

Herramienta para la exploración de tendencias y detección de patrones epidemiológicos en Argentina

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CONICET



I P C S H

Instituto Patagónico de
Ciencias Sociales y Humanas

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del Castillo Bernal”**



**LABORATORIO DE CIENCIAS
DE LAS IMAGENES**

Motivación

El estudio de las causas de los fallecimientos constituye una fuente de información fundamental para la planificación y elaboración de políticas públicas de salud de un país.

Motivación

Saber por qué (a causa de qué) mueren las personas, constituye un insumo fundamental para entender cómo viven las personas.

Objetivos

Analizar el comportamiento temporal y espacial de los fallecimientos por causas específicas en la Argentina.

Objetivos

Analizar el comportamiento temporal y espacial de fallecimientos por causas específicas en la Argentina.

Repetir el análisis de forma sencilla, con diferentes conjuntos de causas específicas de fallecimientos, respetando los sistemas de codificación CIE-9 o CIE-10.

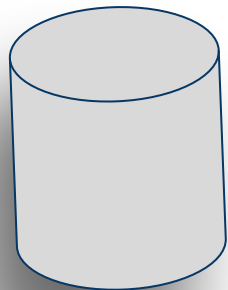
Herramienta

Pipeline, entradas, procesamiento y salidas



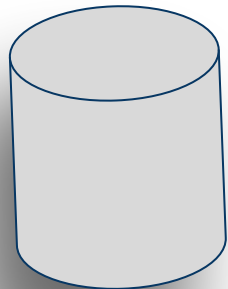
Pipeline, entradas, procesamiento y salidas

Registro de
fallecimientos del
Ministerio de Salud de
la Nación Argentina



Pipeline, entradas, procesamiento y salidas

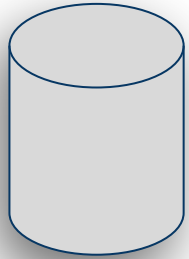
Registro de
fallecimientos del
Ministerio de Salud de
la Nación Argentina



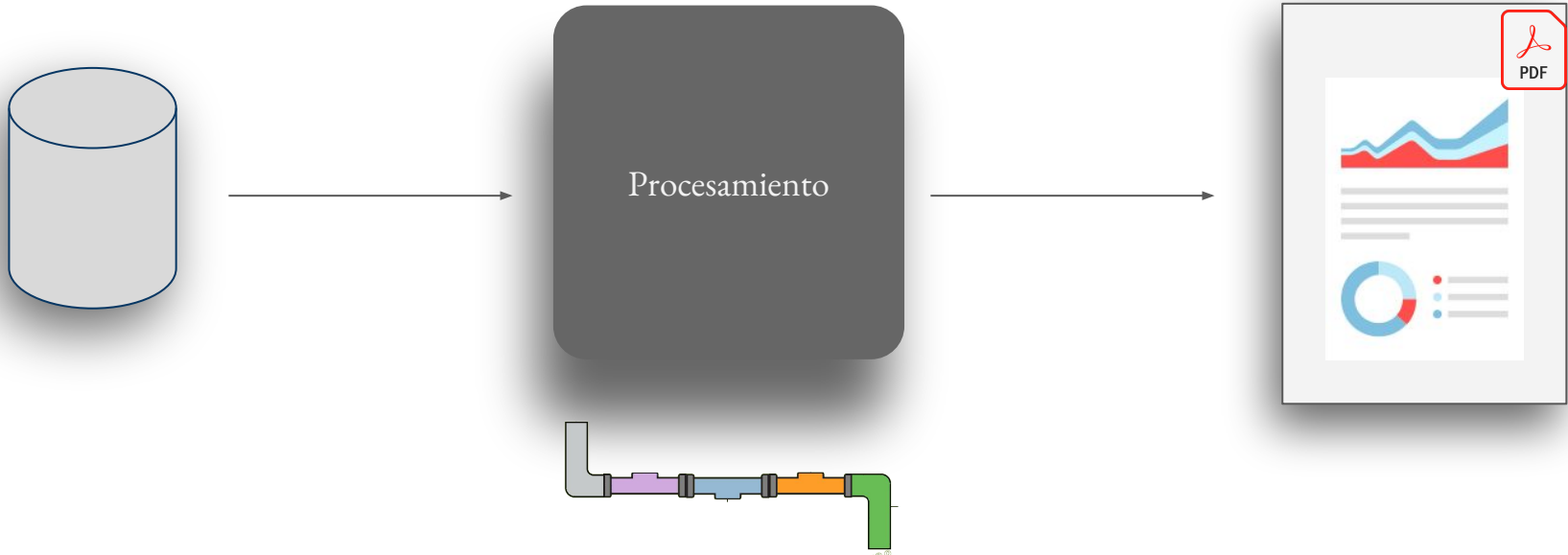
Atributos:
Edad, Sexo,
Provincia, Departamento
Causa (período CIE-9,
período CIE-10)

← | | | | | | | | | | ... | | | | | | | | | | →
1991 2017

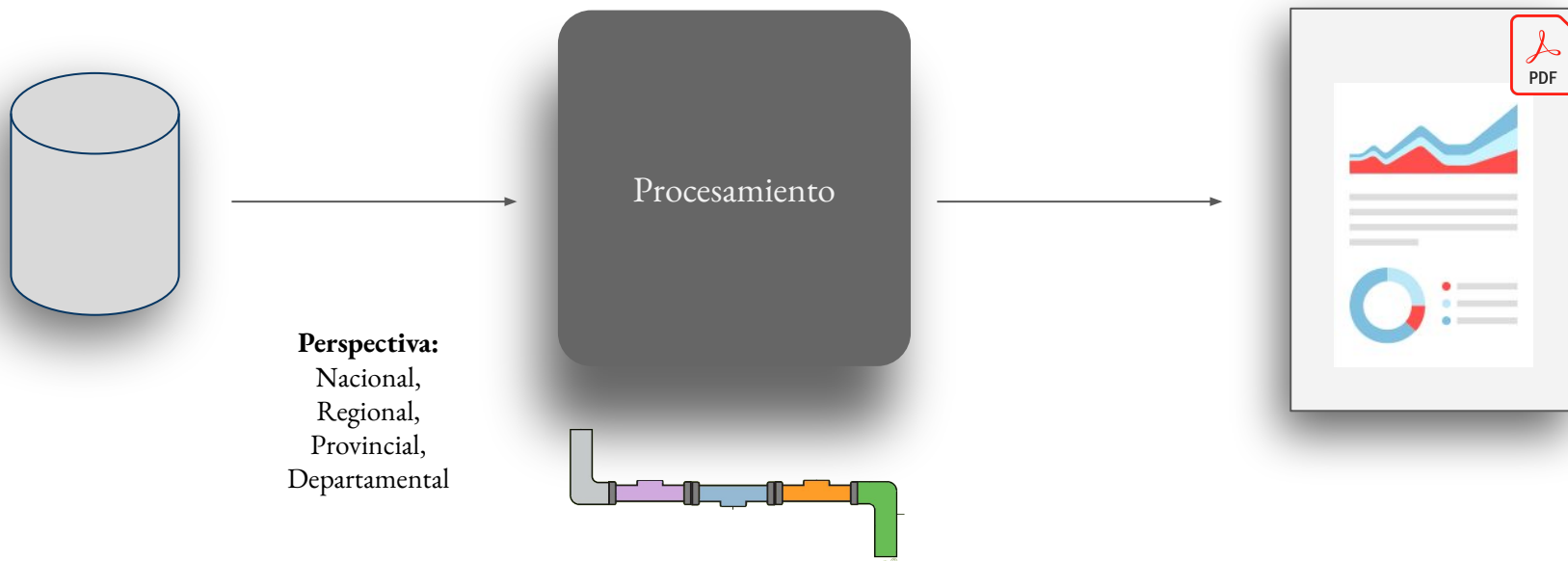
Pipeline, entradas, procesamiento y salidas



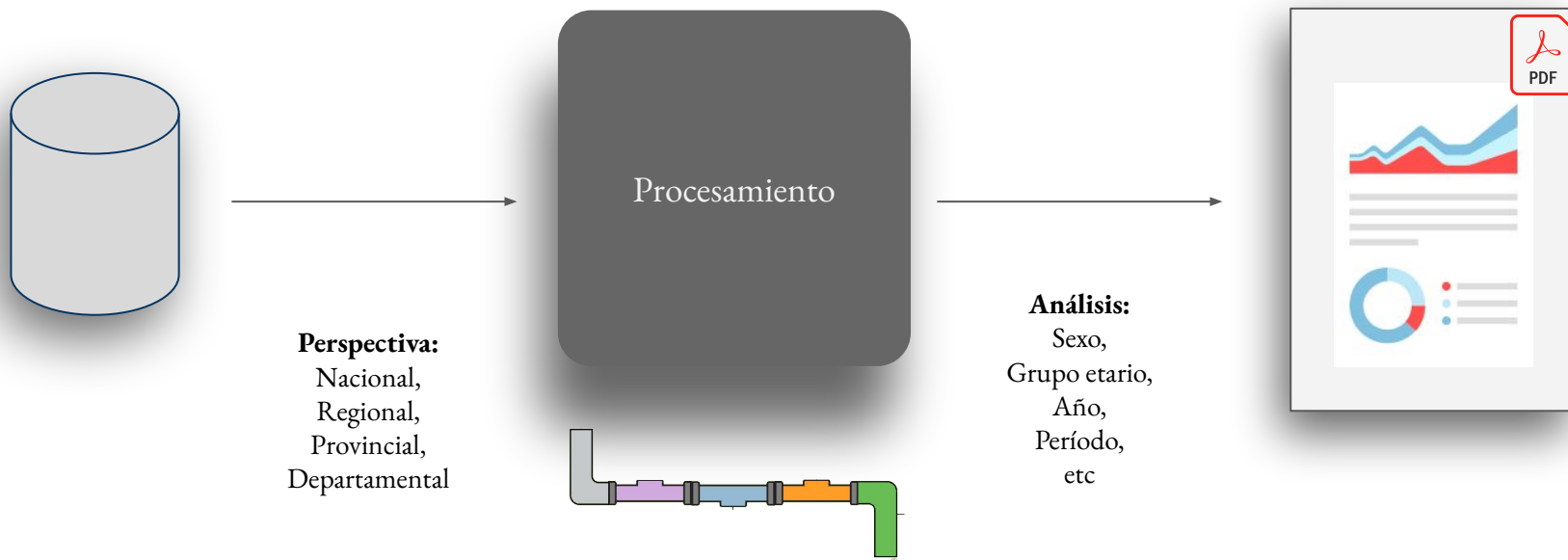
Pipeline, entradas, procesamiento y salidas



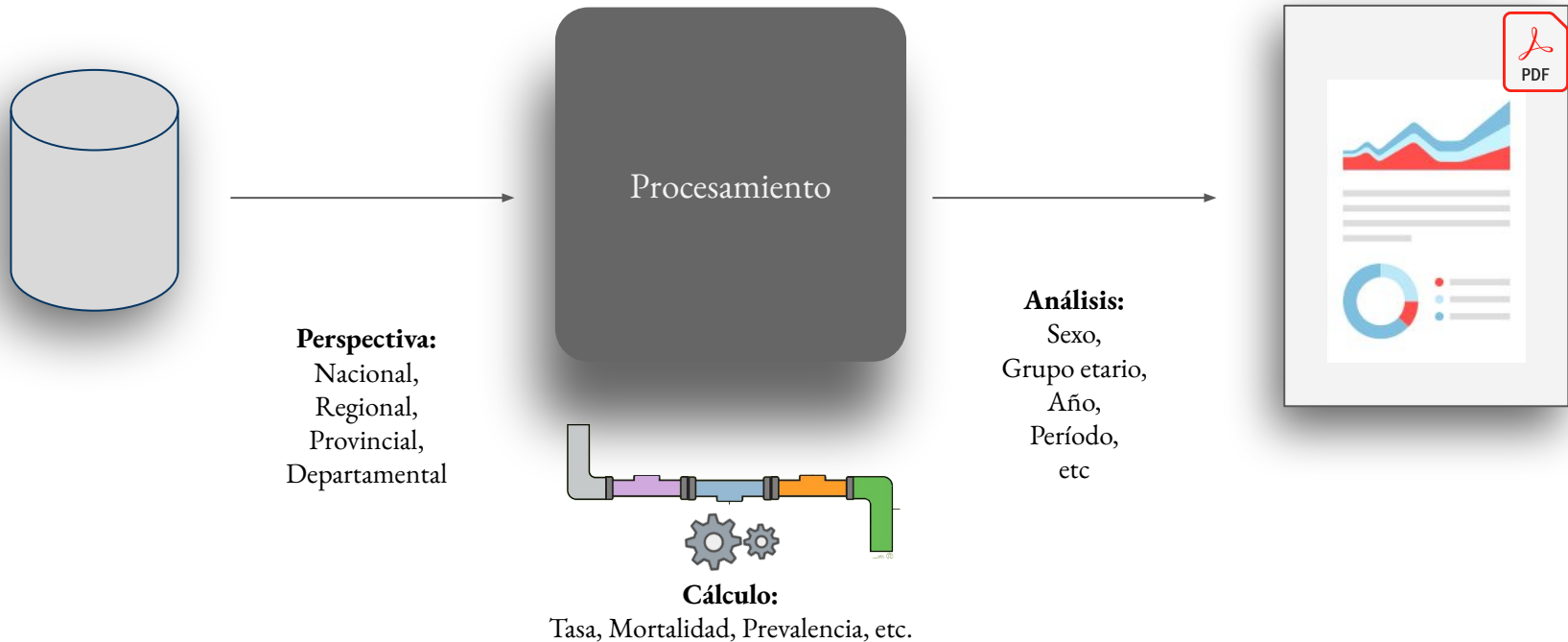
Pipeline, entradas, procesamiento y salidas



