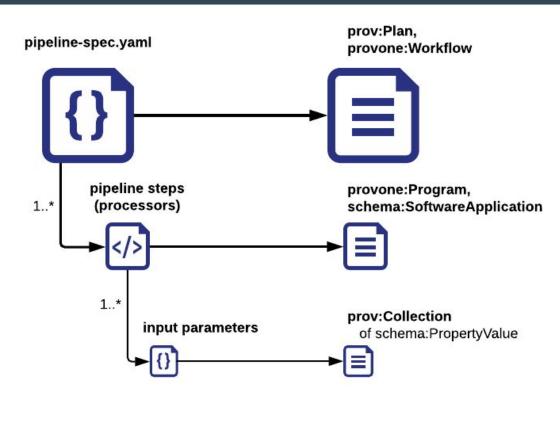
#### From Declarative Workflows to Provenance



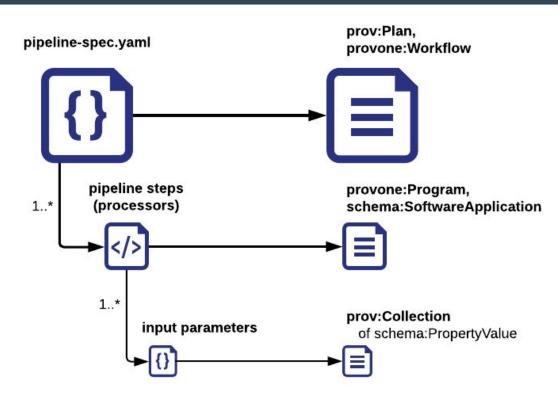




### PROV-O + Schema.org



### PROV-O + Schema.org



**Sub** propertyValueForParameter(parameter) prop\_value = new PropertyValue(parameter.name)

If parameter.value is an Object
prop\_value = propertyValueForParameter(parameter.value)
Else
prop\_value = parameter.value

Return prop\_value End Sub @prefix : <http://data.example.org/id/dataset/1234/v1/> .
@prefix dcterms: <http://purl.org/dc/terms/> .
@prefix voi: <http://www.w3.org/2002/07/vwli> .
@prefix provone: <http://www.w3.org/prov#> .
@prefix provone: <http://www.w3.org/pp0/2/22-rdf-syntax-ns#> .
@prefix rdfs: <http://www.w3.org/2000/01/rdf-schema#> .
@prefix schema: <http://www.w3.org/2001/rdf-schema#> .
@prefix xdfs: <http://www.w3.org/2001/rdf-schema#> .
@prefix schema: <http://www.w3.org/XML/1998/namespace> .
@prefix sd: <http://www.w3.org/2001/XMLSchema#> .

#### : a prov:Bundle,

prov:Entity ;

prov:generatedAtTime "2018-09-21T13:38:10+00:00"^^xsd:dateTime ;
prov:wasAttributedTo :alice .

#### :frictionless-data-pkg a schema:DigitalDocument,

#### prov:Data,

prov:Entity; schema:encodingFormat "application.vnd.datapackage+json"^^xsd:string; schema:url "https://example.org/dataset/1234/v1/datapackage.json"^^xsd:anyURI; prov:qualifiedGeneration [ a prov:Generation; proviactivity :executed-pipeline; prov:endTime "2018-09-21T13:37:53+00:00"^^xsd:dateTime; prov:wsd:GeneratedBy:executed-pipeline.

#### :processed-data a schema:Dataset,

prov:Entity ; schema:distribution [ a schema:DataDownload ; schema:contentUrl "https://example.org/dataset/l234/v1/McMurdoEpifauna.csv"^^xsd:anyURI ; schema:encodingFormat "text/csv"^^xsd:string ] ; proviadFrimarySource :raw-dat ; prov:qualifiedGeneration [ a prov:Generation ; prov:activity :executed-pipeline ; prov:endTime "2018-09-21T13:38:09+00:00"^^xsd:dateTime ; prov:startTime "2018-09-21T13:37:54+00:00"^^xsd:dateTime ] ; prov:wasDerivedFrom :pipeline.pipeline .

