

jupyter Pop-up

Share a Little Bit
of Your Notebooks With Me

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Hello! 🖐️

- Peter Parente (@parente)
- Jupyter Steering Council Member
- Valassis Digital employee
- Purveyor of the analytics user experience

The background of the slide features a dark grey area on the left with a complex network diagram. The diagram consists of numerous teal-colored dots of varying sizes connected by thin, light blue lines, creating a web-like structure that curves from the top left towards the center.

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That title ... 🤔



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Today's Topic

Tools and techniques for sharing notebooks, borne out of personal experience in open communities and private workplaces.



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Just post it on GitHub?

First, consider:

- Audience
- Reach
- Content
- Medium
- Reproducibility
- Longevity
- Privacy
- Technology

The background of the slide features a dark grey area on the left containing a complex network diagram. This diagram consists of numerous teal-colored dots of varying sizes, interconnected by thin, light blue lines. The dots are arranged in a way that suggests a radial or hierarchical structure, with some dots acting as hubs and others as peripheral nodes. The overall effect is a dense, web-like pattern that fills the left side of the frame.

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I'll give 🖐️ examples to you

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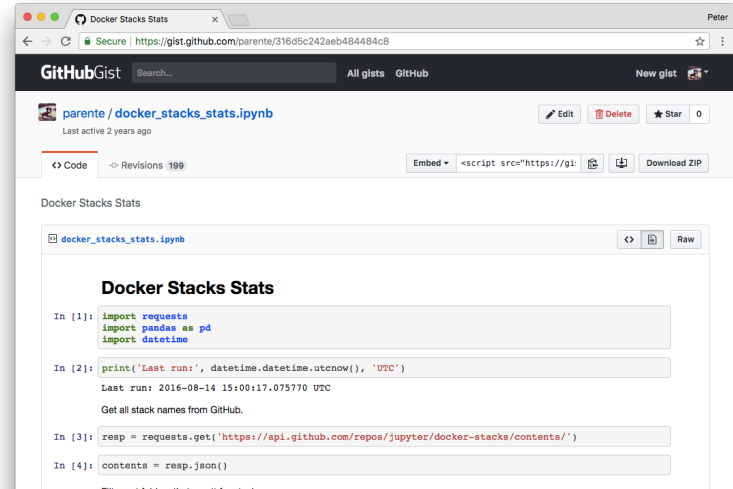
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Publish a one-off document

Consider a gist

Demo: Jupyter Docker Stats





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Publish a one-off document

Consider a gist

1. Install a [gist extension](#).
2. Enter a [GitHub Personal Access token](#).
3. Click a button to publish.

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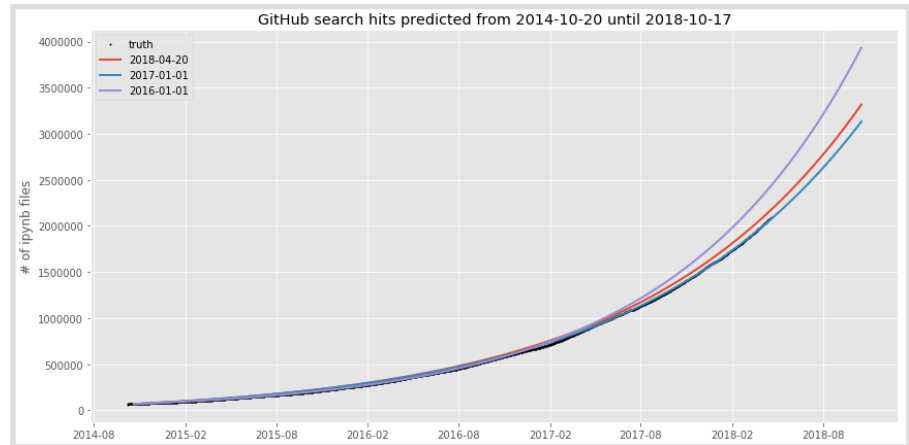
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Regularly update an analysis

Consider scripted notebook execution

Demo: [nbestimate](#)





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Regularly update an analysis

Consider scripted notebook execution

1. Programmatically execute notebooks.
2. Commit and push to a git repository.
3. Automate what you can with bash, Python, [nbflow](#), [papermill](#), [anaconda-project](#), etc.