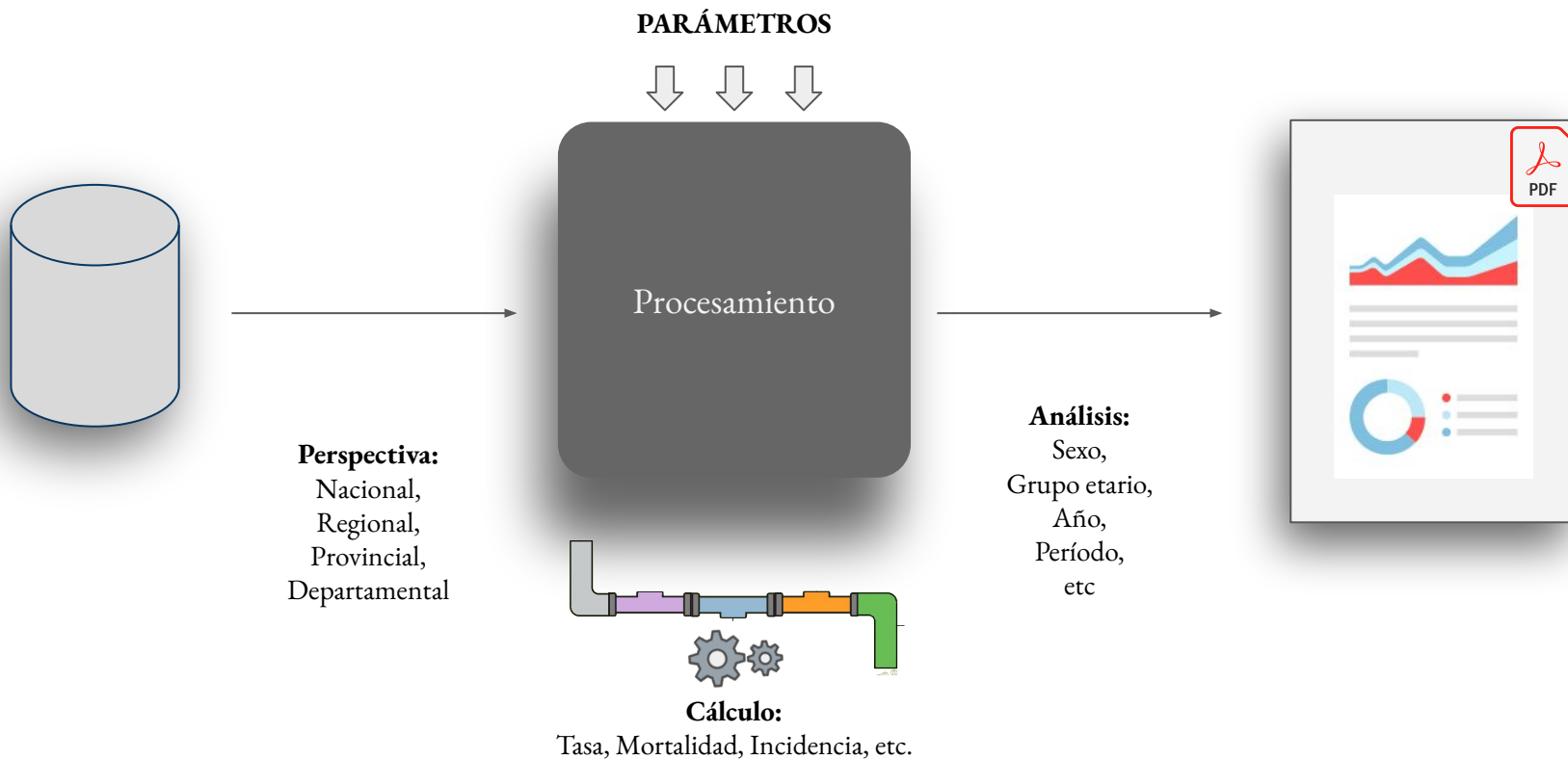
Pipeline, entradas, procesamiento y salidas



Pipeline, entradas, procesamiento y salidas

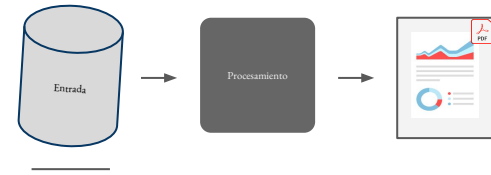


Pipeline, entradas, procesamiento y salidas

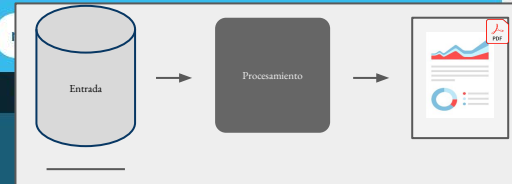


Pipeline, entradas, procesamiento y salidas





Información de entrada



Dirección de Estadísticas e Información de la Salud

Producimos, difundimos y analizamos estadísticas relacionadas con condiciones de vida y problemas de salud.



Indicadores básicos



Publicaciones



Datos



Guías COVID-19

Enfermedad por COVID-19: Guía para la certificación médica de las causas de muerte

[Descargar](#)

Enfermedad por COVID-19: Guía para la codificación de las causas de muerte

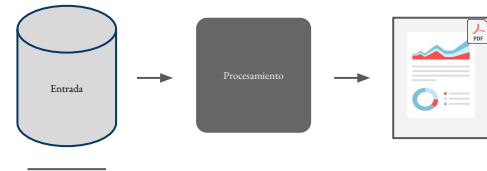
[Descargar](#)



¿Cómo podemos ayudarte?



Información de entrada



Entrada principal

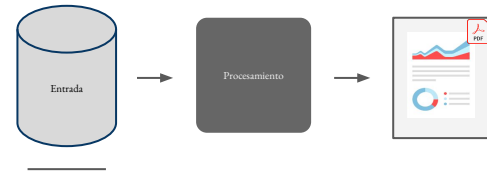
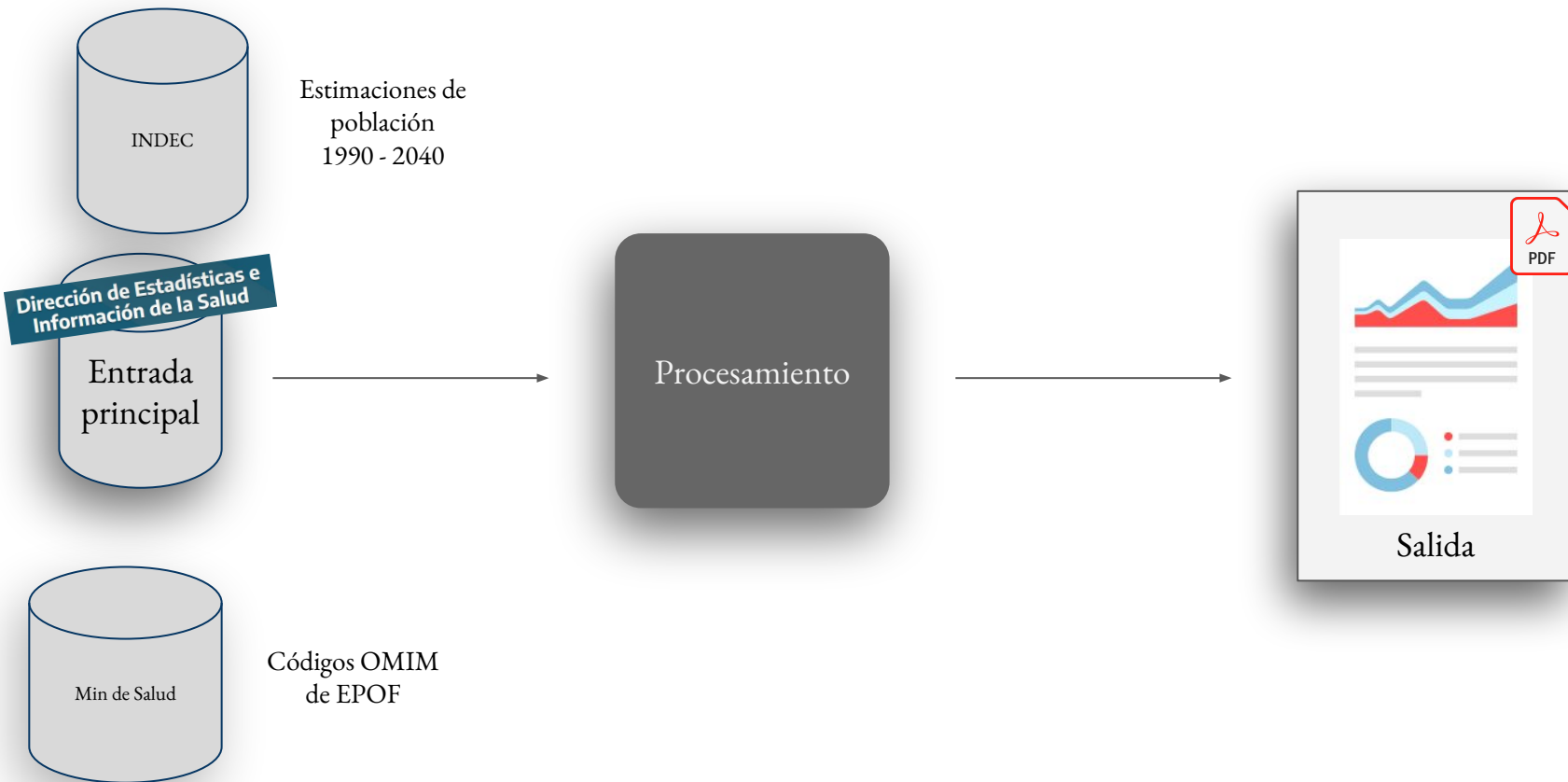


Procesamiento

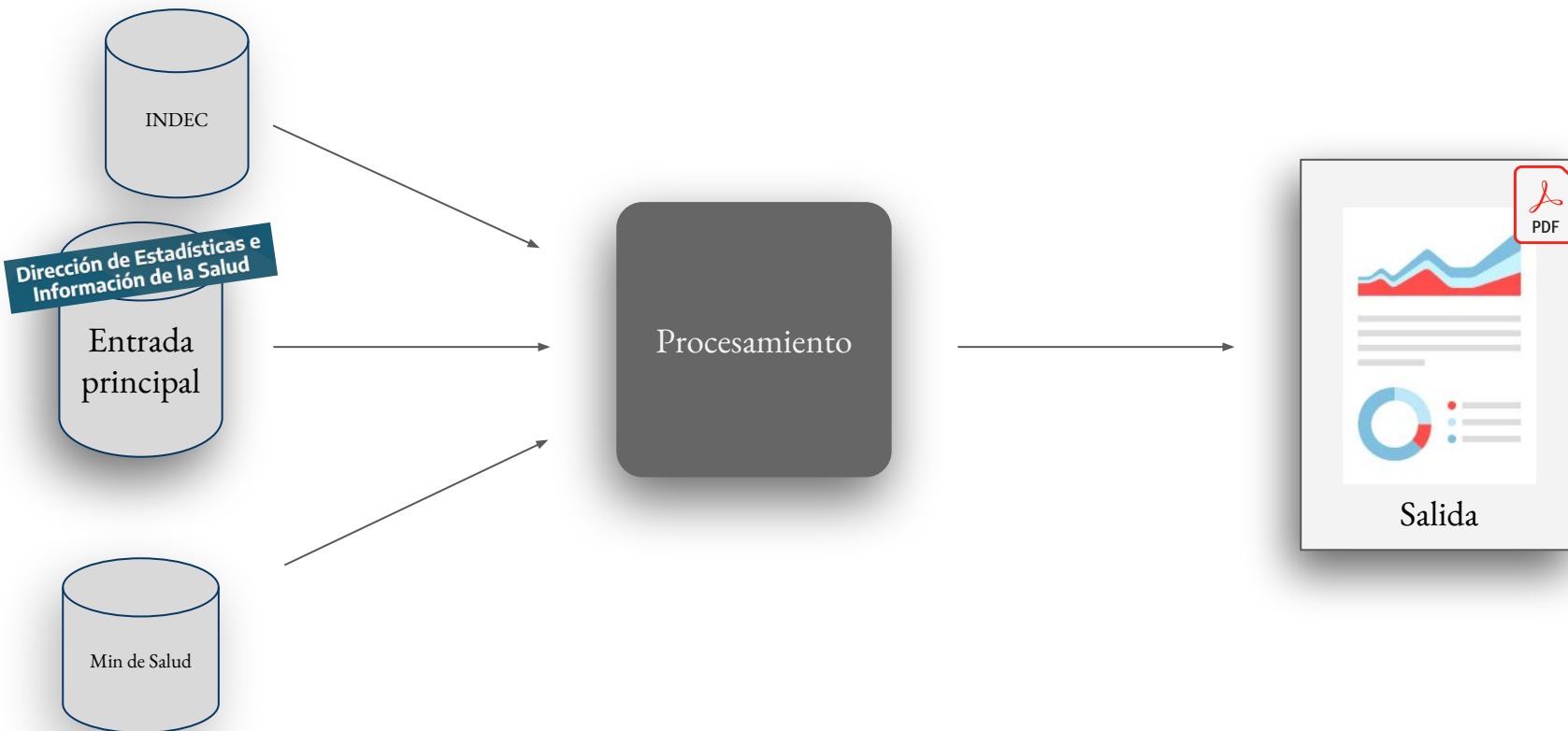
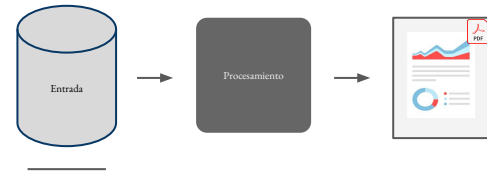


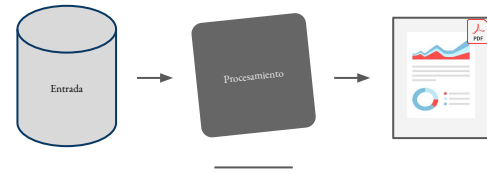
Salida

Información de entrada



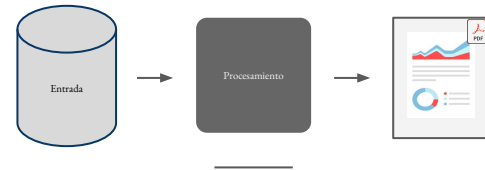
Información de entrada





Procesamiento

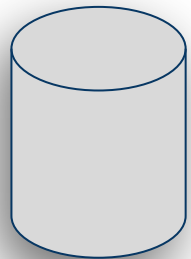
Procesamiento



Tasa de Mortalidad por Causas Específicas (CSMR): Número de fallecimientos por una causa específica / Muertes totales en una población específica, en un tiempo específico.

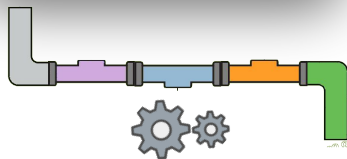
CSMR detalladas: Agrupando, filtrando por grupo etario, sexo, categoría de código de deceso.

Prevalencia: Número de fallecimientos por una causa específica / Población específica, en un punto específico del tiempo.



Perspectiva:
Nacional,
Regional,
Provincial,
Departamental

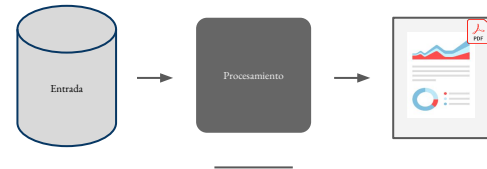
PARÁMETROS



Cálculo:
Tasa, Mortalidad, Incidencia, etc.



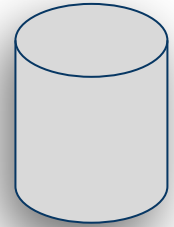
Análisis:
Sexo,
Grupo etario,
Año,
Período,
etc



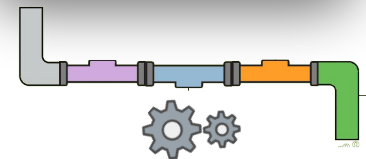
Conjunto de Códigos CIE-10

Configuración de grupos etarios

PARÁMETROS



Perspectiva:
Nacional,
Regional,
Provincial,
Departamental

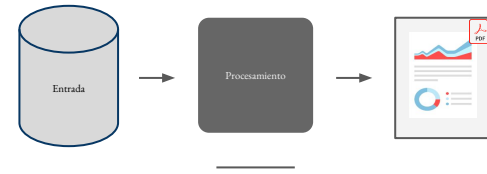


Cálculo:
Tasa, Mortalidad, Incidencia, etc.


