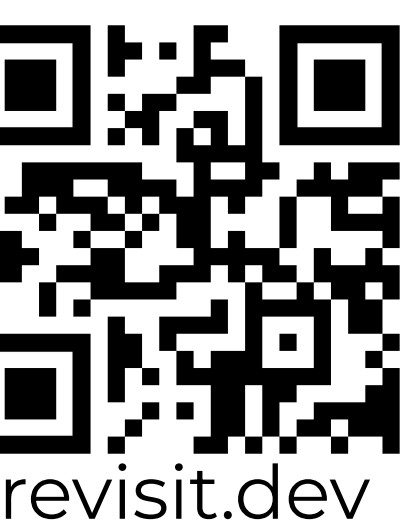


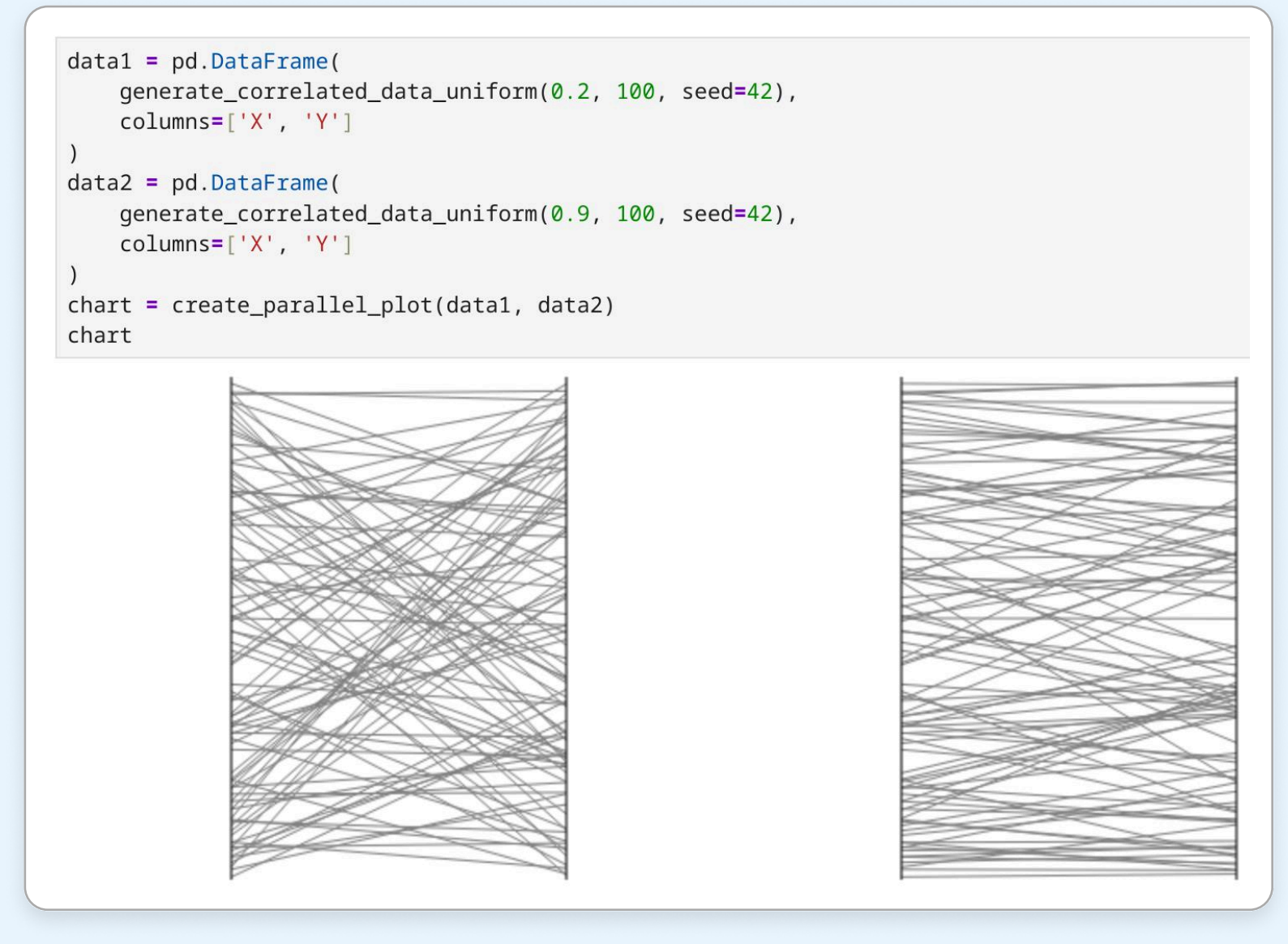
