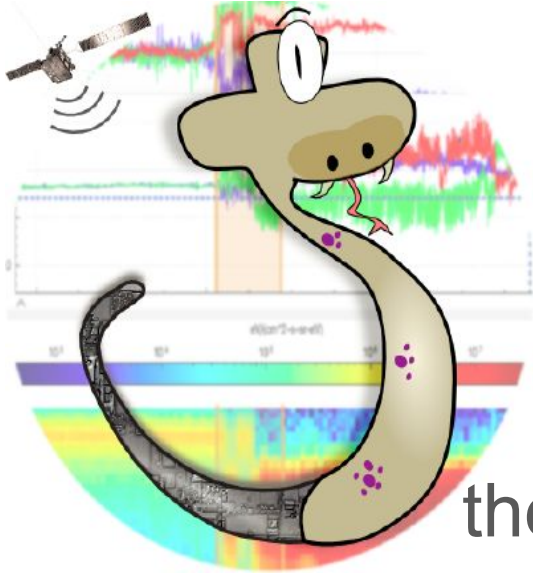




SciQLop:

the **Space Physicist's IDE** Revolution

from archive to science, without leaving your environment

Colloque ATST 2026

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Christian Jacquey², Alexis Rouillard²



Turning data into a plot shouldn't feel like this.