Análisis:
Sexo,
Grupo etario,
Año,
Período,
etc



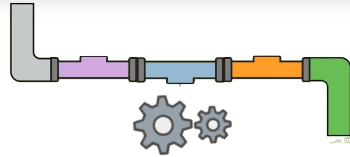
Procesamiento

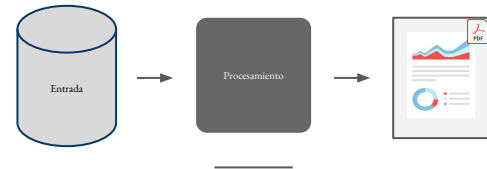


df.head()



	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN





	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN

	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERECIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN



df_regional.head()

	region_nombre	año	fallecimientos
0	Region A	1994	4603
1	Region A	1995	4365
2	Region A	1996	4425
3	Region A	1997	4114
4	Region A	1998	4146

df_provincial.tail()

	provincia_id	año	fallecimientos
667	10	2015	35
668	10	2016	19
669	10	2017	21
670	10	2018	226
671	10	2019	233

df_departamental.head()

	departamento_id	año	fallecimientos
0	99123	2006	3
1	99123	2007	4
2	99123	2010	2
3	88456	1997	6
4	88456	1998	6

	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN



```
df_regional.head()
```

	region_nombre	año	fallecimientos
0	Region A	1994	4603
1	Region A	1995	4365
2	Region A	1996	4425
3	Region A	1997	4207
4	Region A	1998	4118



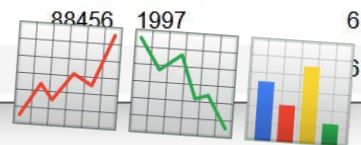
```
df_provincial.tail()
```

	provincia_id	año	fallecimientos
667	10	2015	35
668	10	2016	19
669	10	2017	21
670	10	2018	226
671	10	2019	233



```
df_departamental.head()
```

	departamento_id	año	fallecimientos
0	99123	2006	3
1	99123	2007	4
2	99123	2010	2
3	88456	1997	6
4	88456	1998	3



	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN



```
df_regional.head()
```

	region_nombre	año	fallecimientos
0	Region A	1994	4603
1	Region A	1995	4365
2	Region A	1996	4425
3	Region A	1997	4707
4	Region A	1998	4818



```
df_provincial.tail()
```

	provincia_id	año	fallecimientos
667	10	2015	35
668	10	2016	19
669	10	2017	21
670	10	2018	226
671	10	2019	233

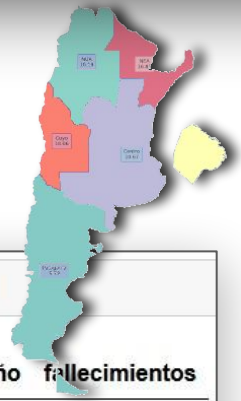


```
df_departamental.head()
```

	departamento_id	año	fallecimientos
0	99123	2006	3
1	99123	2007	4
2	99123	2010	2
3	88456	1997	6
4			3



	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN



```
df_regional.head()
```

	region_nombre	año	fallecimientos
0	Region A	1994	4603
1	Region A	1995	4365
2	Region A	1996	4425
3	Region A	1997	4378
4	Region A	1998	4318



```
df_provincial.tail()
```

provincia_id	año	fallecimientos
667	10 2014	35
668	10 2016	19
669	10 2017	21
670	10 2018	226
671	10 2019	233

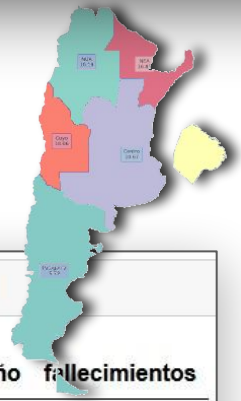


```
df_departamental.head()
```

	departamento_id	año	fallecimientos
0	99123	2006	3
1	99123	2007	4
2	99123	2010	2
3	88456	1997	6
4	88456	1998	5



	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN



```
df_regional.head()
```

	region_nombre	año	fallecimientos
0	Region A	1994	4603
1	Region A	1995	4365
2	Region A	1996	4425
3	Region A	1997	4378
4	Region A	1998	4318



```
df_provincial.tail()
```

	provincia_id	año	fallecimientos
667	10	2017	35
668	10	2016	19
669	10	2017	21
670	10	2018	226
671	10	2019	233

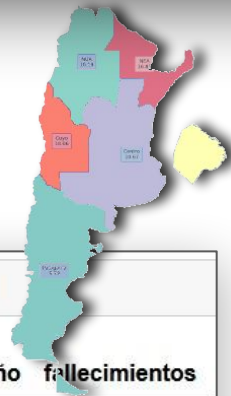


```
df_departamental.head()
```

	departamento_id	año	fallecimientos
0	99123	2006	3
1	99123	2007	4
2	99123	2010	2
3	99123	2011	6
4	99123	1997	3



	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN



```
df_regional.head()
```

	region_nombre	año	fallecimientos
0	Region A	1994	4603
1	Region A	1995	4365
2	Region A	1996	4425
3	Region A	1997	4207
4	Region A	1998	4178

```
df_provincial.tail()
```

	provincia_id	año	fallecimientos
667	10	2015	35
668	10	2016	19
669	10	2017	21
670	10	2018	226
671	10	2019	233

```
df_departamental.head()
```

	departamento_id	año	fallecimientos
0	99123	2006	3
1	99123	2007	4
2	99123	2010	2
3	99123	1997	6
4	99123	1998	5

	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN



df_provincial.tail()

provincia_id	año	fallecimientos
667	10 2016	35
668	10 2016	19
669	10 2017	21
670	10 2018	226
671	10 2019	233

df_regional.head()

region_nombre	año	fallecimientos
0	Region A 1994	4603
1	Region A 1995	4365
2	Region A 1996	4425
3	Region A 1997	4114
4	Region A 1998	4146

df_departamental.head()

departamento_id	año	fallecimientos
0	99123 2006	3
1	99123 2007	4
2	99123 2010	2
3	88456 1997	6
4		3



	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN



df_regional.head()

provincia_id	nombre	año	fallecimientos
0	1999 - 2003	1994	4603
1	Region A	1995	4365
	Region A	1996	4425
	Region A	1997	4114
4	Region A	1998	4146

1994 - 1998

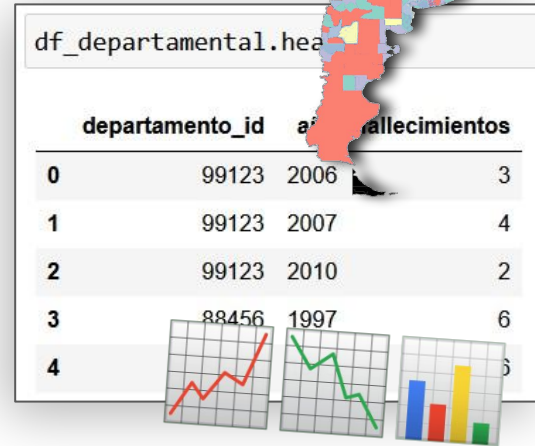
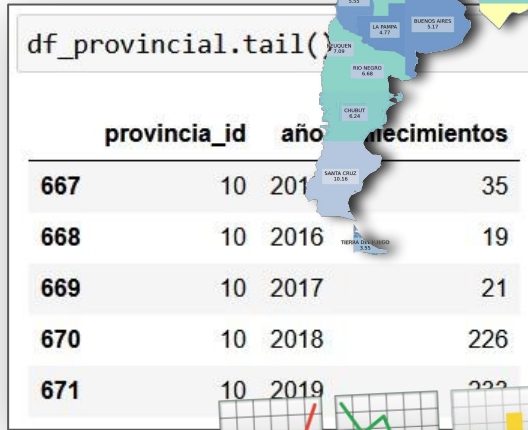
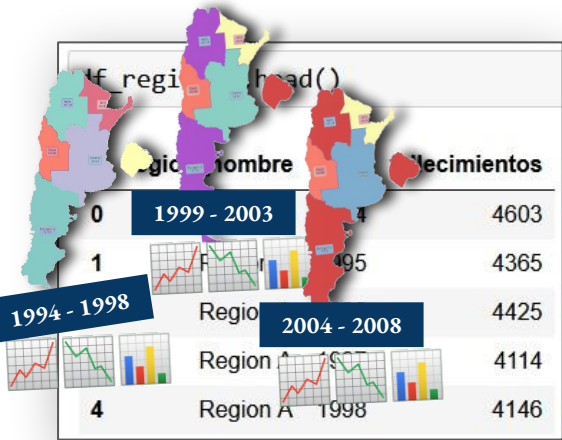
df_provincial.tail()

provincia_id	año	fallecimientos
667	10 201	35
668	10 2016	19
669	10 2017	21
670	10 2018	226
671	10 2019	233

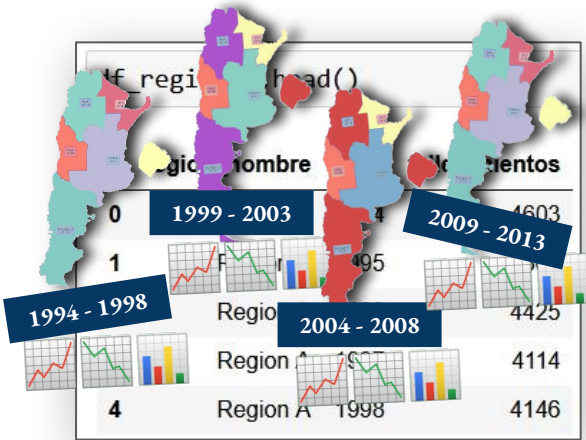
df_departamental.head()

departamento_id	año	fallecimientos
0	99123 2006	3
1	99123 2007	4
2	99123 2010	2
3	88456 1997	6
4		3

	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN



	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN



df_provincial.tail()

provincia_id	año	decesos
667	10 2016	35
668	10 2016	19
669	10 2017	21
670	10 2018	226
671	10 2019	222

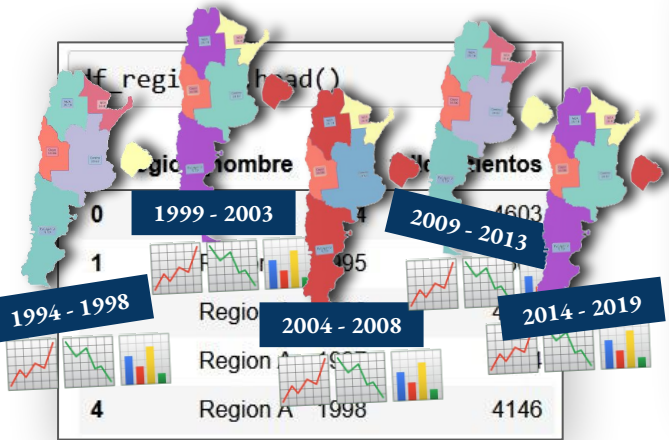


df_departamental.head()

departamento_id	año	decesos
0	99123 2006	3
1	99123 2007	4
2	99123 2010	2
3	88456 1997	6
4		3

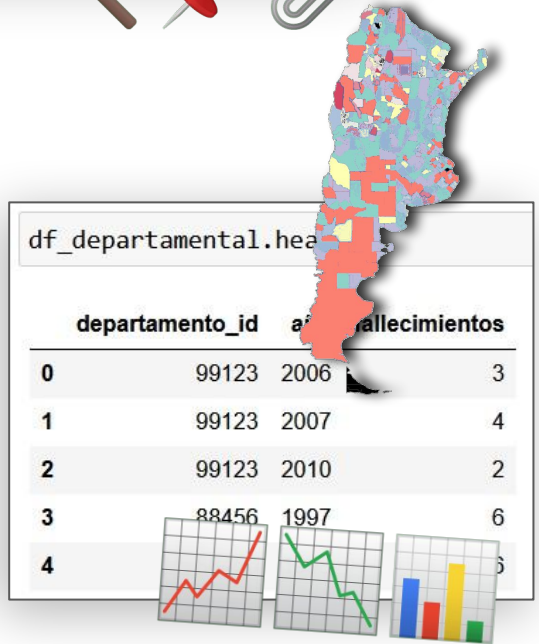


	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN

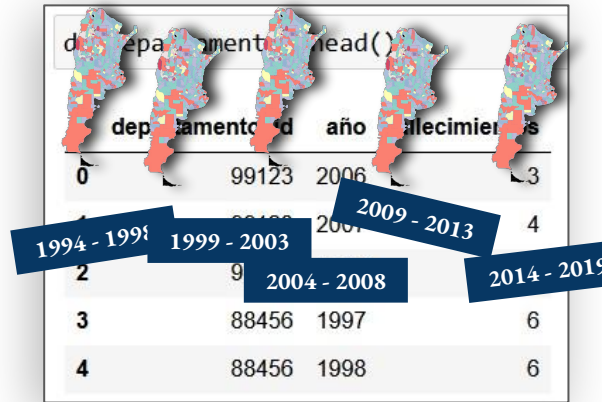
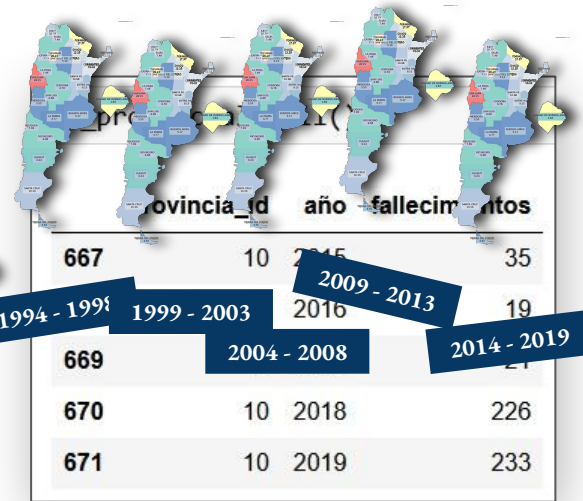
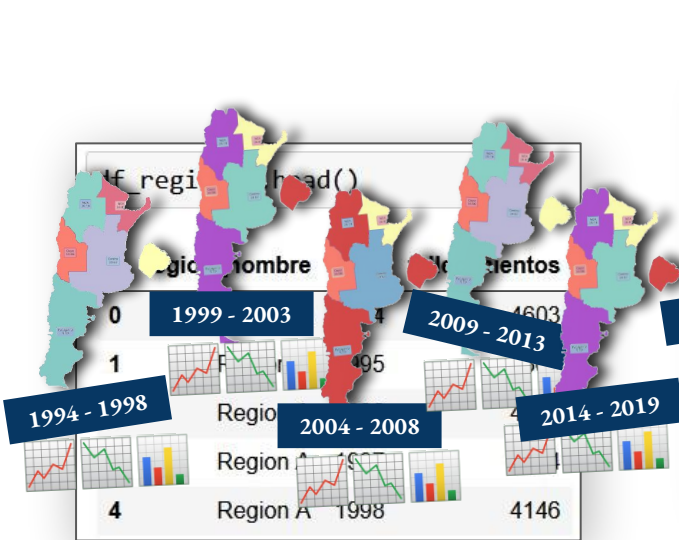


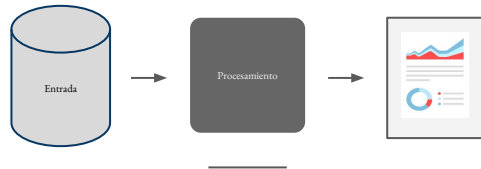
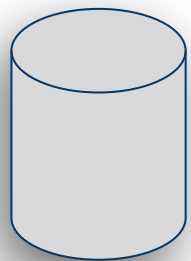
df_provincial.tail()

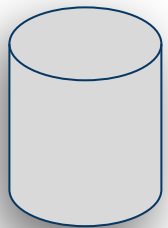
provincia_id	año	decesos	necimientos
667	10	201	35
668	10	2016	19
669	10	2017	21
670	10	2018	226
671	10	2019	233



	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
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3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN



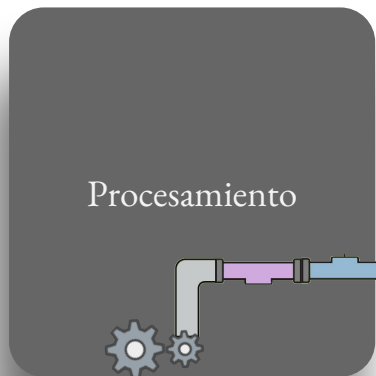




Conjunto
de Códigos
CIE-10

Configuración
de grupos
etarios

PARÁMETROS



	region_nombre	año	fallecimientos
0	Region A	1994	4603
1	Region A	1995	4365
2	Region A	1996	4425
3	Region A	1997	4114
4	Region A	1998	4146

