

Structural Bioinformatics Training Workshop & Hackathon 2018

3D Visualization in Jupyter Notebooks using MMTF

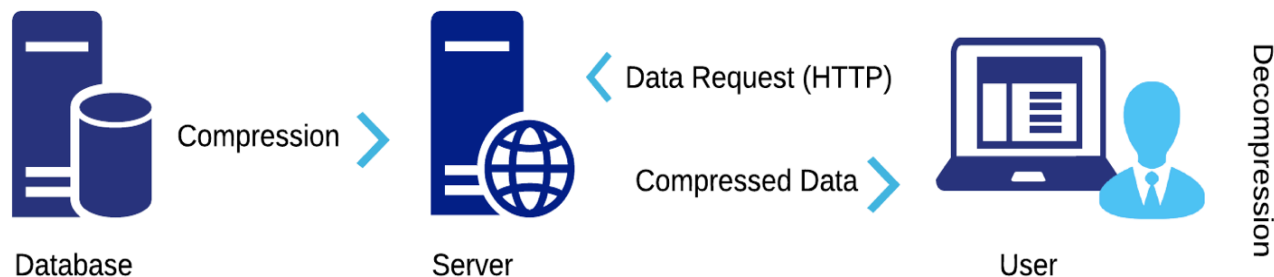
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3D Visualization using MMTF



3D viewer fetches structures using MMTF web services

<http://mmtf.rcsb.org/download.html>

Data are decompressed on the client side using MMTF decoder libraries.

Jupyter Notebook Widgets for 3D Visualization

NGLview

- **Wrapper around NGL**
- WebGL-based JavaScript viewer
- MMTF-JavaScript API
- Subset of NGL features
- Supports visualization of trajectories
- Website: <http://nglviewer.org>
- <https://github.com/arose/nglview>

py3Dmol

- **Wrapper around 3Dmol.js**
- WebGL-based JavaScript viewer
- MMTF-JavaScript API
- Website: <http://3dmol.csb.pitt.edu>
- <https://github.com/3dmol>

Resources

- **NGL**

- Rose AS, et al. (2016) Web-based molecular graphics for large complexes. In Proceedings of the 21st International Conference on Web3D Technology (Web3D '16). ACM, New York, NY, USA, 185-186.
<https://doi.org/10.1145/2945292.2945324>
- Nguyen, H, et al. (2018) NGLview—interactive molecular graphics for Jupyter notebooks, Bioinformatics 34, 1241–1242,
<https://doi.org/10.1093/bioinformatics/btx789>

- **3Dmol.js**

- Rego, N, Koes, D (2015) 3Dmol.js: molecular visualization with WebGL, Bioinformatics 31, 1322–1324, <https://doi.org/10.1093/bioinformatics/btu829>

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