Science is hard enough. **Your tools shouldn't be.**

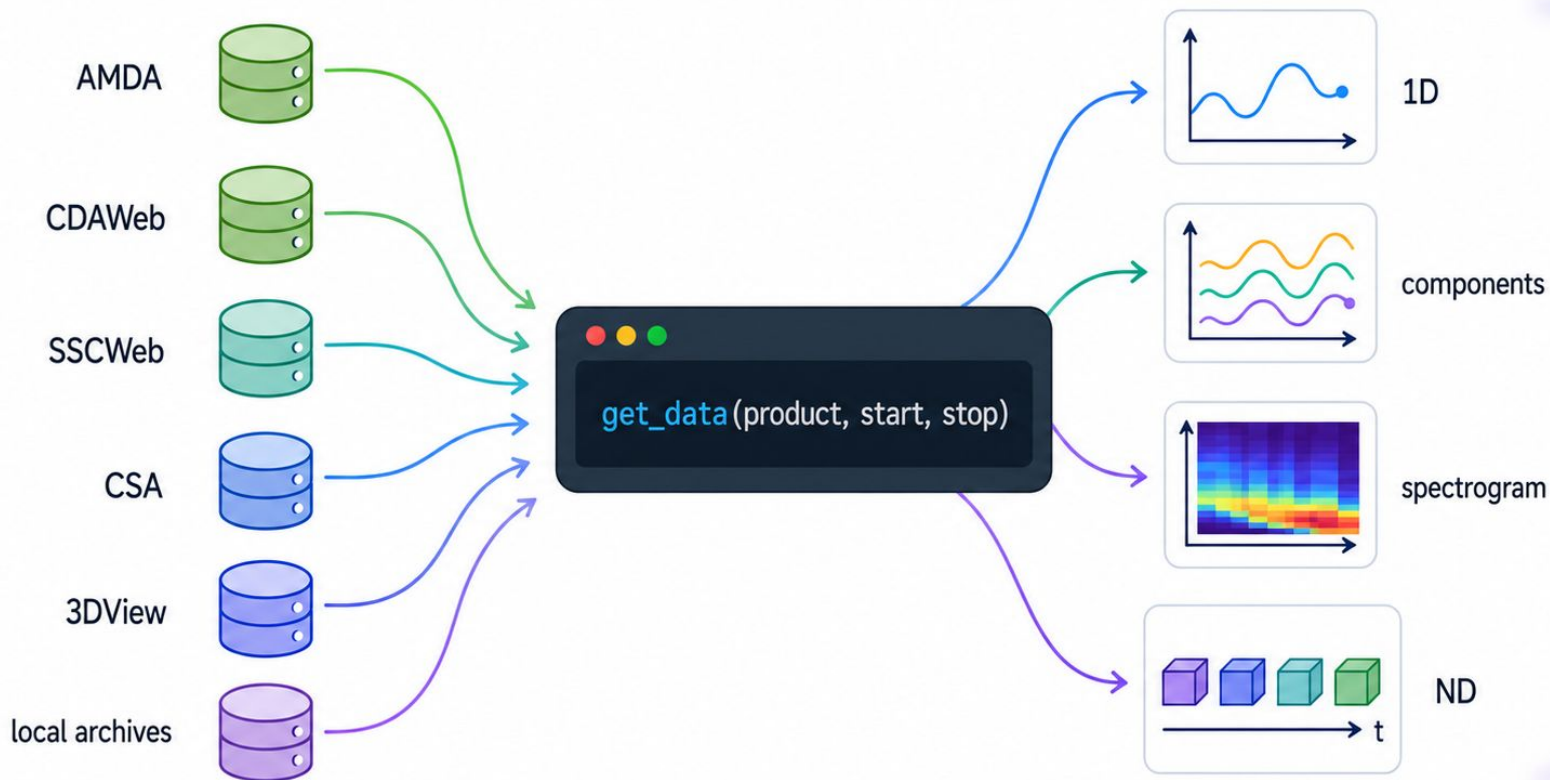
A lot of data in many places

- From **~60 years** of continuous measurements
- **~ 70 space** missions
- **~500TB** of measurements
- Across **~10** different servers
- Many different instruments/products/modes
- Few file formats (CDF, NetCDF, Fits,...)
- Few standards

A lot of use cases

- Quick look?
- Look at an arbitrary set of data?
- Find/discover data?
- Multi-mission analysis?
- Massive statistic study?
- Instrument calibration?
- ...

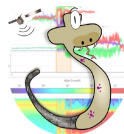
The dream API



The ecosystem

SciQLop

the IDE, browse, visualise, script, annotate, share

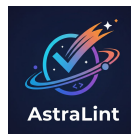


Plugins

AppStore

AstraLint

data quality, validation, standards compliance



```
pip install astralint
```

Speasy

unified access, AMDA, CDA, CSA, SSC...



```
pip install speasy
```

PyCDF++

super-fast C++ CDF reader

```
pip install pycdfpp
```

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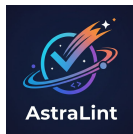