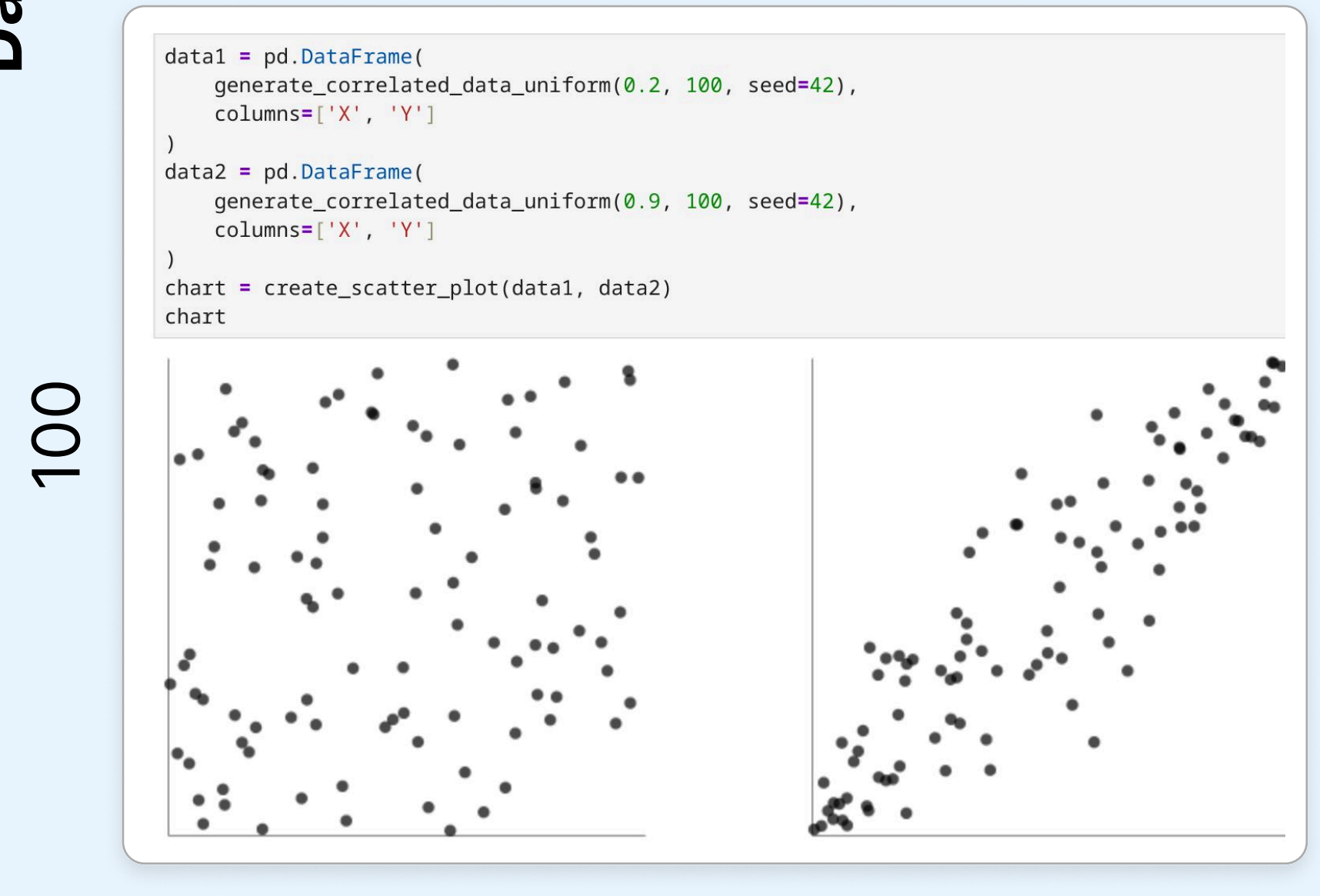
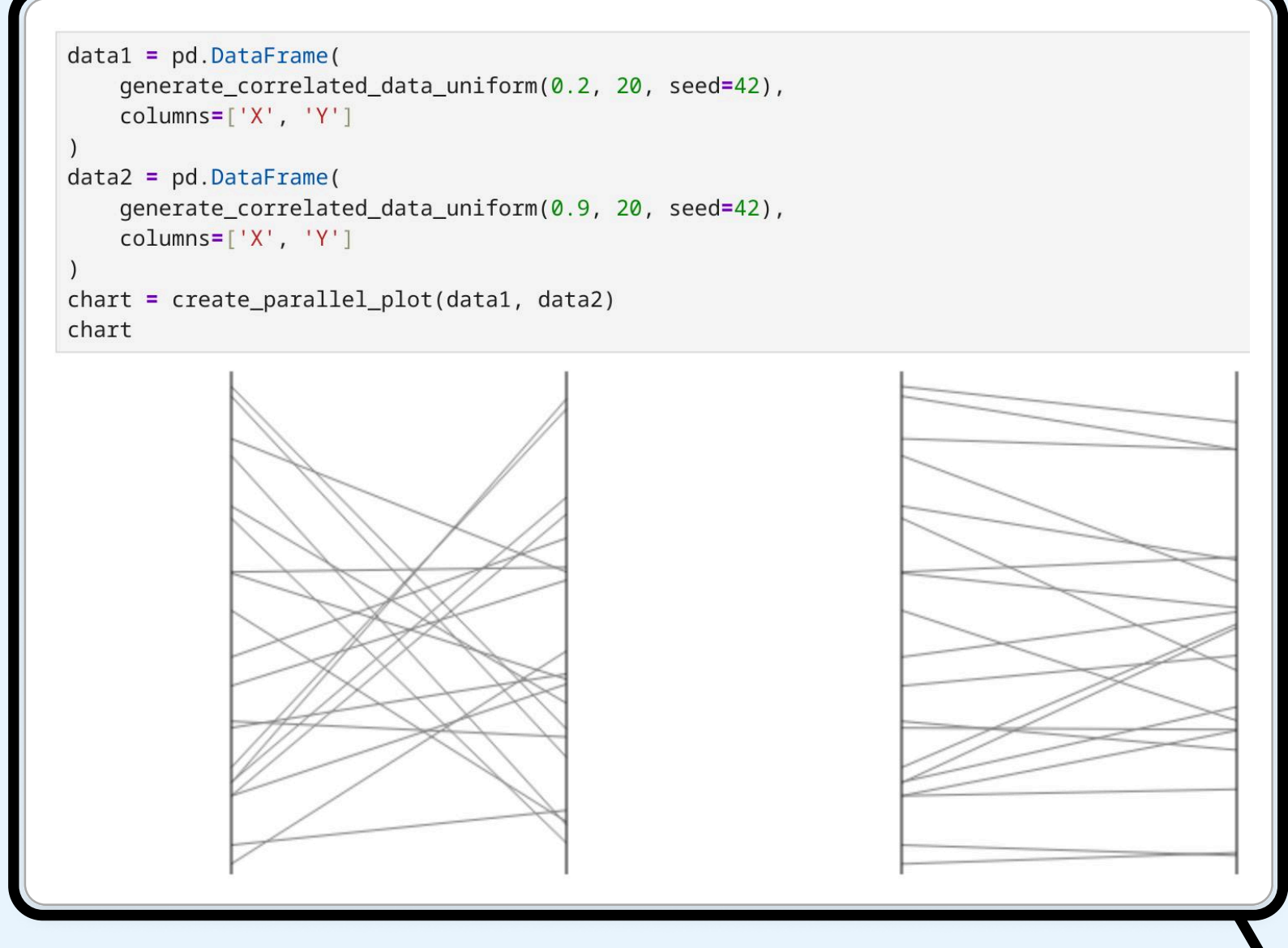
