BioDAC: Challenges of Bio-image Processing at Exascale

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Who are we? What do we do?







- Microscopy imaging (commercial and custom-built microscopes)
- Biological samples
- Image processing and analysis

Light sheet microscopy







Light sheet microscopy



Deskewing:





3D + time data of zebrafish (400 GB) Image courtesy of Dr Clare Buckley and Amelia Race

Image processing challenges:

- Deskewing
- Point-spread function (PSF) correction

PSF correction in light sheet microscopy





Point-spread function (PSF)



Geometry of the light beam



- Spatially-varying PSF
- Non-Gaussian PSF
- + Poisson (emission) and Gaussian (detection) noise

Bespoke Image Analysis Solutions

- Acquisition parameters might be suboptimal for analysis
- Voxel size might be incompatible with existing image analysis methods (especially deep learning-based)
- Interdisciplinary communication is HARD



Weigert et al. In Proceedings of the IEEE/CVF WACV. 2020.



Computational requirements



- Optimisation locally on down-sampled.
- Data transfer is a bottleneck Globus.

Gibbs et al. Front Cell Dev Biol, 9, 2021, 10.3389/fcell.2021.739079

Computational requirements and the Exascale Test Bed

- Reaching memory limits in processing pipeline.
 - Use high-memory nodes, or:
 - Code redesign for distributed memory system

- To improve I/O performance we can use the Exascale Test Bed:
 - o 1PB NVMe storage
 - Access to CSD3's compute
 - o 500GB/s bandwidth

https://excalibur.ac.uk/projects/exascale-data-testbed



Conclusions

- Light sheet microscopy enables the acquisition of high-resolution 3D+time images
- Complex image processing algorithms and customised image analysis pipelines are often necessary to answer the relevant scientific question
- Multiple levels (PC to cluster) of advanced hardware and software are required to store, transfer, and fully process these images
- Improved communication with the biologist teams might help alleviate some of the challenges

Thank you for your attention!