Detail: Using nbconvert to execute notebooks

```
import nbformat
from nbconvert.preprocessors import ExecutePreprocessor

with open('my_notebook.ipynb') as fh:
    nb = nbformat.read(fh, 4)

exp = ExecutePreprocessor(timeout=60)
rerun_nb, _ = exp.preprocess(nb, {})
```


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Create a learning playground

Consider Binder

Demo: [Wild Wolf Watch](#)



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Create a learning playground

Consider Binder

1. Push a notebook and requirements to GitHub.
2. Enter your details at mybinder.org.
3. Get a link or button to share.

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Museum Life+Science
@lifeandscience



New red wolf pups! 🎉🐾 The Animal Department counted three: 2 male & 1 female! All pups were noted as being in good health. On the Keeper blog: behind-the-scenes pics, their first vet check, & more! buff.ly/2qTfQjg #redwolves #newarrivals #lifeandscience #durhamnc

10:00 AM - Apr 22, 2018



44



17

people are talking about this

twitter.com/lifeandscience/status/988054922355793921

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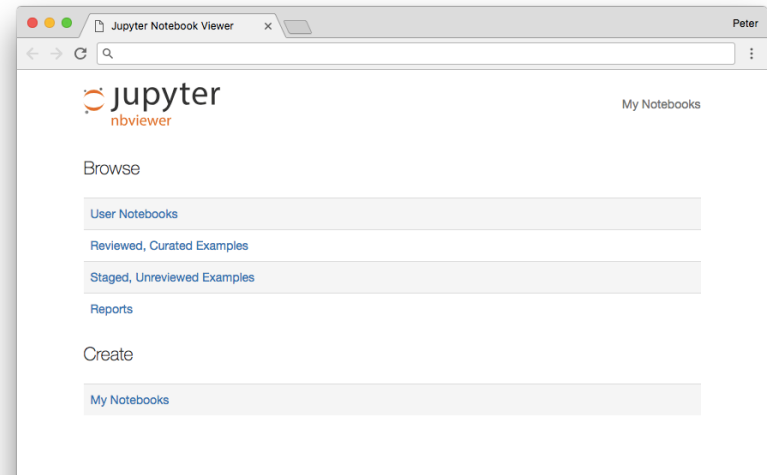
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Access team work-in-progress

Consider running a local notebook viewer

Demo: nbviewer in JupyterHub





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Access team work-in-progress

Consider running a local notebook viewer

1. Install [nbviewer](#) or [commuter](#).
2. Point it to user directories or S3 buckets.
3. Secure it (e.g., [as a JupyterHub service](#)).

Detail: nbviewer as a JupyterHub service

```
c.JupyterHub.services = [  
    {  
        'name': 'nbviewer',  
        'url': 'http://127.0.0.1:9000',  
        'cwd': '/home',  
        'command': ['python', '-m', 'nbviewer', '--localfiles']  
    }  
]
```


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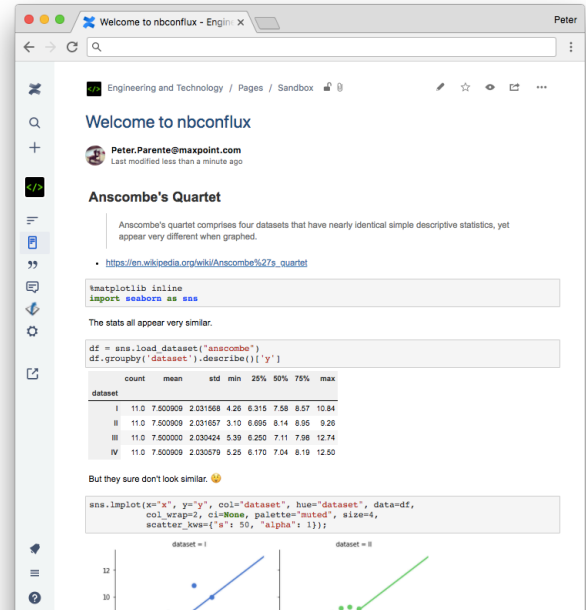
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Share with a broader org

Consider knowledge base integration

Demo: *nbconflux*





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Share with a broader org

Consider knowledge base integration

1. Use [nbconvert preprocessors and exporters](#).
2. Package conversion scripts and APIs.
3. Create [extensions for one-click publishing](#).



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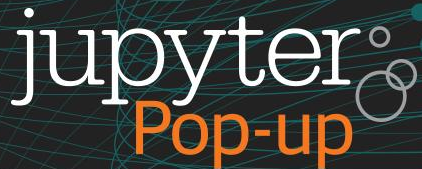
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Come along!

- Jupyter notebooks are JSON files
- They have a [formal schema](#)
- nbconvert uses Python + [jinja](#) to transform them
- You can use whatever tools / languages you like
 - e.g., [@nteract/notebook-preview](#) is JS+React

Detail: A simple, valid notebook

```
import nbformat
simple = {
    "nbformat": 4,
    "nbformat_minor": 4,
    "metadata": {
        "kernelspec": {
            "name": "python3",
            "display_name": "Python 3"
        }
    },
    "cells": []
}
nbformat.validate(simple, version=4)
```

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Going home

Remember:

- Many tools exist to help you share notebooks
- It's OK to use them as-is or extend them to fit your workflow
- Consider audience, longevity, privacy, reproducibility, etc. in your selection