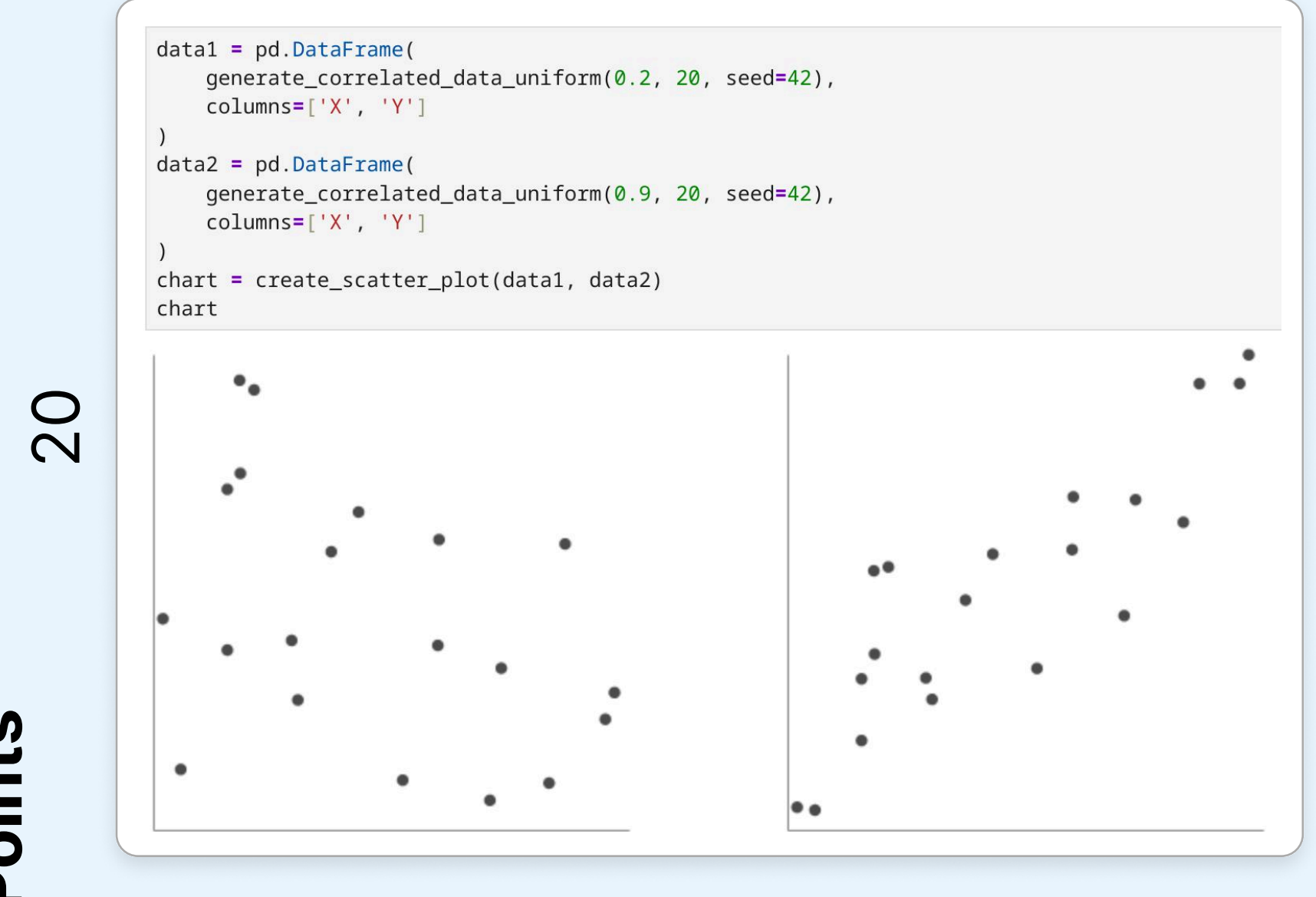