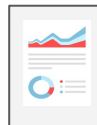
provincia_id	año	fallecimientos
667	10 2015	35
668	10 2016	
669	10 2017	
670	10 2018	
671	10 2019	

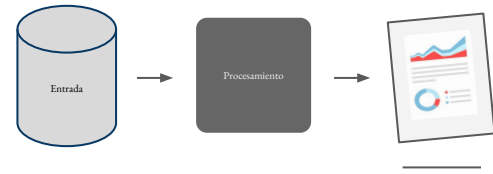


departamento_id	año	fallecimientos
0	99123 2006	3
1	99123 2007	4
2	99123 2010	2
3	88456 1997	6
4	88456 1998	6

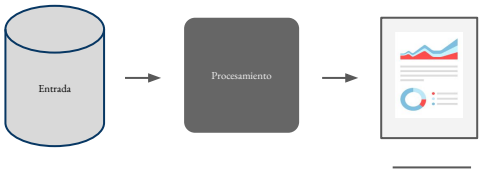
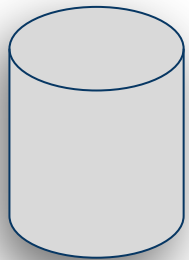


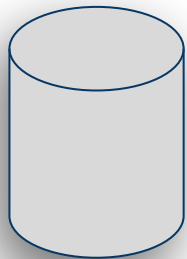
Procesamiento



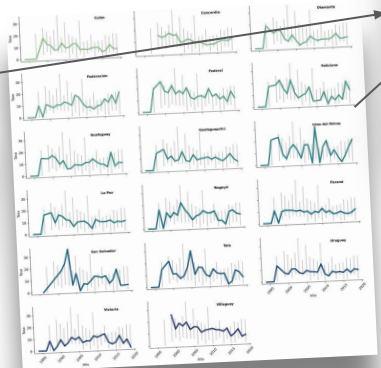
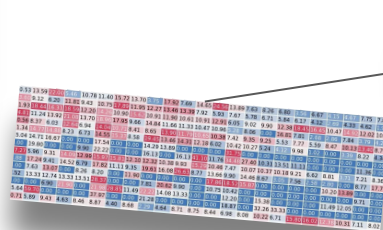
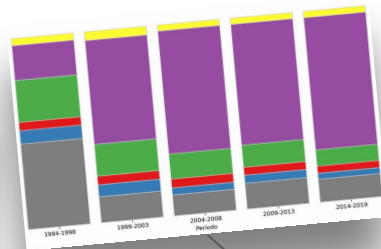
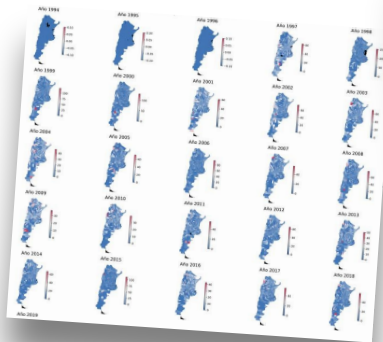


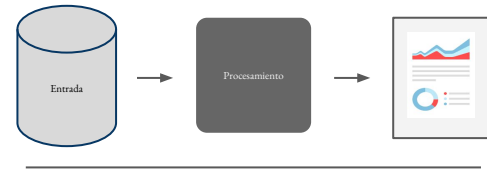
Salida



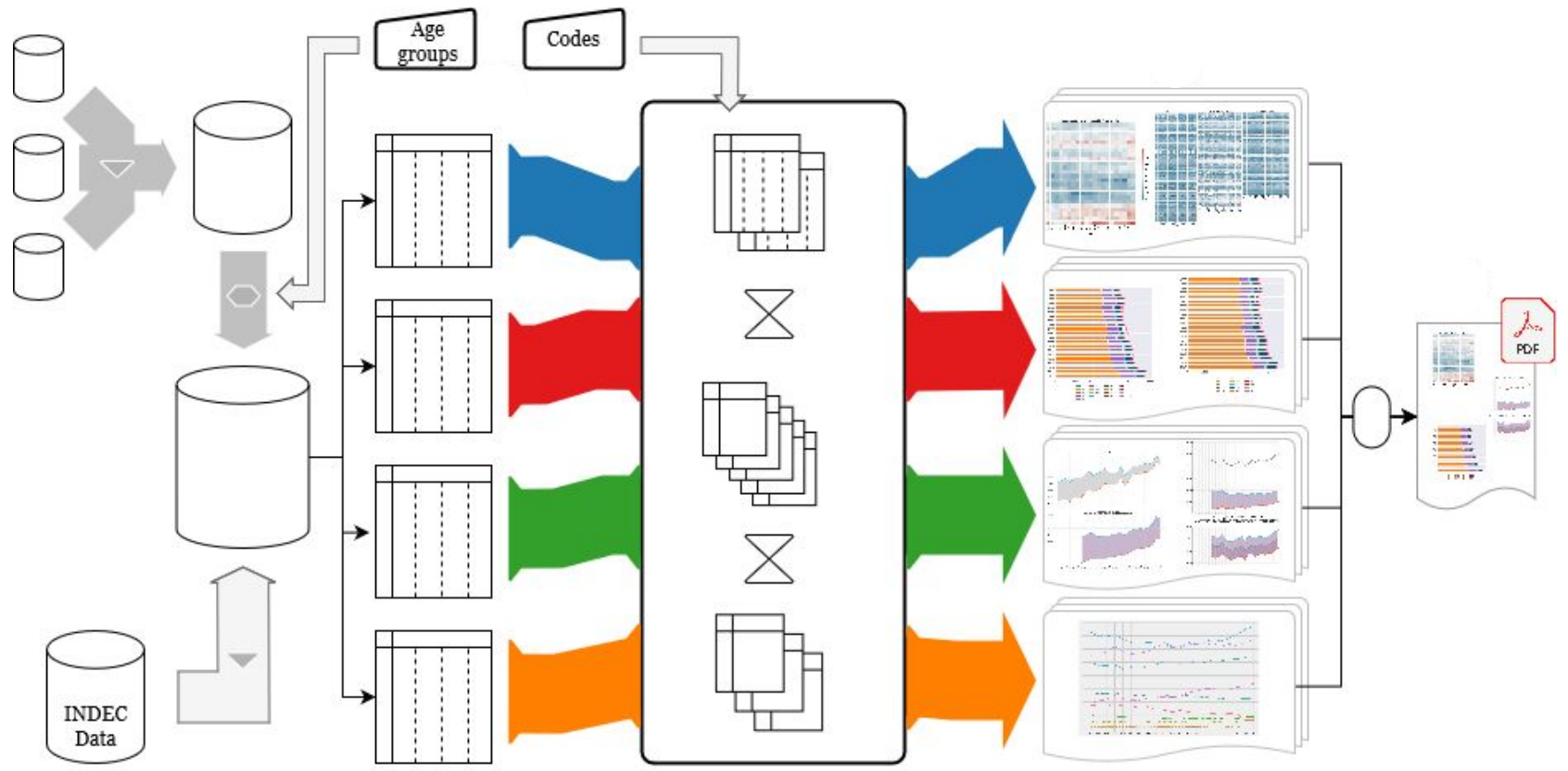
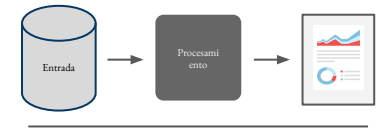


Procesamiento





Pipeline



Implementación

The Jupyter logo features a stylized orange smiley face with two curved lines for eyes and a wide, upward-curving mouth. Four grey circles are positioned at the corners of a square around the smiley face. The word "jupyter" is written in a dark grey, lowercase, sans-serif font across the center of the smiley face's face.

jupyter


```
File Edit View Insert Cell Help
In [147]:
# Copiar
# Cargar las bases que corresponden a nivel regional. Crear 5 mesas (1 por cada quinquenio). Dejar variable para poder cambiar las columnas
# Es necesario entonces tener con dos bases de datos:
# - Bases de datos (valores)
# - Bases de datos (valores)
# Crear un dataset general con granularidad el departamento, el de provincia, nombre de region + recuentos + nuevas mesas.
# Al dataset anterior se debe agregar una columna TASA, que se calcula según la siguiente fórmula:
# (nueve total / nacidos vivos) * 10000

In [147]: import pandas
import numpy
import sys
import pandas
import matplotlib.pyplot as plt
import seaborn
import time

In [148]: ONE_PREFIX = 'YPI'
TWO_PREFIX = 'YPI'

Carga
Caso del mapa de Argentina. URL para obtener la información de departamentos, provincia y region estandarizada.

In [149]: from functools import partial
def get_departments_info(provincia, nombre_archivo):
    url = 'https://datos.bancomundial.org/indicadores/Argentina/gestion'
    response = requests.get(url)
    data = response.json()
    return data

registros = [
    {'provincia': 'Buenos Aires',
     'nombre': 'Buenos Aires',
     'lat': 34.6037,
     'lon': -58.3816},
    {'provincia': 'Córdoba',
     'nombre': 'Córdoba',
     'lat': 31.3772,
     'lon': -64.2445},
    {'provincia': 'Entre Ríos',
     'nombre': 'Entre Ríos',
     'lat': 32.7350,
     'lon': -58.1180},
    {'provincia': 'Mendoza',
     'nombre': 'Mendoza',
     'lat': 32.8895,
     'lon': -68.8110},
    {'provincia': 'Salta',
     'nombre': 'Salta',
     'lat': 26.7825,
     'lon': -65.1760},
    {'provincia': 'Santiago del Estero',
     'nombre': 'Santiago del Estero',
     'lat': 27.6540,
     'lon': -64.2445},
    {'provincia': 'Tucumán',
     'nombre': 'Tucumán',
     'lat': 26.8083,
     'lon': -65.2147},
    {'provincia': 'Chaco',
     'nombre': 'Chaco',
     'lat': 27.3548,
     'lon': -58.9972},
    {'provincia': 'Formosa',
     'nombre': 'Formosa',
     'lat': 26.3103,
     'lon': -58.1750},
    {'provincia': 'Misiones',
     'nombre': 'Misiones',
     'lat': 27.3548,
     'lon': -54.3850},
    {'provincia': 'Paraná',
     'nombre': 'Paraná',
     'lat': 31.7333,
     'lon': -55.8333},
    {'provincia': 'Rosario',
     'nombre': 'Rosario',
     'lat': 32.9697,
     'lon': -58.7822},
    {'provincia': 'San Luis',
     'nombre': 'San Luis',
     'lat': 33.2375,
     'lon': -64.2445},
    {'provincia': 'Santa Fe',
     'nombre': 'Santa Fe',
     'lat': 31.5375,
     'lon': -59.2575},
    {'provincia': 'Tierra del Fuego',
     'nombre': 'Tierra del Fuego',
     'lat': 54.8071,
     'lon': -72.2683},
    {'provincia': 'Uruguay',
     'nombre': 'Uruguay',
     'lat': 34.8778,
     'lon': -56.1635},
    {'provincia': 'Venezuela',
     'nombre': 'Venezuela',
     'lat': 6.2808,
     'lon': -54.6876},
    {'provincia': 'Argentina',
     'nombre': 'Argentina',
     'lat': 34.0,
     'lon': -54.0}
]

In [150]: get_nacidos_vivos = pandas.read_csv(
    'https://datos.bancomundial.org/indicadores/Argentina/gestion/nacidos_vivos_1994-2003.csv',
    usecols=(PROVINCIAS, DEPARTAMENTOS, CUESTA), skiprows=1, skipcols=1)

In [151]: get_nacidos_vivos.head()

Out[151]:
   PROVINCIA  DEPARTAMENTO  CUESTA
0  1994      2           1  100
1  1994      2           2  150
2  1994      2           3  80000
3  1994      2           4  20000
4  1994      2           5  1950

In [152]: get_nacidos_vivos[CUESTA] = get_nacidos_vivos[CUESTA].astype('int')

In [153]: get_nacidos_vivos.head()

Out[153]:
   PROVINCIA  DEPARTAMENTO  CUESTA
0  1994      2           1      1
1  1994      2           2      1
2  1994      2           3      801
3  1994      2           4      993
4  1994      2           5      1

In [154]: def clean_department_code(code):
    pad = 3 - len(str(code))
    return str(code).zfill(pad)

# Función para limpiar el código de provincia
def clean_province_code(code):
    return str(code) if len(str(code)) == 1 else str(code)

print(clean_department_code(1))
print(clean_department_code(1))
print(clean_province_code(1))
print(clean_province_code(1))

4 801
41 801
413 801
```



PLOOMBER



On writing clean Jupyter notebooks

10 recommendations for writing readable and maintainable notebooks

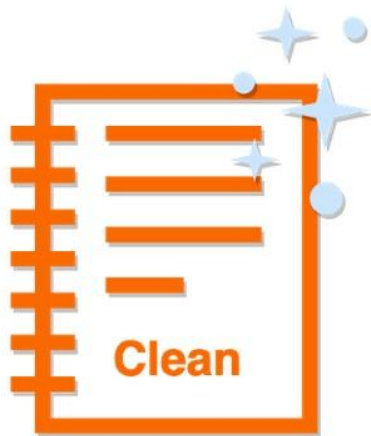


Image by author

Notebooks are a magnificent tool to explore data, but such a powerful tool can become hard to manage quickly. Ironically, the ability to interact with our data rapidly (modify code cells, run, and repeat) is the exact reason why a notebook may become an obscure entanglement of variables that are hard to understand, even to the notebook's author. But it doesn't have to be that way. This post summarizes my learnings over the past few years on writing clean notebooks.

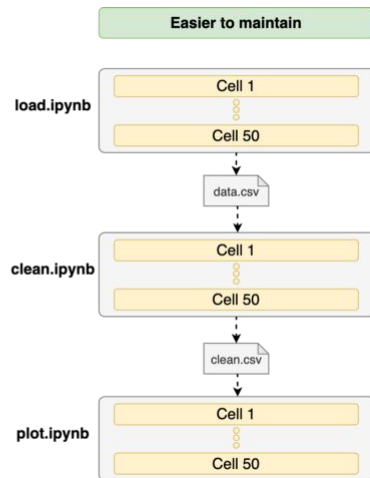
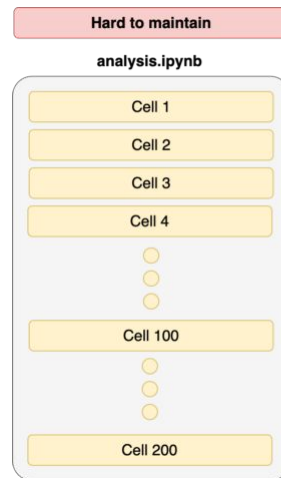
On writing clean Jupyter notebooks

10 recommendations for writing readable and maintainable notebooks



Image by author

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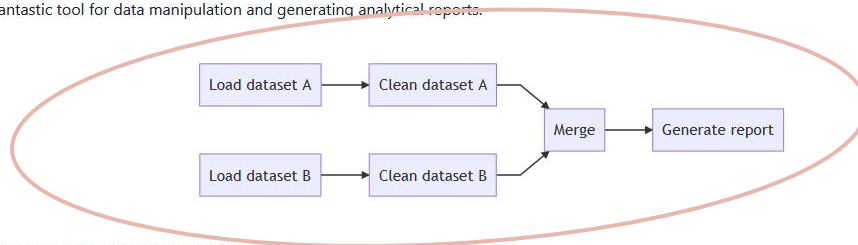
Get Started

Use Cases

[Machine Learning](#)[Research Projects](#)[Analytics](#)[SQL Pipelines](#)[User Guide](#)[Cloud](#)[Deployment](#)[Cookbook](#)[API Reference](#)[Community](#)

Analytics

Ploomber is a fantastic tool for data manipulation and generating analytical reports.