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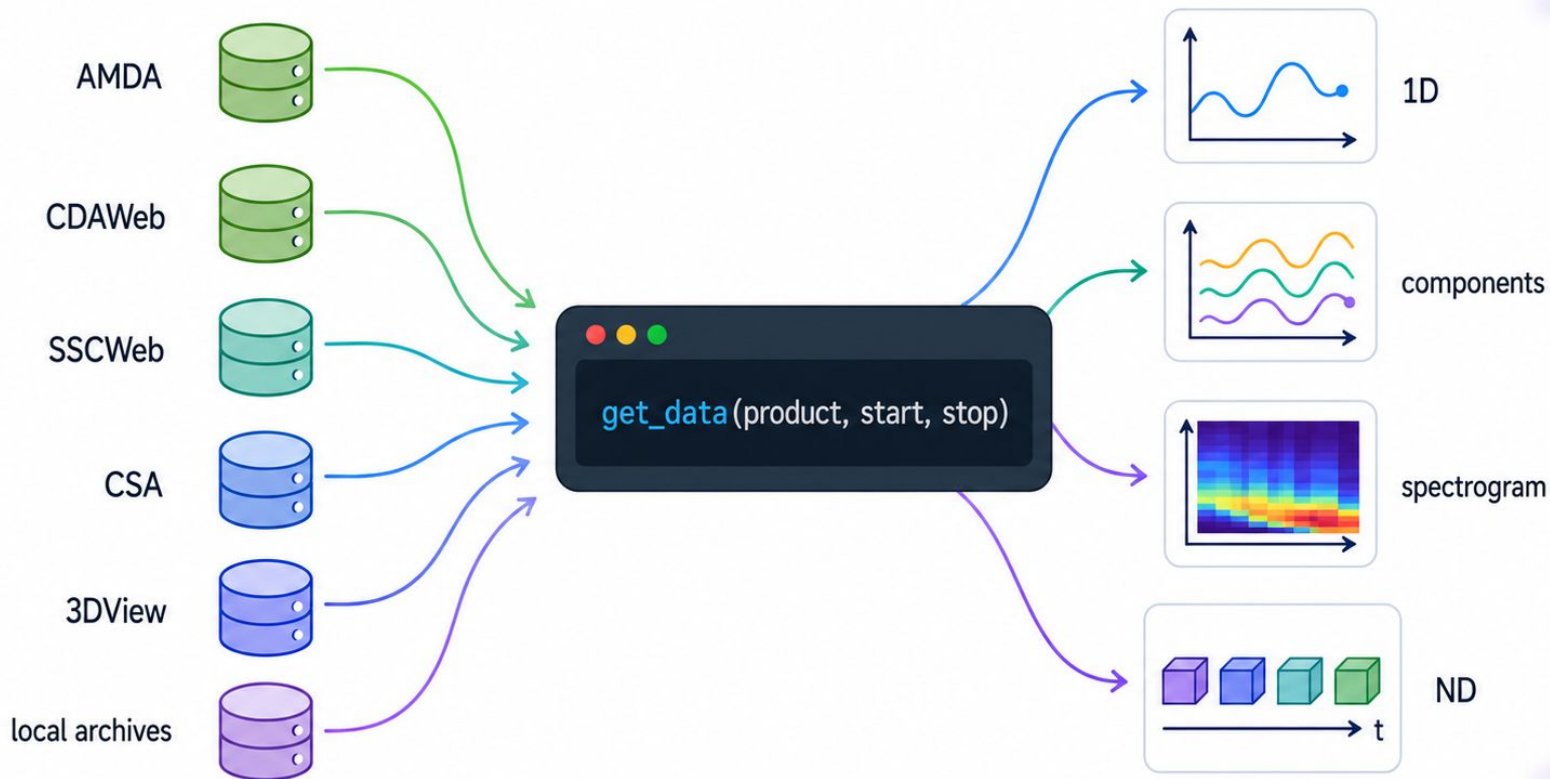
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The dream API



Speasy: one API to rule them all



Speasy: one API to rule them all

70+ missions

AMDA · CDA · CSA · SSC · local

65,000+ products

one call, any archive

Tab-completable

discover data in IPython

Get

```
import speasy as spz  
mag = spz.get_data(  
    'amda/imf',  
    "2016-6-2",  
    "2016-6-5")
```

-> **SpeasyVariable**

Compute

```
import speasy as spz  
import numpy as np  
  
mag_total =  
np.linalg.norm(mag,  
axis=1)
```

-> **SpeasyVariable**

Align

```
from speasy.signal import  
resampling  
  
Tperp_interp, Tpara_interp =  
resampling.interpolate(  
    b, [Tperp, Tpara])
```