# ReVISitPy: Python Bindings for the reVISit Study Framework

H. Shrestha, J. Wilburn, B. Bollen, A. M. McNutt, A. Lex, L. Harrison



Data Points



Scatterplot

Parallel Coordinates Plot

### Visualization Types

ReVISitPy is a Python package that enables experiment designers to **create and test visualization studies** with potentially **enormous configuration files** for the reVISit study framework.

- Correlation 0.1 vs. 0.2
- Correlation 0.1 vs. 0.3
- Correlation 0.1 vs. 0.4
- Co-relation 0.1 vs. 0.5
- Correlation 0.2 vs. 0.9**
- Correlation 0.9 vs. 1.0

```
{
  "components": {
    "parallelCoords-0.1.0.0.0.0": {
      "allowFailedTraining": null,
      "config": {
        "$schema": "https://open.vega.io/sch",
        "background": "white",
        "padding": 5,
        "height": 300,
        "data": {
          "signals": [
            "layout": {
              "marks": [
                "scales": [
                  "config": {
                    "correctAnswer": null,
                    "description": null,
                    "helpTextPathOverride": null,
                    "instruction": null,
                    "instructionLocation": null,
                    "meta": null,
                    "nextButtonDisableTime": null,
                    "nextButtonEnableTime": null,
                    "nextButtonLocation": null,
                    "nextButtonText": null,
                    "provideFeedback": null,
                    "recordAudio": null,
                    "response": {
                      "type": "radio"
                    }
                  }
                ]
              }
            }
          }
        }
      }
    }
  }
}
```

