Programmatic search and analysis using the CSD Python API

CCDC Virtual Workshop Spring 2021 – Session 3

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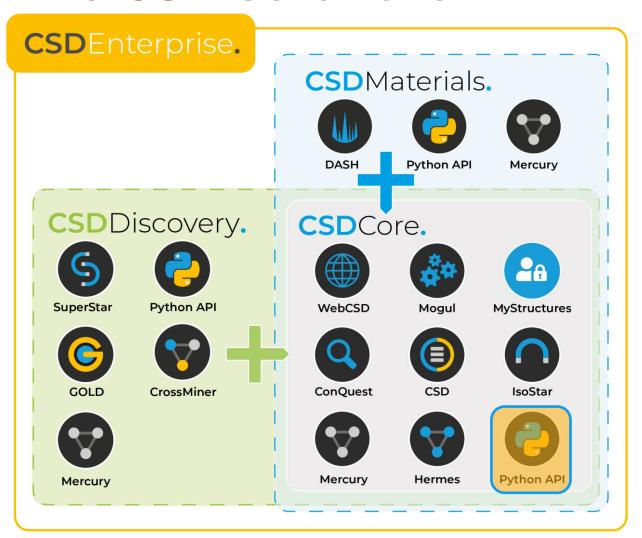


Learning outcomes for today

- Familiarise yourself with how CSD entries are represented in the CSD Python API.
- Learn how to access CSD entries.
- Learn how to read different file formats.
- Learn how to run a search and output results.



The CSD software





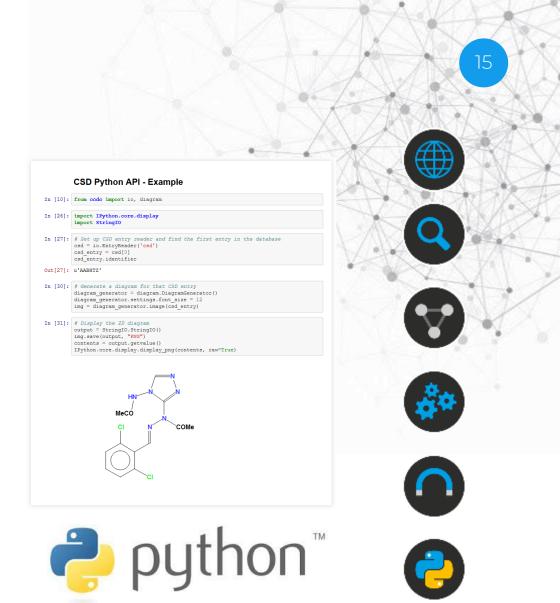




CSD Python API

The CSD-Python API
(Application Programming Interface)

- Enables you to use many capabilities of the CSD-Core without being bound by graphical interfaces
- You can readily create CSD-driven analyses and workflows tailored to your needs
- Our Mercury/Hermes interface allows you to easily access your scripts and run analysis

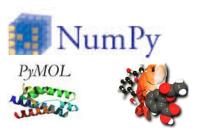




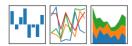
The CSD Python API – Why?

- In CSD-Core:
 - Many things easy to do for one entry/structure, hard/tedious for 100s or 1000s
 - Custom analyses
 - Quick/simple tools and tweaks
- Use with other tools & packages











Benefits of scripting workflows

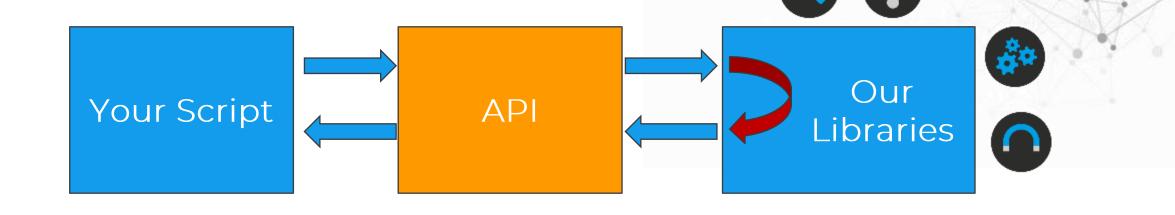


- Automating reporting and analyses saves time
- Results will be consistent and predictable
- Having a scripted workflow makes it easy to share knowledge with colleagues



What is an API?