- ▶ Example: BigQuery and Cloud Storage pipeline
- ▼ Example: Exploratory data analysis pipeline

```
Terminal (shell) Click to copy  
$ pip install ploomber  
$ ploomber examples -n templates/exploratory-analysis -o exploratory-analysis
```

Modularize your project

Instead of coding everything in a single notebook (which is difficult to maintain and collaborate), you can quickly break down your analysis into multiple parts.

Faster iterations

Finding data insights is an iterative process, with Ploomber's [incremental builds](#) you can rapidly iterate on your data since the framework skips redundant computations and only executes tasks whose source code has changed since the last execution.

Automated report generation

Once your pipeline is ready, you can easily create HTML reports from your scripts/notebooks. Just change the extension of the task, and Ploomber will automatically convert the output for you.

Analytics

[Modularize your project](#)[Faster iterations](#)[Automated report generation](#)

```
→ exploratory-analysis tree
```

```
├── environment.yml  
├── pipeline.yaml  
├── README.ipynb  
├── README.md  
├── requirements.txt  
├── scripts  
│   ├── clean.py  
│   ├── custom.py  
│   ├── get.py  
│   ├── profile-clean.py  
│   └── profile-raw.py  
└── _source.md
```

```
1 directory, 11 files
```

```
→ exploratory-analysis tree
```

```
.  
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│   ├── profile-clean.py  
│   └── profile-raw.py  
└── _source.md
```

```
1 directory, 11 files
```

```
tasks:
  # get raw data
  - source: scripts/get.py
    product:
      nb: products/get.html
      data: products/raw.csv

  # clean raw data
  - source: scripts/clean.py
    product:
      nb: products/clean.html
      # clean data
      data: products/clean.csv

  # quick clean data profiling
  - source: scripts/profile-clean.py
    # html report
    product: products/report-clean.html

  # custom plots
  - source: scripts/custom.py
    product: products/custom.html
```

```
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```

tasks:
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    product:
      nb: products/get.html
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  # clean raw data
  - source: scripts/clean.py
    product:
      nb: products/clean.html
      # clean data
      data: products/clean.csv

  # quick clean data profiling
  - source: scripts/profile-clean.py
    # html report
    product: products/report-clean.html

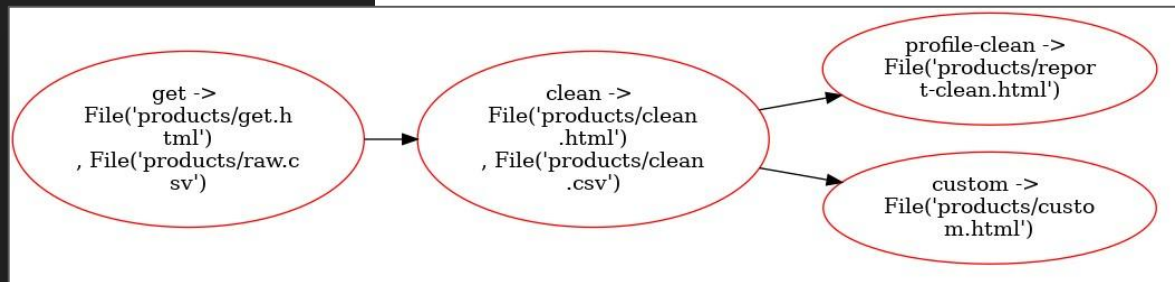
  # custom plots
  - source: scripts/custom.py
    product: products/custom.html

```

```

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1 directory, 11 files

```



```
# ---
# jupyter:
#   jupyter:
#     text_representation:
#       extension: .py
#       format_name: percent
#       format_version: '1.3'
#       jupyter_text_version: 1.13.6
#     kernelspec:
#       display_name: Python 3 (ipykernel)
#       language: python
#       name: python3
# ---
```

```
# %% [markdown]
# Get penguins data.
```

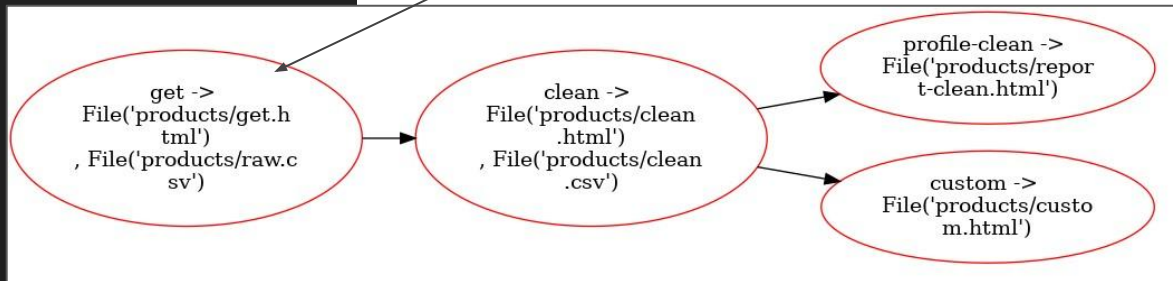
```
# %%
import seaborn as sns
```

```
# %% tags=["parameters"]
upstream = None
product = None
```

```
# %%
df = sns.load_dataset('penguins')
```

```
# %%
df.to_csv(product['data'], index=False)
```

```
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└── get.py
    ├── profile-raw.py
    └── _source.md
1 directory, 11 files
```



```
# %% [markdown]
# Clean penguins data.

# %%
import pandas as pd

# %% tags=["parameters"]
upstream = ['get']
product = None

# %%
df = pd.read_csv(upstream['get']['data'])

# %%
print(f'Raw data has {len(df)} rows...')

# %%
df.head()

# %%
# really simple cleaning, remove if the row has nas...
clean = df[df.isna().sum(axis=1) == 0]

# %%
print(f'Clean data has {len(clean)} rows...')

# %%
clean.to_csv(product['data'], index=False)
```

```
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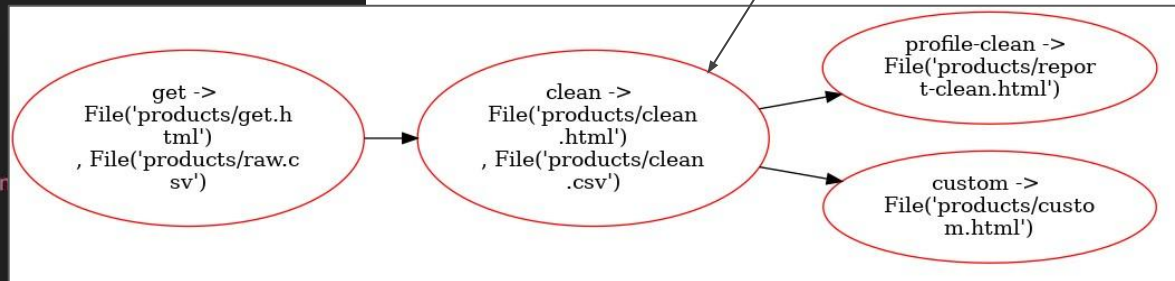
# %%
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# %%
# really simple cleaning, remove if the row has r
clean = df[df.isna().sum(axis=1) == 0]

# %%
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# %%
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```

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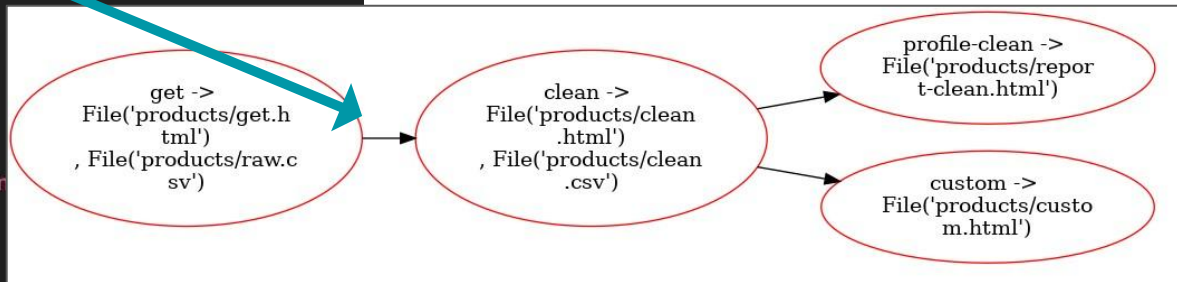
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# %%
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# %%
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# %%
clean.to_csv(product['data'], index=False)
```

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```



```
# %% [markdown]
# Quick clean data profiling.

# %%
import pandas as pd
from pandas_profiling import ProfileReport

# %% tags=["parameters"]
upstream = ['clean']
product = None

# %%
df = pd.read_csv(upstream['clean']['data'])

# %%
ProfileReport(df, title="Clean Data Profiling Report")

# %%
```

```
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├── scripts
│   ├── clean.py
│   ├── custom.py
│   └── get.py
├── profile-clean.py
├── profile-raw.py
└── _source.md

1 directory, 11 files
```

```

# %% [markdown]
# Quick clean data profiling.

# %%
import pandas as pd
from pandas_profiling import ProfileReport

# %% tags=["parameters"]
upstream = ['clean']
product = None

# %%
df = pd.read_csv(upstream['clean']['data'])

# %%
ProfileReport(df, title="Clean Data Profiling Report")

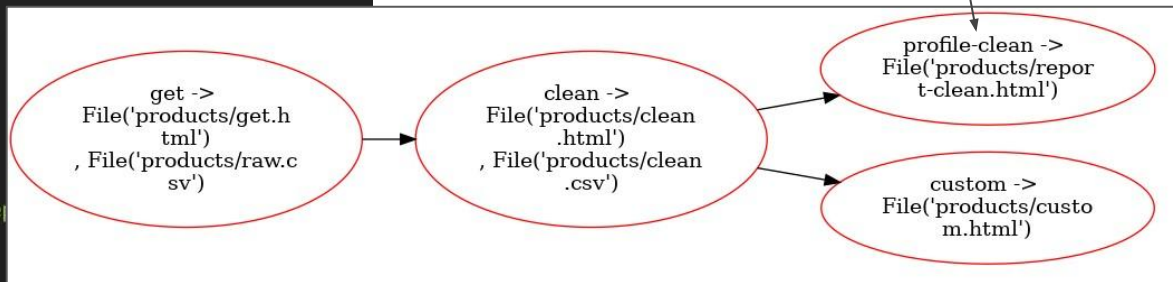
# %%

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exploratory-analysis tree
├── environment.yml
├── pipeline.yaml
├── README.ipynb
├── README.md
├── requirements.txt
├── scripts
│   ├── clean.py
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├── profile-raw.py
├── _source.md
└── 1 directory, 11 files

```



```
# %% [markdown]
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from pandas_profiling import ProfileReport

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product = None
```

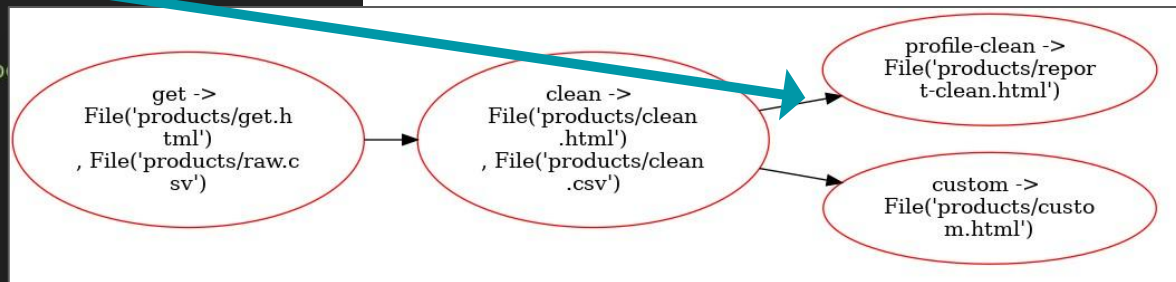
```
# %%
df = pd.read_csv(upstream['clean']['data'])

# %%
ProfileReport(df, title="Clean Data Profiling Report")

# %%
```

```
exploratory-analysis tree
├── environment.yml
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├── README.md
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├── profile-raw.py
└── _source.md

1 directory, 11 files
```



```
# %% [markdown]
# Create custom plots.

# %%
import seaborn as sns
import pandas as pd
import plotly.express as px
import matplotlib.pyplot as plt
import matplotlib

# %%
# prettier plots
plt.style.use('ggplot')
# larger plots
matplotlib.rc('figure', figsize=(15, 10))
# larger fonts
sns.set_context('notebook', font_scale=1.5)

# %% tags=["parameters"]
upstream = ['clean']
product = None

# %%
df = pd.read_csv(upstream['clean']['data'])

# %%
df.head()

# %%
_ = df['species'].value_counts().plot(kind='bar')

# %%
fig = px.histogram(df, x="bill_length_mm")
fig.show()

# %%
sns.pairplot(df, hue="species", height=3,diag_kind="hist")

# %%
```

```
→ exploratory-analysis tree
├── environment.yml
├── pipeline.yaml
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├── README.md
├── requirements.txt
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1 directory, 11 files
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```

# %% [markdown]
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# %%
sns.pairplot(df, hue="species", height=3,diag_kind="hist")

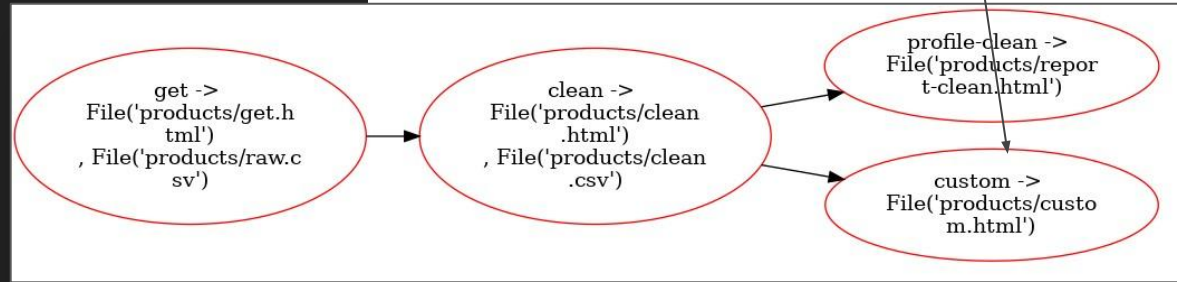
# %%

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```

exploratory-analysis tree
├── environment.yml
├── pipeline.yaml
├── README.ipynb
├── README.md
├── requirements.txt
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└── 1 directory, 11 files