Generated Trials: 45

~232,000 lines of code for the entire study!

```
import itertools

# Generate all combinations of two values between 1 and 10
combinations = itertools.combinations(range(1, 11), 2)

# Create the dataset with values divided by 10
dataSet = [{"corrValues": [x / 10, y / 10]} for x, y in combinations]

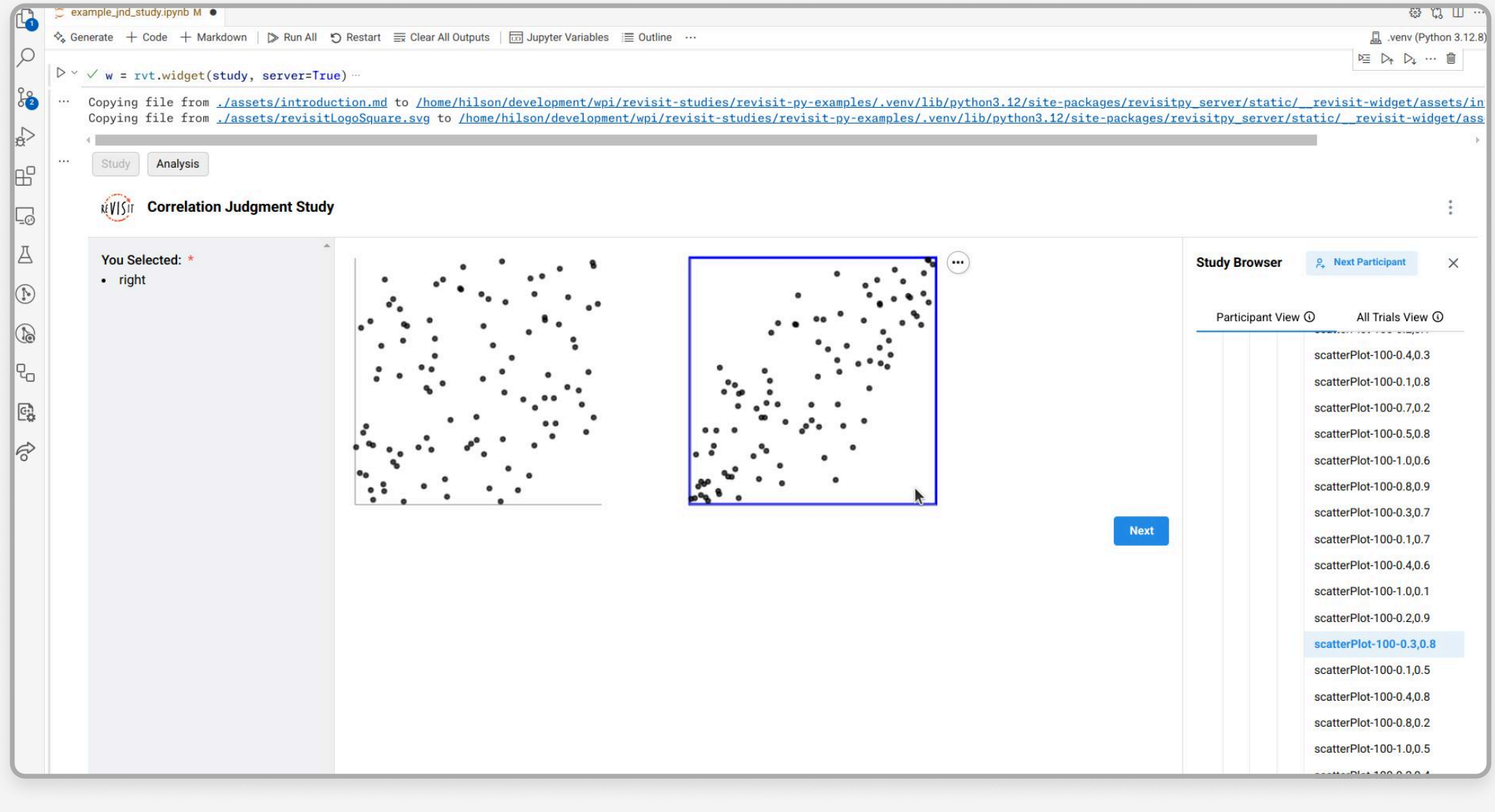
main_sequence = rvt.sequence(order='fixed')

main_sequence.permute(
    factors=[('visType': 'scatterPlot'), ('visType': 'parallelCoords')],
    order='latinSquare',
).permute(
    factors=[('numPoints': 20), ('numPoints': 100)],
    order='fixed',
).permute(
    factors=dataSet,
    order='random',
).component(component_function)

sequence = rvt.sequence(order='fixed', components=[introduction]) + main_sequence

study = rvt.studyConfig(
    schema="https://raw.githubusercontent.com/revisit-studies/study/v2.0.0-rc1/src/parser/StudyConfigSchema.json",
    uiConfig=ui_config,
    studyMetadata=study_metadata,
```