• <u>Application Programming Interface</u>

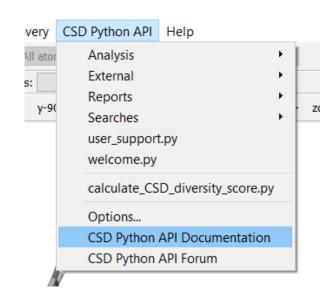


? python™



CSD Python API: Documentation

Documentation is easy to navigate and highly descriptive





Next topic
Conditions of Use
Ouick search

The CSD Python API

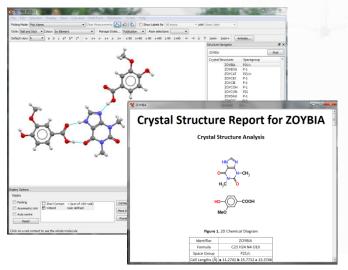
- · Conditions of Use
- Release notes
 - Overview
 - · Citing the CSD Python API
 - Licensed Features
 - Change Log
 - . ^ ^ A

- · Descriptive documentation
 - · Quick primer to using the CSD Python API
 - Using the CSD Python API with Mercury and Hermes
 - · Reading and writing molecules and crystals
 - · Working with entries
 - · Working with crystals
 - Working with molecules, atoms and bonds
 - Editing molecules
 - Search philosophy
 - Text-numeric searching
 - Substructure searching
 - Similarity searching
 - Reduced cell searching
 - Combined searches
 - Conformer generation and molecular minimisation
 - · Field-based virtual screening
 - Working with proteins
 - · Working with cavities
 - Docking and scoring
 - Pharmacophore searching
 - Protein-ligand searching
 - SMARTS implementation
 - Molecular geometry analysis
 - Analysing molecular interactions preferences
 - Interaction Maps
 - Crystal packing similarity
 - · Generating 2D diagrams of molecules
 - Descriptors
 - Graph Sets
 - HBond Propensities
 - · HBond Coordination
 - Morphology
 - Utilities



CSD Python API: Interface

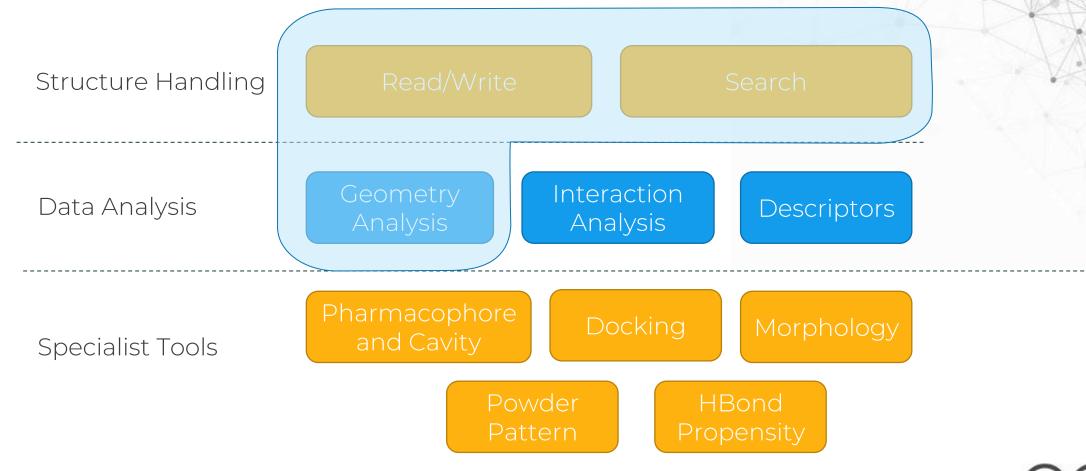
Mercury/ Hermes includes a "CSD Python API" menu



- Allows direct running of built-in or user-generated Scripts from the Mercury graphical user interface
- Scripts menu is designed to work from the currently loaded structure in the visualiser (e.g. CSD entry, CIF or MOL2)

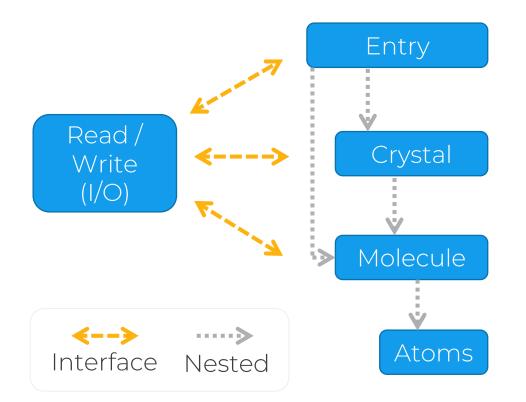


Python API Structure





Structure Handling



Submitted Properties – Habit, Colour, etc. Journal Source Info

Crystallographic Data – Full Structure

Molecular Component of Crystal

Individual Atoms and Their Properties