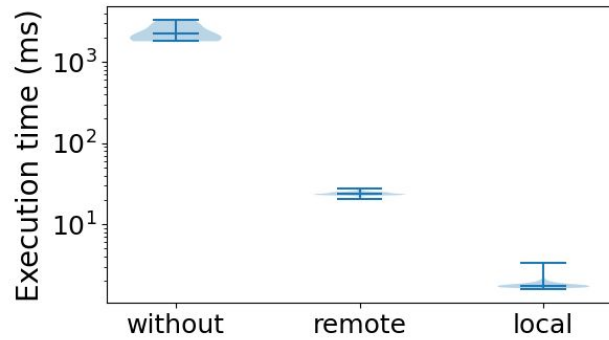
-> on **b**'s time axis

No boilerplate

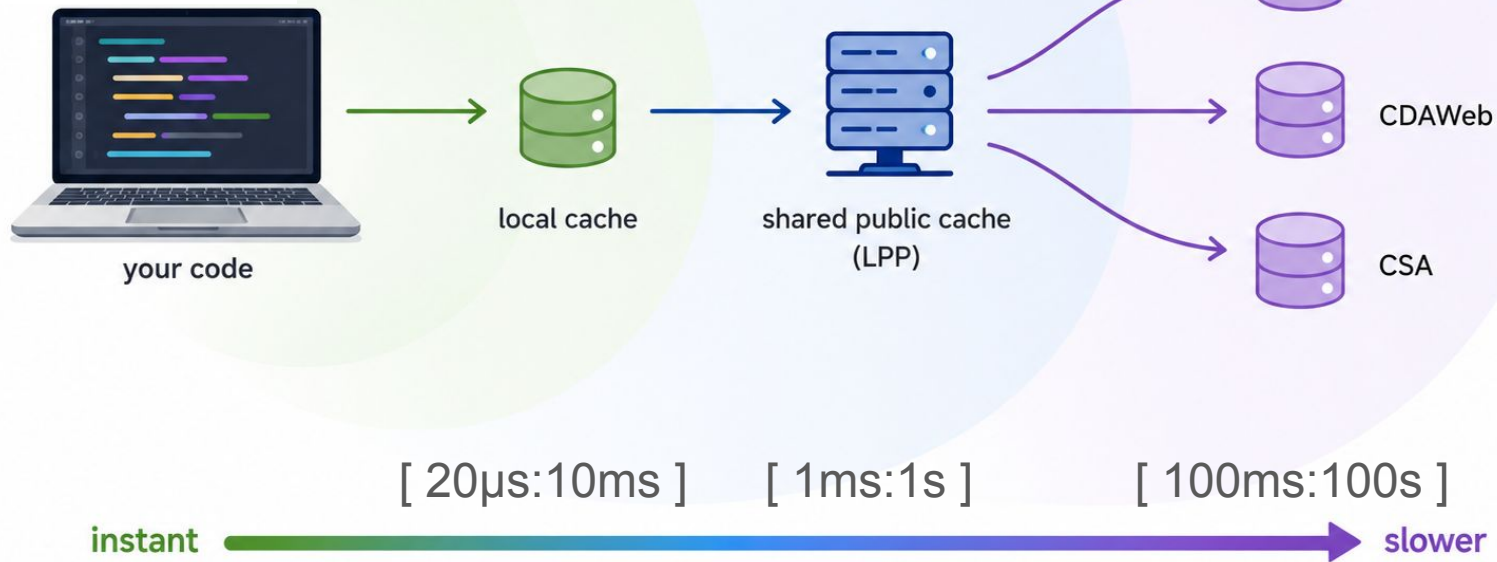
NumPy just works

scipy-compatible, time-aware

Comparison plot



Super fast access to data



```
1 def filtered_B(start, stop):
2     b = spz.get_data("cda/MMS1_FGM_SRVY_L2/mms1_fgm_b_gse_srvy_l2",
3                       start, stop)
4     if b is None:
5         return None
6     return filtering.sosfiltfilt(sos, b)
```

```
def mms1_cos_theta_vb(start, stop):
    B,V=spz.get_data([
        spz.inventories.tree.cda.MMS.MMS1.FGM.MMS1_FGM_BRST_L2.mms1_fgm_b_gse_brst_l2_clean,
        spz.inventories.tree.cda.MMS.MMS1.DIS.MMS1_FPI_BRST_L2_DIS_MOMS.mms1_dis_bulkv_gse_brst
    ], start, stop)

    B = B['Bx GSE', 'By GSE', 'Bz GSE']
    V = interpolate(B,V)
    cos_theta_vb = np.einsum('ij, ij->i', V, B)
                    /(np.linalg.norm(B, axis=1)*np.linalg.norm(V, axis=1))

    return cos_theta_vb
```

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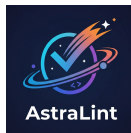