```




```

# %% [markdown]
# Create custom plots.

# %%
import seaborn as sns
import pandas as pd
import plotly.express as px
import matplotlib.pyplot as plt
import matplotlib

# %%
# prettier plots
plt.style.use('ggplot')
# larger plots
matplotlib.rcParams['figure', figsize=(15, 10))
# larger fonts
sns.set_context('notebook', font_scale=1.5)

# %% tags=["parameters"]
upstream = ['clean']
product = None

# %%
df = pd.read_csv(upstream['clean']['data'])
# %%
df.head()

# %%
_ = df['species'].value_counts().plot(kind='bar')

# %%
fig = px.histogram(df, x="bill_length_mm")
fig.show()

# %%
sns.pairplot(df, hue="species", height=3, diag_kind="hist")

# %%

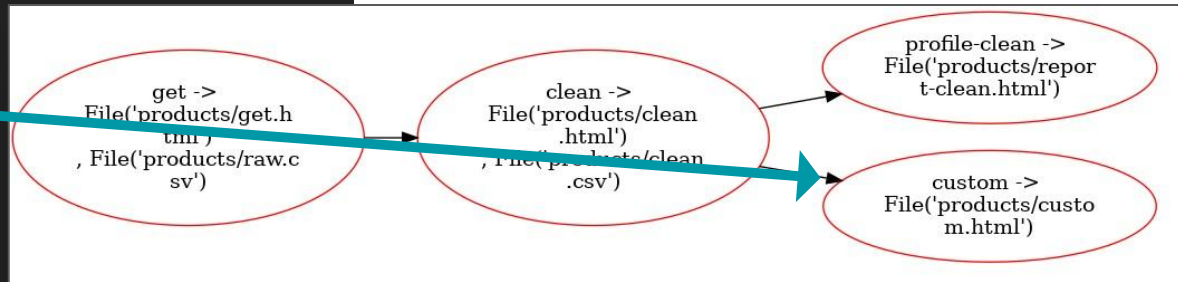
```

```

exploratory-analysis tree
├── environment.yml
├── pipeline.yaml
├── README.ipynb
├── README.md
├── requirements.txt
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└── _source.md

1 directory, 11 files

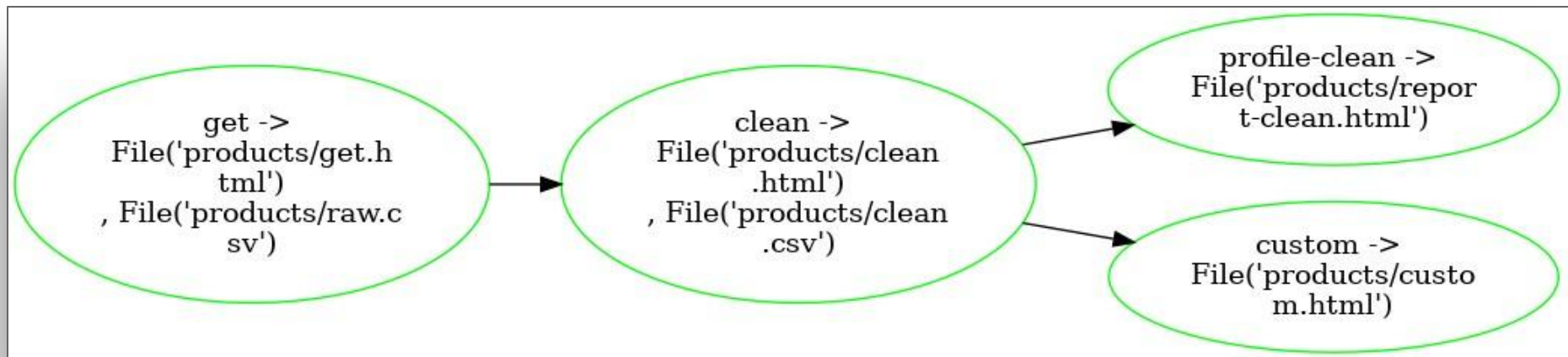
```




```
$ ploomber build
```

name	Ran?	Elapsed (s)	Percentage
get	True	1.80789	12.0805
clean	True	1.10755	7.40075
profile-clean	True	7.13636	47.6859
custom	True	4.91354	32.8328

name	Ran?	Elapsed (s)	Percentage
get	True	1.80789	12.0805
clean	True	1.10755	7.40075
profile-clean	True	7.13636	47.6859
custom	True	4.91354	32.8328



```
→ exploratory-analysis tree
```

```
*  
├── environment.yml  
├── pipeline.png  
├── pipeline.yaml  
├── products  
│   ├── clean.csv  
│   ├── clean.html  
│   ├── custom.html  
│   ├── get.html  
│   ├── raw.csv  
│   └── report-clean.html  
├── README.ipynb  
├── README.md  
├── requirements.txt  
├── scripts  
│   ├── clean.py  
│   ├── custom.py  
│   ├── get.py  
│   ├── profile-clean.py  
│   └── profile-raw.py  
└── _source.md
```

```
2 directories, 18 files
```

```

→ exploratory-analysis tree
├── environment.yml
├── pipeline.png
├── pipeline.yaml
├── products
│   ├── clean.csv
│   ├── clean.html
│   ├── custom.html
│   ├── get.html
│   └── raw.csv
├── report-clean.html
├── README.ipynb
├── README.md
├── requirements.txt
├── scripts
│   ├── clean.py
│   ├── custom.py
│   ├── get.py
│   ├── profile-clean.py
│   ├── profile-raw.py
│   └── _source.md
└── 2 directories, 18 files

```

Clean Data Profiling Report

Overview Variables Interactions Correlations Missing values Sample

Overview

Overview Alerts 0 Reproductions

Dataset statistics		Variable types	
Number of variables	7	Categorical	3
Number of observations	333	Numeric	4
Missing cells	0		
Missing cells (%)	0.0%		
Duplicate rows	0		
Duplicate rows (%)	0.0%		
Total size in memory	18.3 KB		
Average record size in memory	56.4 B		

Variables

species: Distinct 3, Missing 0, Memory size 2.7 KB

island: Distinct 3, Missing 0, Memory size 2.7 KB

