# Needs for Diverse Translational Biomedical Interoperable Computational Tools

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2023-12-05

# The Team!













































































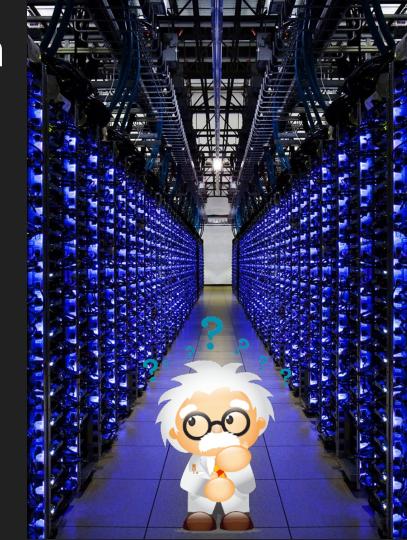






# Foundational Problem

How can a scientist harness petabytes of data to derive quantitative measurements and gain meaningful insights?







# Polus Platform - Open Source Adobe Creative Cloud... for Science



#### **Constant Processing Time For Increasing Data Sizes**

• Web-based solution to access cloud/server/ HPC computational and storage resources



#### Collaborative, Visual, and Interactive

· Web based UI; Simple point-click interface



#### **Traceable and Reproducible Science**

 Environment, packages, functions, etc. all recorded automatically; automated hyperlinked provenance to raw data



#### **Analytical**

 Validated measurements/algorithms; Supports latest in machine learning, deep learning, and statistics



#### Flexible and Modular

 Open source Plugins/Applications/Codebase; Interoperable container-based. Shared Metadata, Storage, and AuthN/AuthZ



#### Secure

Private research, patient data/PII, and IP is securely and safely stored





# Diverse Biomedical Data & Tooling

#### Began with Imaging

- Microscopy light, fluorescent, electron, pathology, high content screening
- Clinical Fundus images (RGB, fluorescent), X-Rays, CTs, MRIs, OCTs, etc.

#### **Expanded Out!**

 Cheminformatics, molecular modeling, real world evidence (electronic health records, insurance/payer claims, mortality, etc.)

#### On the Horizon

 Omics - genomics, transcriptomics, proteomics, metabolomics, etc.





## **Diverse Tool Needs**

#### Language Support

Python, Java, R, C/C++, and Rust

#### Diverse Libraries

- Traditional, Tensorflow, Pytorch, Jax, etc.
- Diverse driver requirements i.e. CUDA

#### **Diverse Hardware**

- CPU x86 and ARM
  - AVX/AVX2
- GPU Nvidia
  - Historical code written across multiple Nvidia compute capabilities
- RAM 8GB  $\rightarrow \infty$



**OPyTorch** 











### Diverse Users Needs

#### **Programmers**

SDKs and libraries

#### Technical

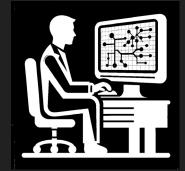
- Command line tools & terminal
- Direct edit YAML/JSON

#### Advanced Domain Experts

- **Complex GUIs**
- Configure/optimize hyperparameters
- Some "Macro" scripting

#### **Domain Experts**

- Simple GUIs
- Single button if possible!











# Diverse Data Sizes and Scaling Needs

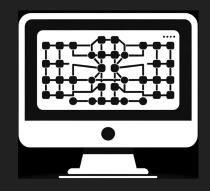
#### **Dataset Scales Processed**

- < 1GB Text and image data</li>
- > 100s of TBs Imaging

#### Scaling Paradigms

- Split on data
  - Wells of plates in HCS
  - Molecules binding to proteins
- Scale on computation
  - Molecular modeling
  - Feature extraction
  - EHR/Claims extraction, transformation, loading, filtering, modeling









# Interoperable Analysis Tools

#### 100+ Scalable Interoperable Tools!

• Background correction, Table Merge, etc.

#### Deep Learning Training/Inference Models

- Point/click interface for training & inference
- Over 400+ classification networks
- Over 4200+ segmentation networks (Video→)

#### **Custom Deep Learning Models**

- Image segmentation models MESMER, CellPose, Spline/Star-Dist, etc.
- LLMs Llama 2s, Mistral, Falcon, etc.
- Chemi-informatic MolGAN, diffDock, etc.
- MM OnionNet-pose, AlphaFold-2, BioExcel/BioBB, etc.

#### Trained/Implemented

• 4000+ image segmentation models on diverse images



# Questions?