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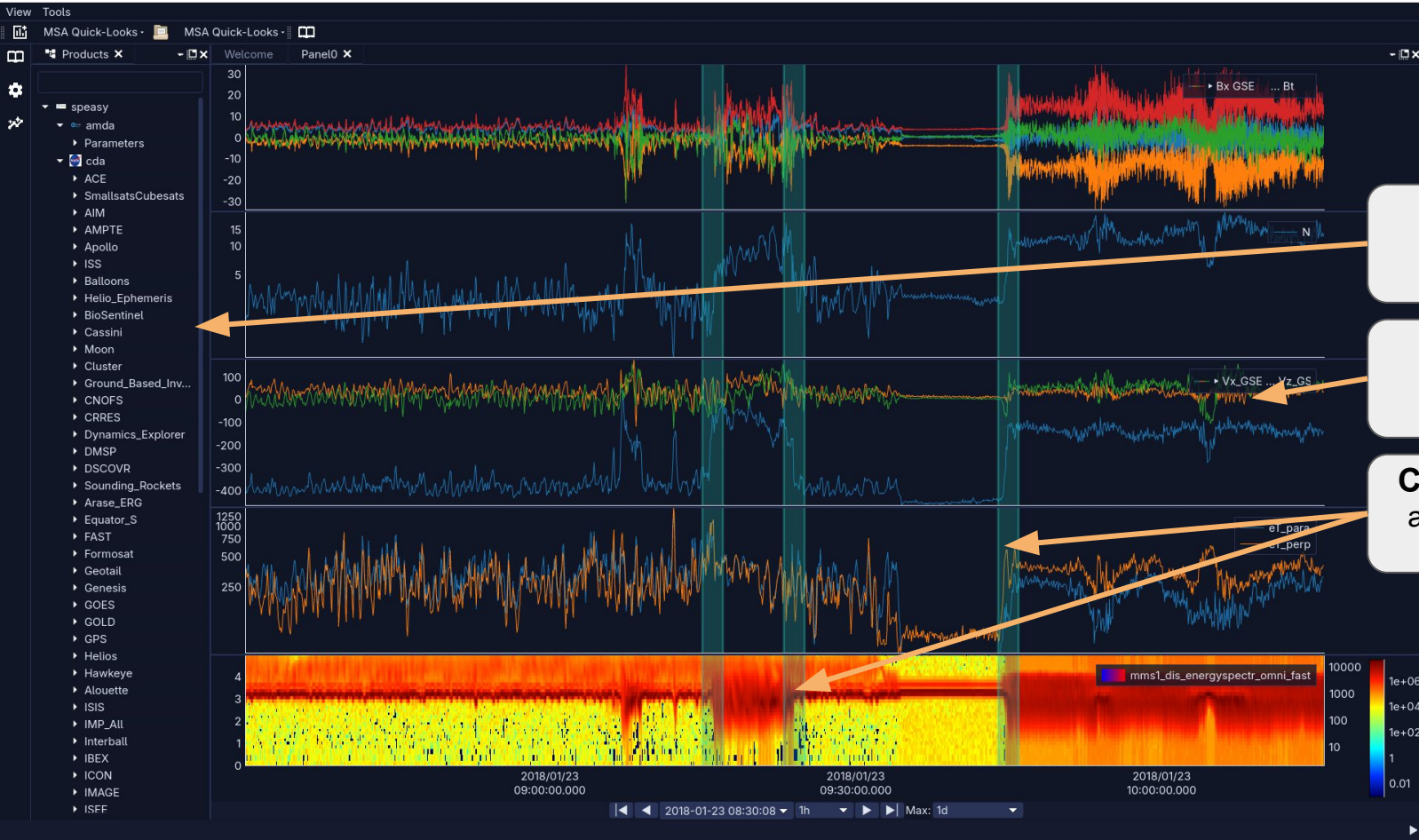
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SciQLop: the Integrated Development Environment (IDE)