bill_length_mm: Distinct 163, Minimum 52.1, Maximum 59.6

Interactions

Correlations

Species's p Pearson's r Kendall's tau Cramer's V (adj) Toggle correlation descriptions

Missing values

Count Matrix

A simple visualization of nullity by column.

→ exploratory-analysis tree

environment.yml

pipeline.png

pipeline.yaml

products

clean.csv

clean.html

custom.html

get.html

raw.csv

report-clean.html

README.ipynb

README.md

requirements.txt

scripts

clean.py

custom.py

get.py

profile-clean.py

profile-raw.py

_source.md

2 directories, 18 files

Code custom.py

```
1 import pandas as pd
2 import numpy as np
3 import matplotlib.pyplot as plt
4 import seaborn as sns
5 import sys
```

```
6 # read data
7 df = pd.read_csv('raw.csv')
```

```
8 # filter data
9 df = df[df['species'] != 'Chondro']
```

```
10 # clean data
11 df = df[df['bill_length_mm'] > 0]
```

```
12 # save data
13 df.to_csv('clean.csv', index=False)
```

```
14 # generate report
15 report = generate_report(df)
```

```
16 # save report
17 report.save('report-clean.html')
```

```
18 # exit
19 sys.exit(0)
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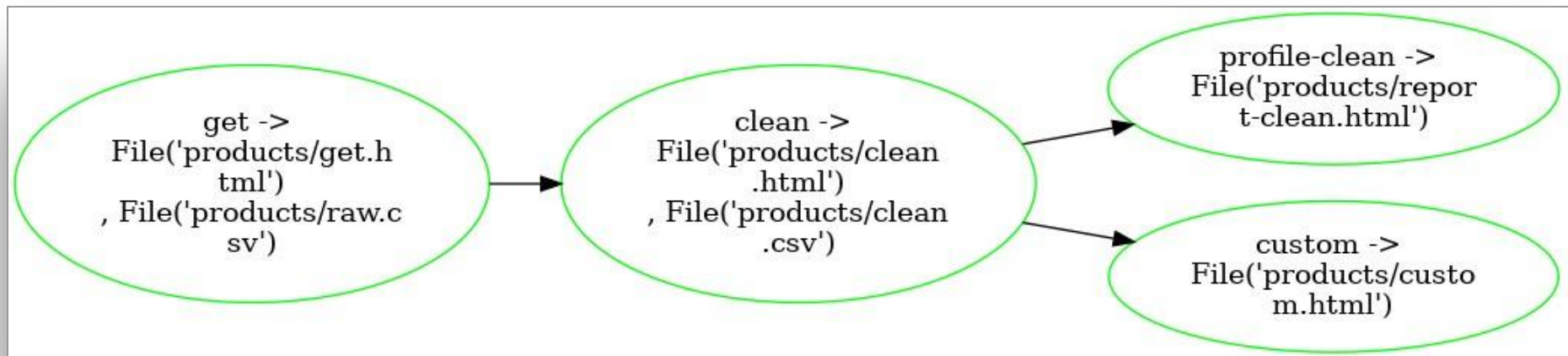
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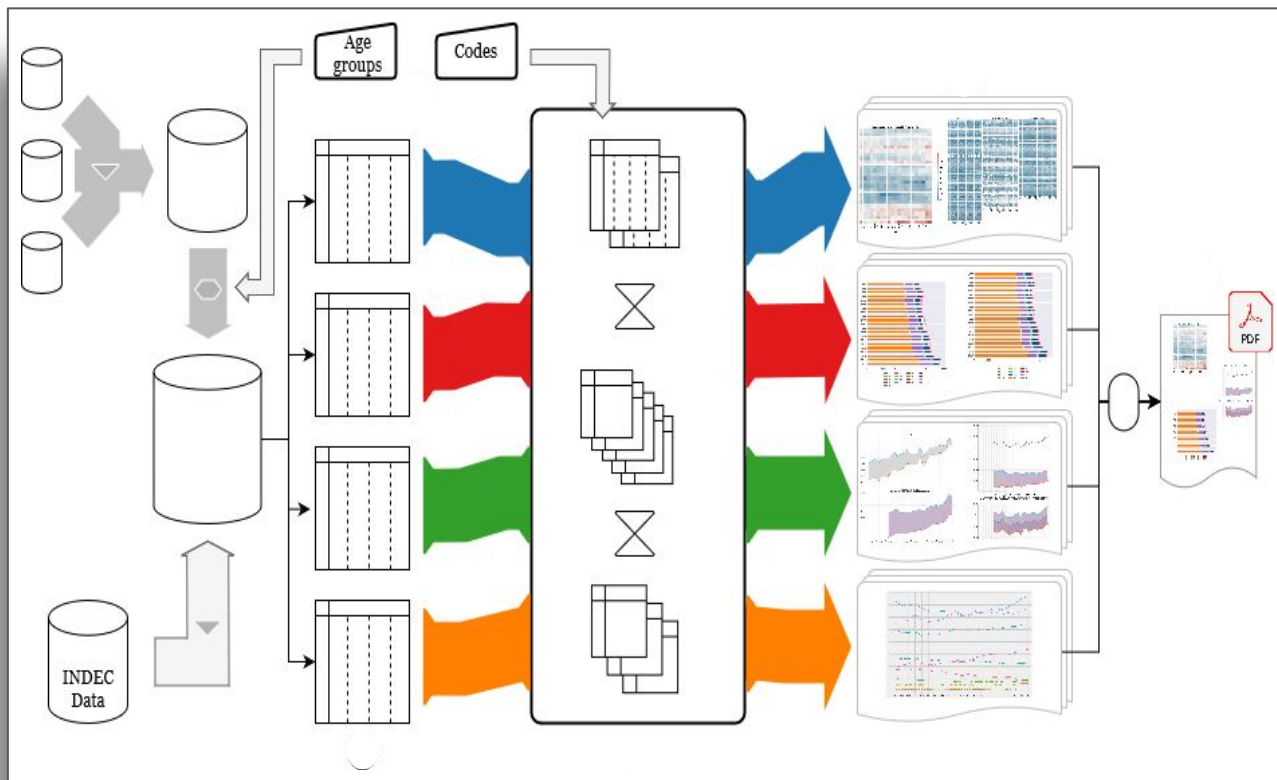


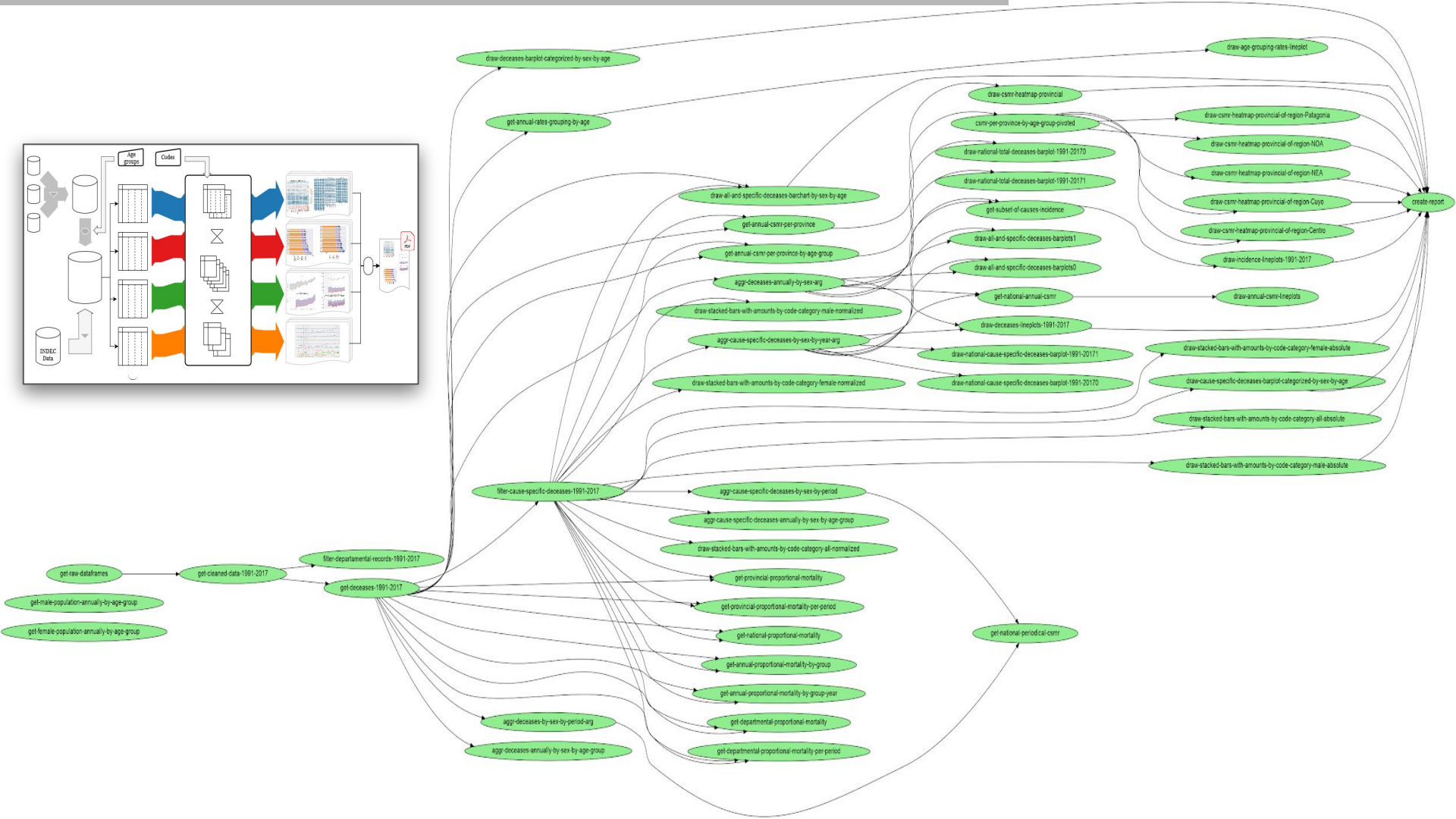
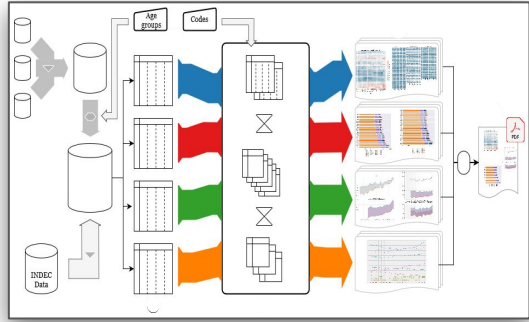
get ->
File('products/get.html')
, File('products/raw.csv')

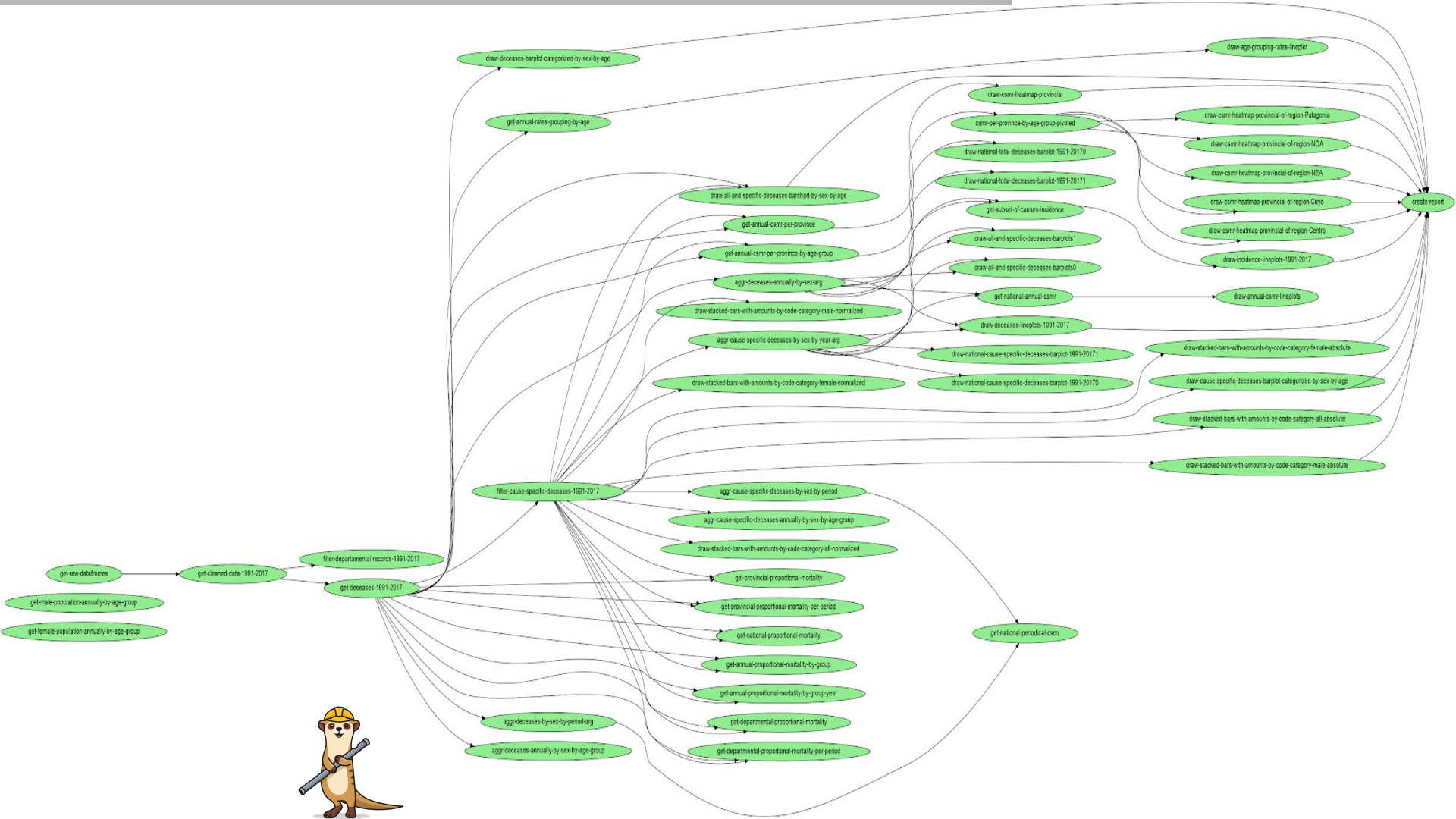
clean ->
File('products/clean.html')
, File('products/clean.csv')

profile-clean ->
File('products/report-clean.html')

custom ->
File('products/custom.html')







Conclusiones

1) Modularización → mantenimiento

Modularización → mantenimiento

	AÑO	PROVRES	DEPRES	CUENTA
0	1994.0	2.0	1.0	1380.0
1	1994.0	2.0	2.0	1321.0
2	1994.0	2.0	3.0	881.0
3	1994.0	2.0	4.0	933.0
4	1994.0	2.0	5.0	1951.0

Modularización → mantenimiento

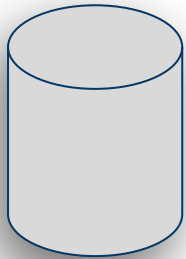
	AÑO	PROVRES	DEPRES	CUENTA
0	1994.0	2.0	1.0	1380.0
1	1994.0	2.0	2.0	1321.0
2	1994.0	2.0	3.0	1832.0
3	1994.0	2.0	4.0	933.0
4	1994.0	2.0	5.0	1951.0

1832.0

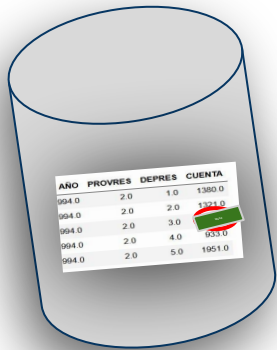
Modularización → mantenimiento

	AÑO	PROVRES	DEPRES	CUENTA
0	1994.0	2.0	1.0	1380.0
1	1994.0	2.0	2.0	1321.0
2	1994.0	2.0	3.0	1832.0
3	1994.0	2.0	4.0	933.0
4	1994.0	2.0	5.0	1951.0

1832.0



Modularización → mantenimiento



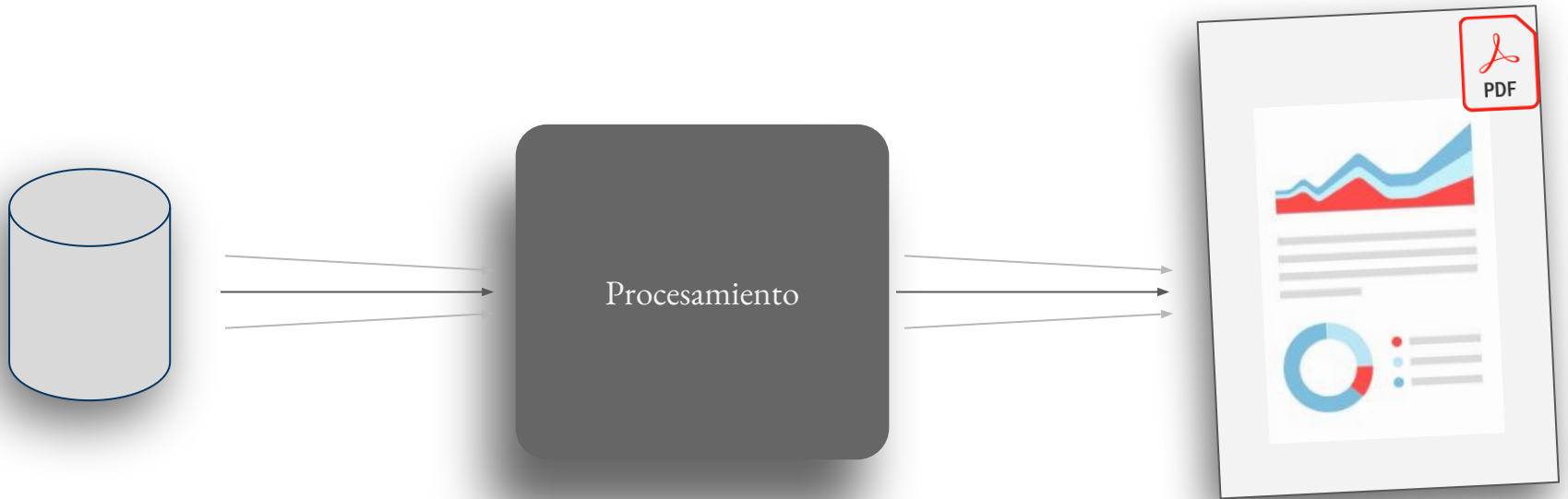
A 3D cylinder representing a database. On its side is a table with the following data:

AÑO	PROVRES	DEPRES	CUENTA
994.0	2.0	1.0	1380.0
994.0	2.0	2.0	1921.0
994.0	2.0	3.0	65.0
994.0	2.0	4.0	633.0
994.0	2.0	5.0	1951.0



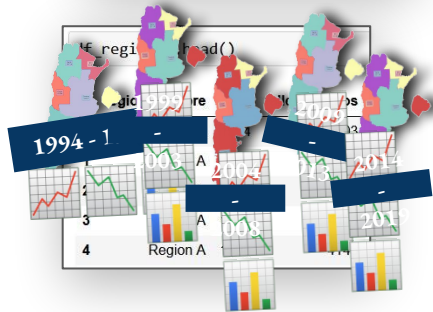
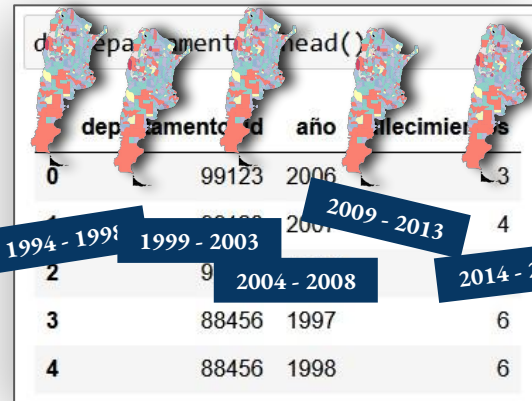
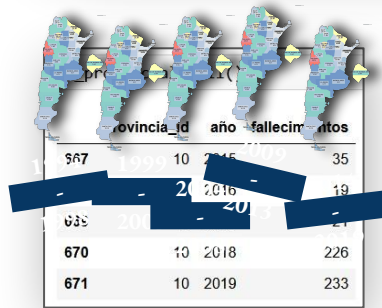