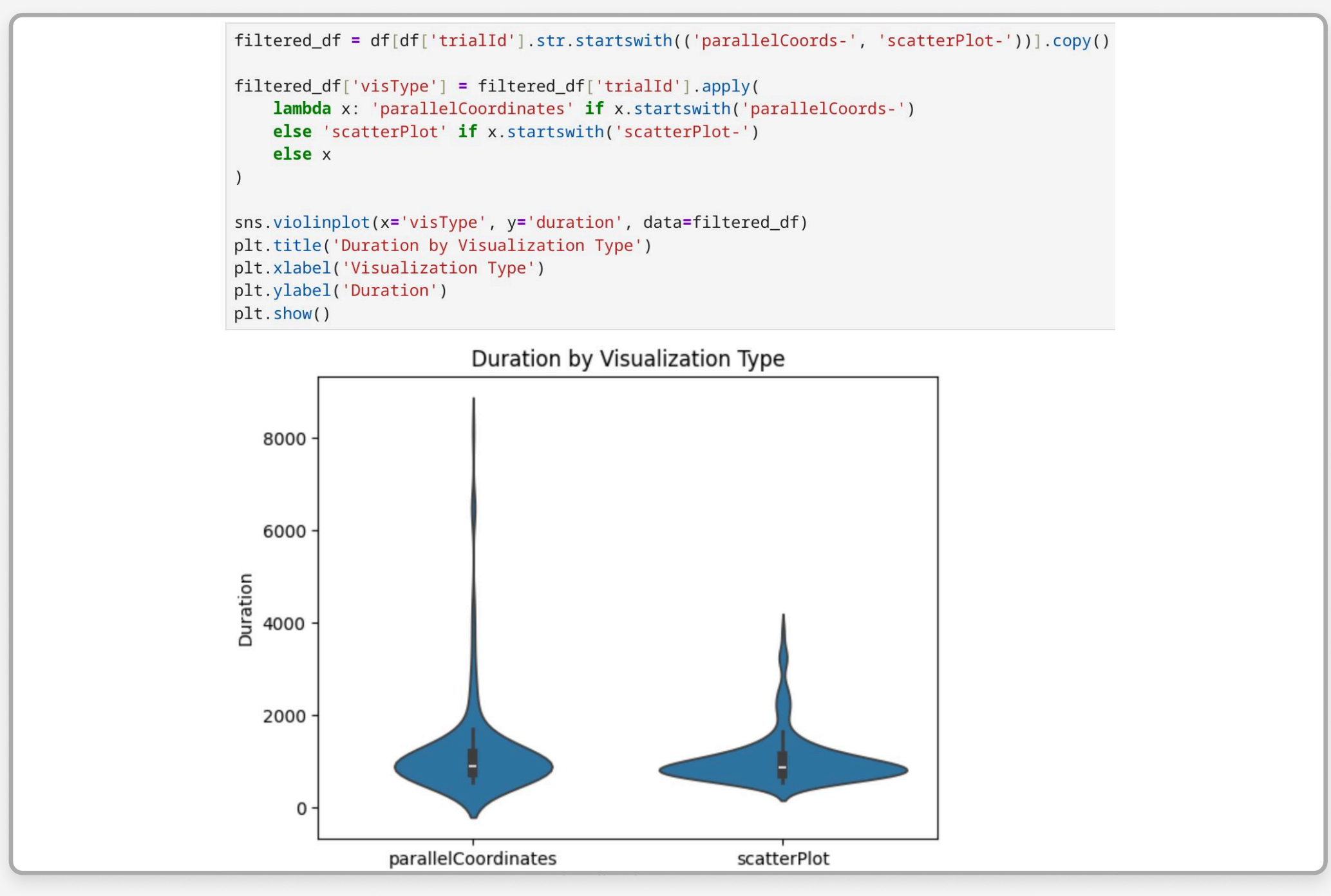
A Create reVISit specification



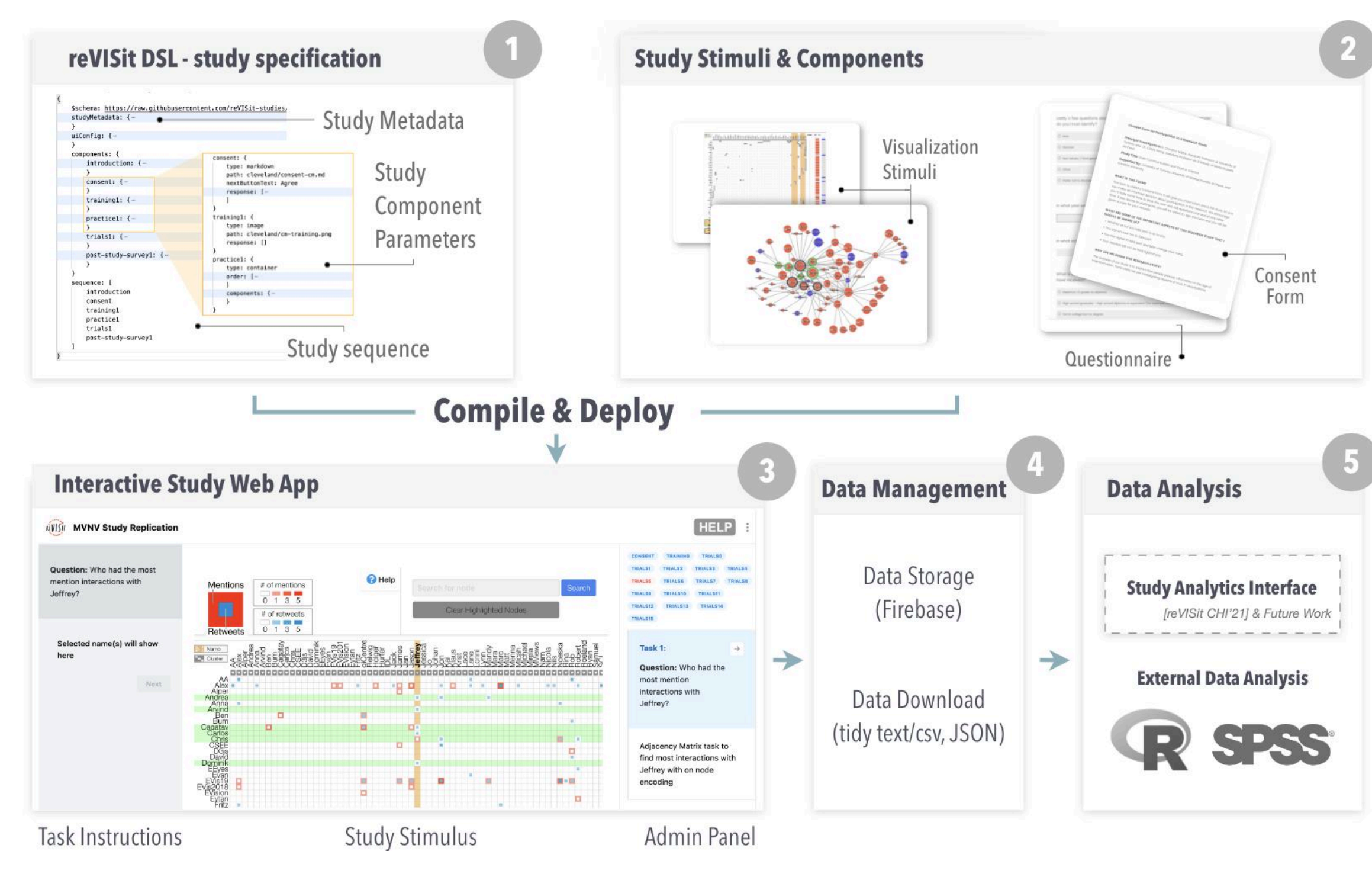
B Preview study in the anyWidget powered reVISit interface

id	participantId	trialId	order	response	status	responseTime	responseTime	description	instruction	responseTarget	answer	correctAnswer	responseTime	startTime
0	01W46C00	01W46C00	0	introduction	0	completed	NAN	NAN	100.00	NAN	NAN	NAN	NAN	2025.00
1	01W46C00	01W46C00	1	vegAndMeatResponse	1	completed	NAN	NAN	100.00	NAN	NAN	NAN	NAN	2025.00
2	01W46C00	01W46C00	2	vegAndMeatResponse	2	completed	NAN	NAN	100.00	NAN	NAN	NAN	NAN	2025.00
3	01W46C00	01W46C00	3	vegAndMeatResponse	3	completed	NAN	NAN	100.00	NAN	NAN	NAN	NAN	2025.00
4	01W46C00	01W46C00	4	vegAndMeatResponse	4	completed	NAN	NAN	100.00	NAN	NAN	NAN	NAN	2025.00

C Export data from reVISit's analysis interface



D Perform preliminary analysis of the data



ReVISit is an **open-source software toolkit** and framework for creating, deploying, and monitoring user studies. Running quality empirical studies can be demanding and resource-intensive, requiring significant time, cost, and expertise. ReVISit addresses these challenges by introducing a domain-specific language for designing experiments

ReVISitPy provides easy-to-use functions for developing, testing and performing preliminary analysis of an entire studies from within a **single python notebook**.



Award Number 2213756 & 2213757