#### :step-1-add-resource a provone:Program,

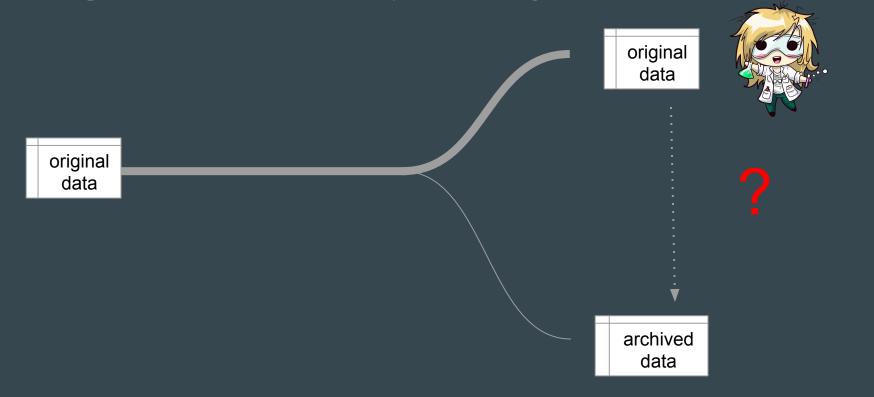
prov:Entity ;
schema:supportingData :step-1-add-resource-inputs .

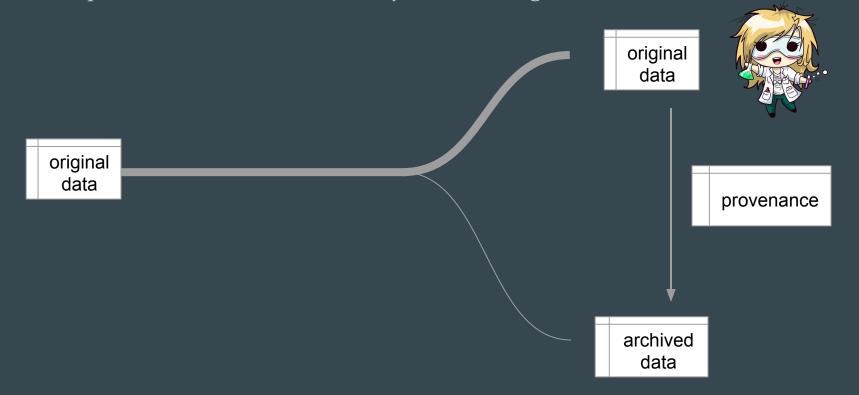
```
:step-1-add-resource-inputs a schema:DataFeed ;
```

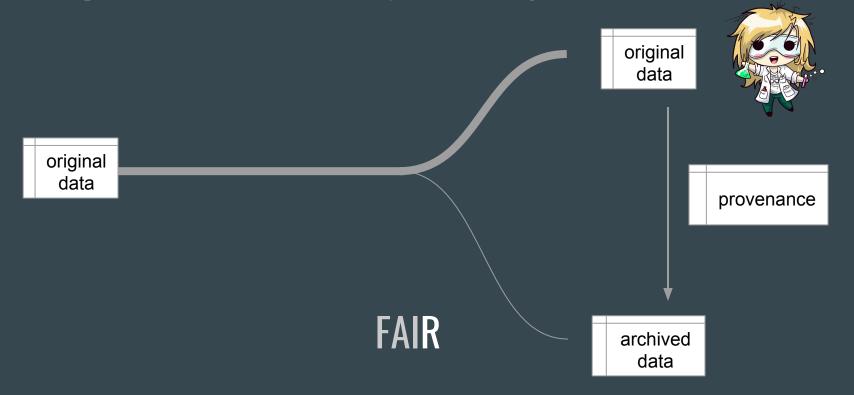
#### [ a schema:PropertyValue ;

#### Workflow to PROV

#### :step-1-add-resource-inputs a schema:DataFeed ; schema:dataFeedElement [ a prov:Collection ; rdfs:comment "A single step in pipeline."@en-US ; prov:hadMember [ a provone:Data, schema: PropertyValue, prov:Entity ; schema:name "run"^^xsd:string ; schema:value "add resource"^^xsd:string ], [ a provone:Data, schema: PropertyValue, prov:Entity ; schema:name "parameters"^^xsd:string ; schema:value [ a schema:PropertyValue ; schema:name "headers"^^xsd:string ; schema:value 1 1. [ a schema: PropertyValue ; schema:name "name"^^xsd:string ; schema:value "mcmurdo\_epifauna"^^xsd:string ], [ a schema: PropertyValue ; schema:name "url"^^xsd:string ; schema:value "https://example.org/dataset/1234/original/20180921T123456Z/McMurdoEpifauna.xlsx"^2 [ a schema: PropertyValue ; schema:name "format"^^xsd:string ; schema:value "xlsx"^^xsd:string ], [ a schema: PropertyValue ; schema:name "sheet"^^xsd:string ; schema:value "animals"^^xsd:string ] ] ] .







## Workflow Tools at BCO-DMO



frictionlessdata.io

github.com/frictionlessdata/datapackage-pipelines

pypi.org/project/dataflows

w3.org/TR/prov-o/

purl.dataone.org/provone-v1-dev

schema.org