```
$ ploomber build
```

Modularización → mantenimiento

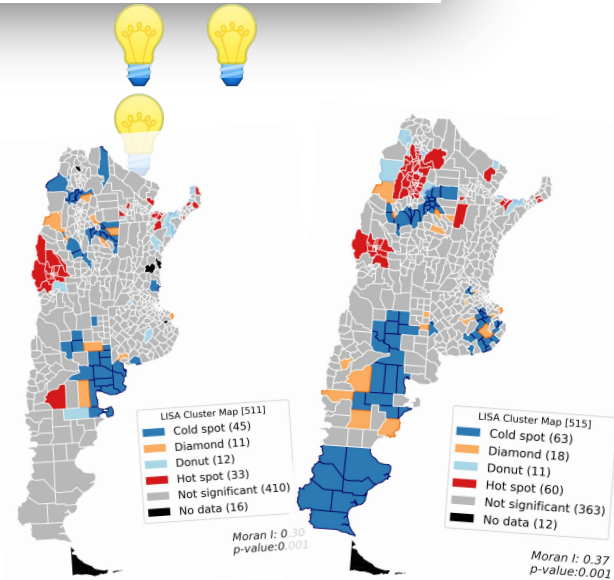
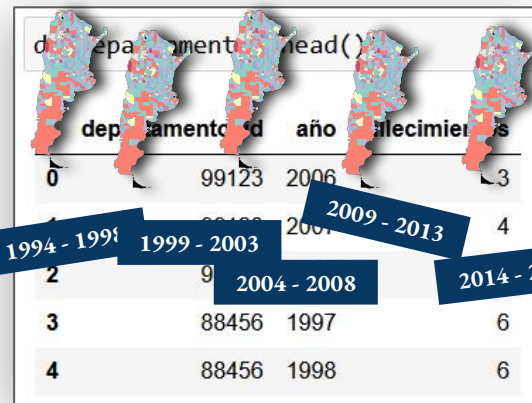
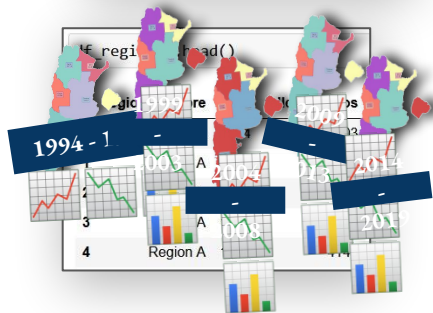
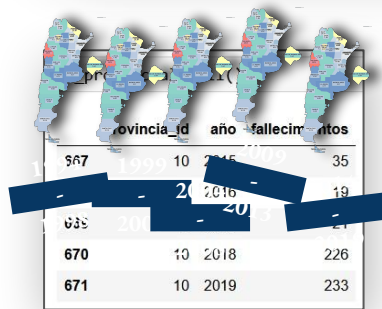


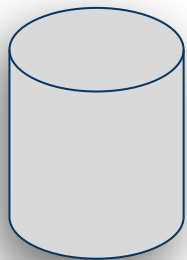
2) Modularización → evolución

	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN

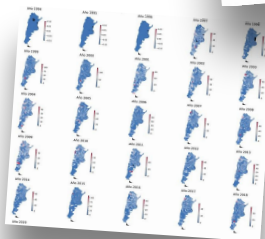


	AÑO	JURIREG	PROVRES	DEPRES	CAUSAMUERCIE10	TIEMGEST	PESOFETO	Unnamed: 7	Unnamed: 8
0	1994.0	62.0	62.0	NaN	A41	20.0	300.0	NaN	NaN
1	1994.0	62.0	62.0	NaN	A41	35.0	2600.0	NaN	NaN
2	1994.0	6.0	6.0	NaN	A50	34.0	3050.0	NaN	NaN
3	1994.0	6.0	6.0	NaN	A50	33.0	2000.0	NaN	NaN
4	1994.0	6.0	6.0	NaN	A50	37.0	3070.0	NaN	NaN

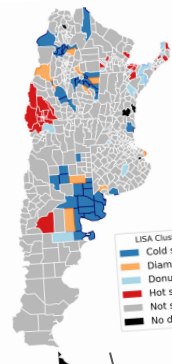
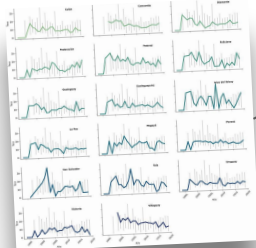




Procesamiento

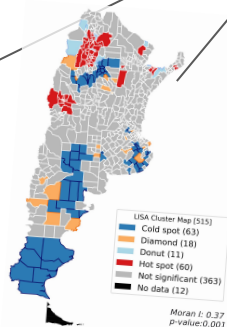


Year	Value	Year	Value	Year	Value	Year	Value	Year	Value
2004-2005	10.28	2005-2006	10.28	2006-2007	10.28	2007-2008	10.28	2008-2009	10.28
2009-2010	10.28	2010-2011	10.28	2011-2012	10.28	2012-2013	10.28	2013-2014	10.28
2014-2015	10.28	2015-2016	10.28	2016-2017	10.28	2017-2018	10.28	2018-2019	10.28
2019-2020	10.28	2020-2021	10.28	2021-2022	10.28	2022-2023	10.28	2023-2024	10.28



LISA Cluster Map [511]
Cold spot (45)
Diamond (11)
Donut (12)
Hot spot (33)
Not significant (410)
No data (16)

Moran I: 0.30
p-value: 0.001

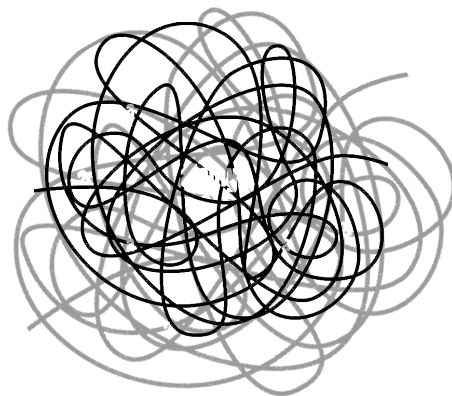


LISA Cluster Map [515]
Cold spot (63)
Diamond (18)
Donut (11)
Hot spot (60)
Not significant (363)
No data (12)

Moran I: 0.37
p-value: 0.001

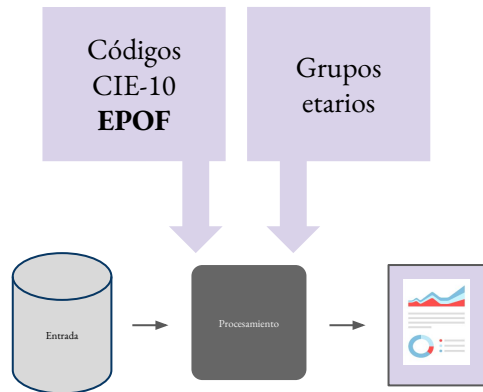


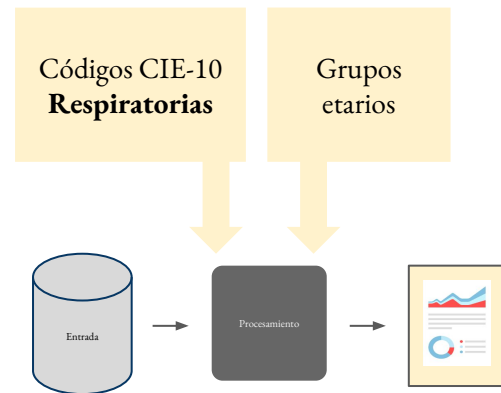
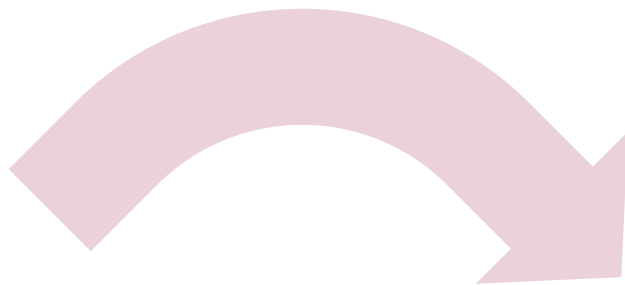
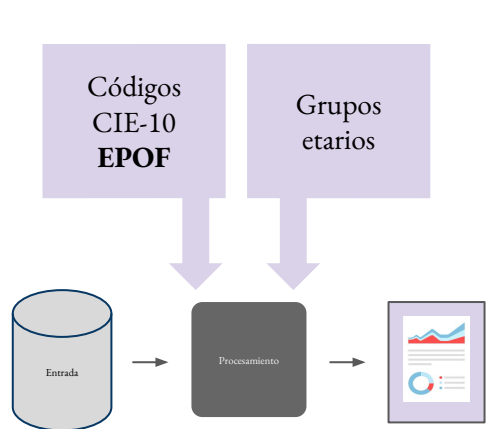
3) Mejora la interacción del equipo

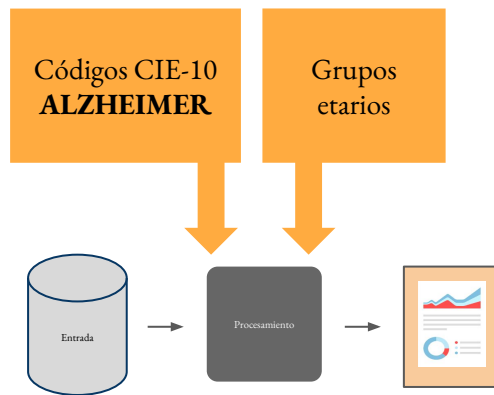
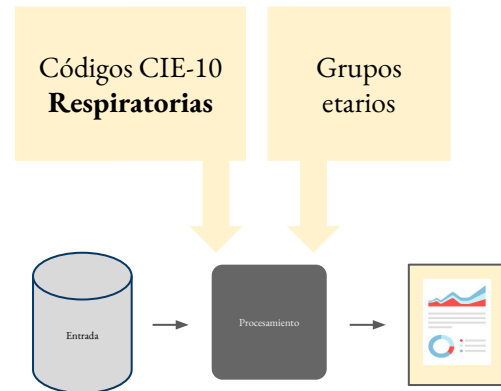
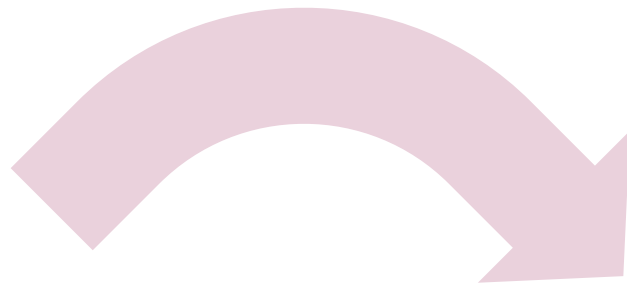
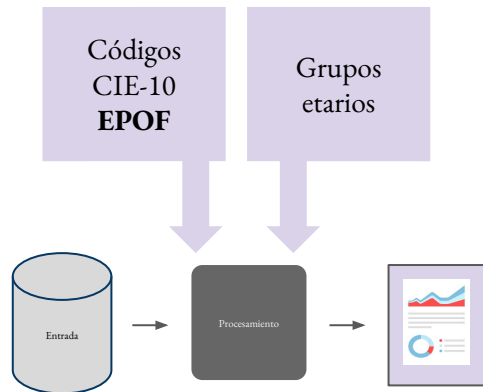


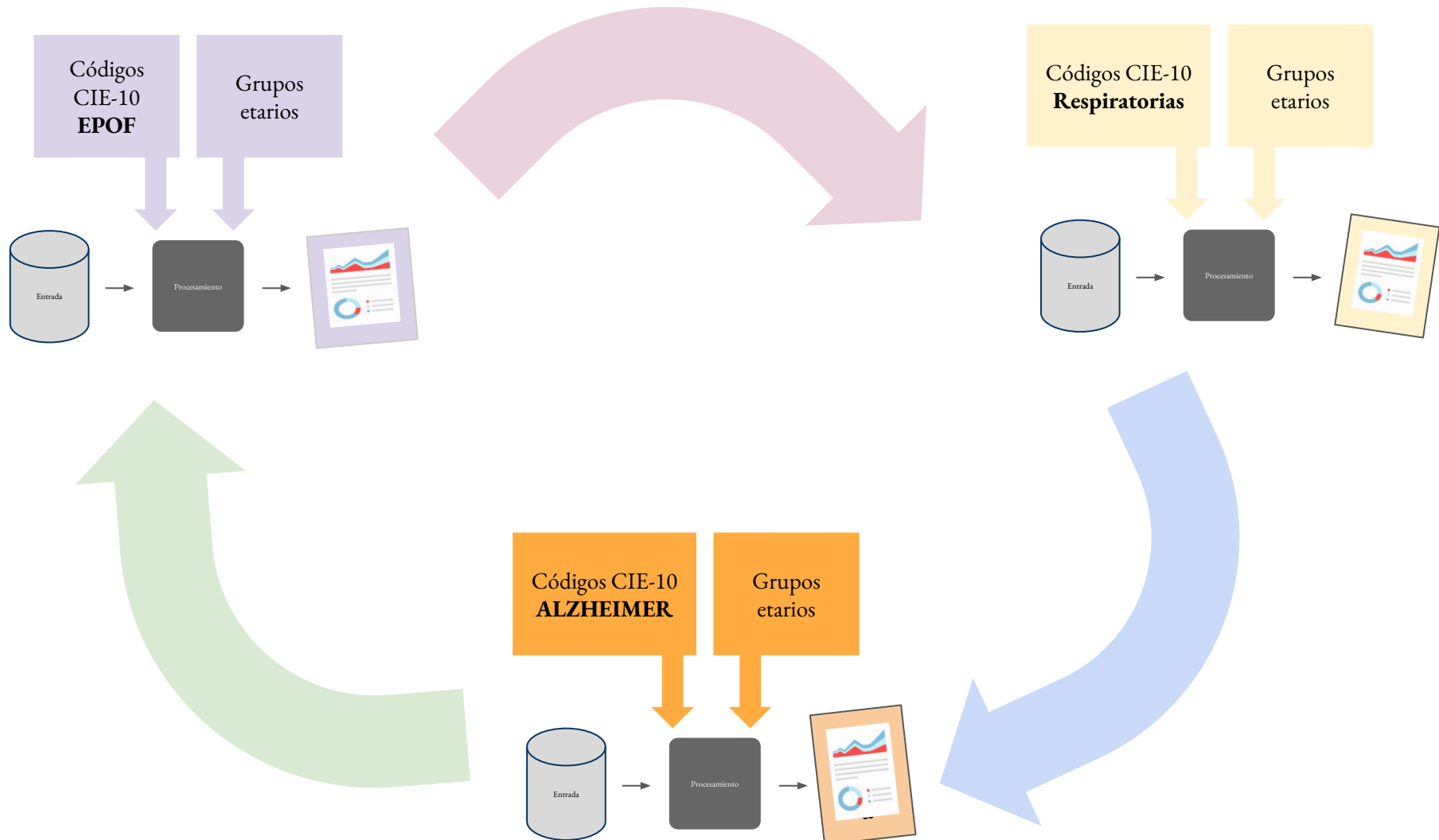


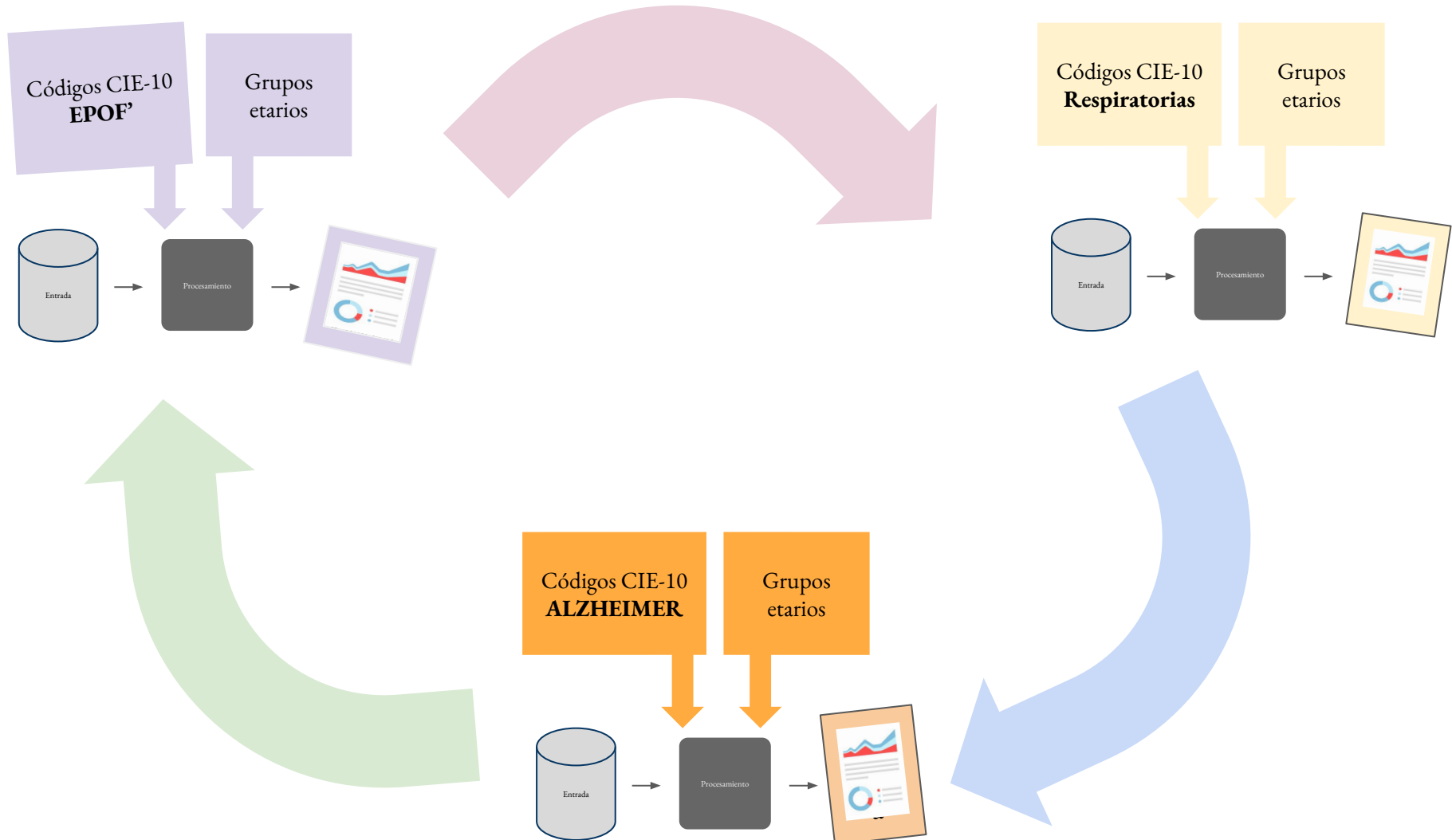
4) Parametrización → Reutilización

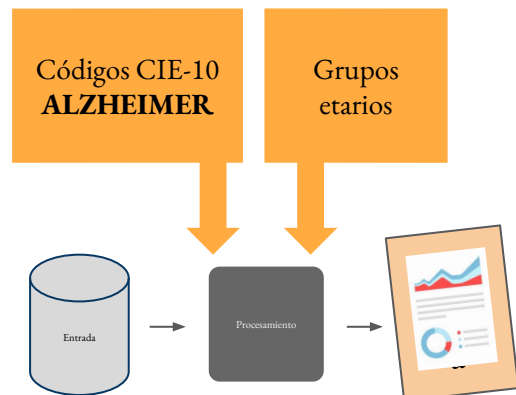
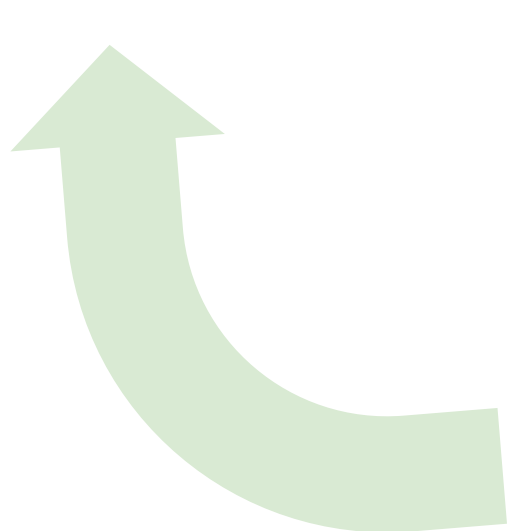
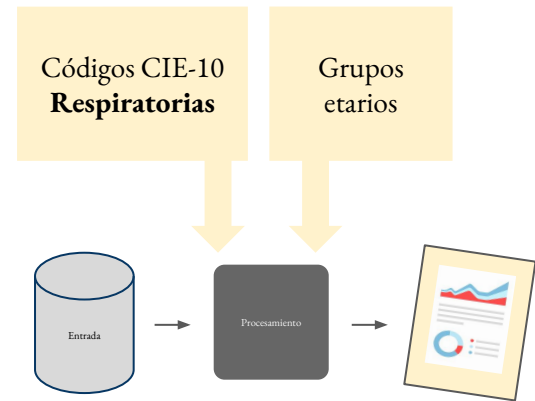












Herramienta para la exploración de tendencias y detección de patrones epidemiológicos en Argentina

Morales Arturo Leonardo, Figueroa Marcelo , Delrieux Claudio, Ramallo Virginia, Dipierri José Edgardo

Muchas gracias