Product tree
File explorer for
datasets

Plots panels

Catalogs overlay
annotate and share
events

SciQLop: the Integrated Development Environment (IDE)

```
%%vp --path "mms/mirror"
```

```
def mirror_mode_threshold(start: float, stop: float) -> Scalar["Mirror mode threshold"]:  
    mms1 = spz.inventories.data_tree.cda.MMS.MMS1  
    products = [  
        mms1.DIS.MMS1_FPI_FAST_L2_DIS_MOMS.mms1_dis_temppara_fast,  
        mms1.DIS.MMS1_FPI_FAST_L2_DIS_MOMS.mms1_dis_tempperp_fast,  
        mms1.FGM.MMS1_FGM_SRVY_L2.mms1_fgm_b_gse_srvy_l2,  
        mms1.DIS.MMS1_FPI_FAST_L2_DIS_MOMS.mms1_dis_numberdensity_fast,  
    ]  
    tpara, tperp, b, n = spz.get_data(products, start, stop)  
    if any(v is None for v in (tpara, tperp, b, n)):  
        return None  
  
    b = interpolate(tperp, b)  
    Pperp = tperp * n * 1e6  
    beta_perp = 2 * cst.mu_0 * cst.e * Pperp / (b["Bt"] * 1e-9) ** 2  
    return beta_perp * 1000. * (tperp / tpara - 1)
```


SciQLop: the Integrated Development Environment (IDE)

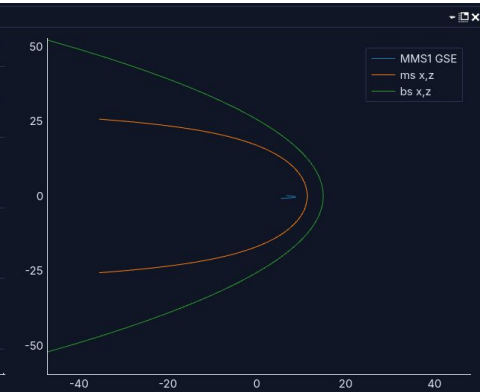
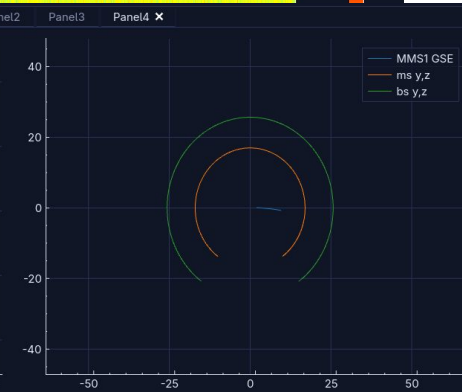
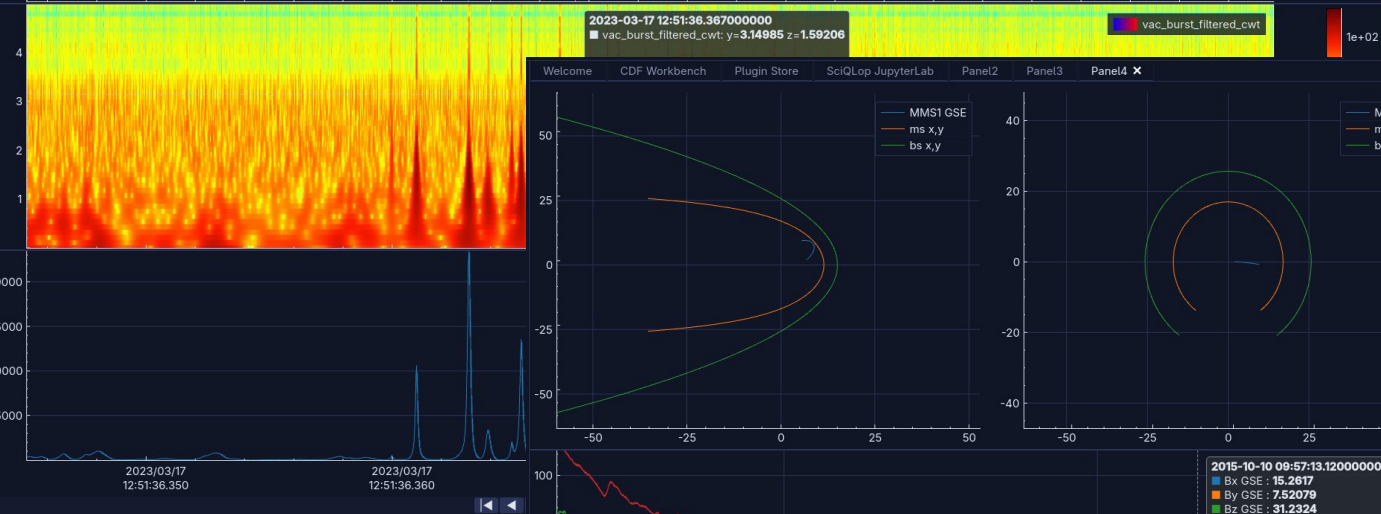
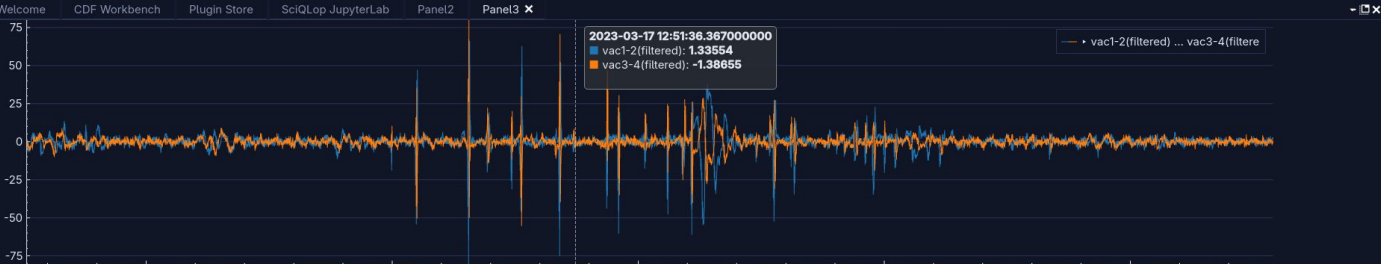
The screenshot displays the SciQLop IDE interface, which is a JupyterLab environment. The interface is divided into several panels:

- Left Panel:** A file explorer showing a project structure with folders like 'dependencies', 'MMS', 'scripts', 'Simple_Virtual_P...', 'Tutorials', 'image.png', 'pyproject.toml', 'uv.lock', 'workspace_spec...', 'workspace.json', 'workspace.json...', and 'workspace.sciqlop'.
- Center Panel:** A code editor showing a Python script named 'Untitled.ipynb'. The script contains several cells of code, including a function definition for 'magnetic_field' and a function for 'magnetic_field2'. The code is written in a dark theme with syntax highlighting.
- Right Panel:** A debug console showing the output of the code execution. It includes a status bar at the top indicating 'VP Debug: magnetic_field' and a plot area showing a time series plot of magnetic field data. The plot has a y-axis ranging from -150,000 to 100,000 and an x-axis showing time from 2020/01/01 to 2020/01/01. The plot is titled 'VP Debug: magnetic_field2'.

Arrows point from the code editor to the debug console, indicating the flow of data and the execution of the code.

JupyterLab
Write your own code

Debug plot
Quick debug
feedback



Plugins

- Calude, OpenCode, Github Copilot agents
 - Create/”see” plot panels
 - Find products
 - Create/modify Jupyter Notebooks
- CDF explorer
 - Plot any variable
- Radio astronomy
 - Curated list of space missions/instruments products
 - Sunpy/Radio Spectra integration
- Bepicolombo MSA
 - Custom panels

Conclusion

<https://github.com/SciQLop>



In v 0.12 now

- 65k+ products
- Workspaces
- JupyterLab
- Built-in DSP functions
- AI assistance
- Plugins
- AppStore
- Catalogs (Local, Remote, *collaborative*)

build for your instrument

- Plugins SDK
- Custom data providers
- Instrument specific panels
- Custom pipelines
- Share on AppStore

On the roadmap

- Workspaces sharing
- More UX features
- More plugins
- Collaborative JupyterLab/UI sessions
- More data sources

open source · MIT/GPL license · supported by CDPP · contributions welcome

Workshop SciQLop, 15-17/09/2026, Jussieu (Paris)



- Talks
 - SciQLop success stories
 - SciQLop use cases
- Hands on

