DEVELOPING DISTRIBUTED HIGH-PERFORMANCE COMPUTING CAPABILITIES OF AN OPEN SCIENCE PLATFORM FOR ROBUST EPIDEMIC ANALYSIS (OSPREY)

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OUTLINE

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- Data Streams as a Service (DSaaS)
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WHY OSPREY?
67 Days to Lockdown

From Chicago’s first known COVID-19 case to the governor’s historic shelter-in-place order, here’s how those critical, frenzied early weeks of the pandemic unfolded, in the words of those at the center of the crisis.


MONDAY, MARCH 9

The State’s Disaster Proclamation

One day after a Cook County man was announced as Illinois’s seventh case and its first known instance of “community spread” — the spread of a disease to people who have had no known contact with other infected people — Pritzker issued a disaster proclamation that marshaled state resources to combat the virus. Four additional cases were revealed this day, all of them in Cook County. The next day saw eight more cases, including the first two outside Cook, bringing the state total to 19. The numbers were rising fast now, just as experts had predicted.

Argonne National Lab built what’s called an agent-based model specifically for Chicago that predated COVID and mimics the movement of its 2.7 million residents. All the gyms, schools, and businesses are built in, and it [takes into account] how people interact throughout the day. So pretty early on, we connected with them to start thinking about using that Chicago model for COVID.
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WHY OSPREY?

- During COVID, individual research groups generally worked independently trying to exploit advances in HPC, data management, ML/AI, and automation methods to develop, calibrate, modify, verify, and validate their epidemiologic models.
  - Required large amounts of heroic, overlapping work
  - Risked lacking robustness, security, scalability, or efficiency
- Heterogeneous, changing, and incomplete data required complex integration across diverse and novel surveillance signals
  - Created significant challenges for use in epidemiologic workflows
- The Open Science Platform for Robust Epidemic analYsis (OSPREY) seeks to lower the barriers to and automate epidemiologic model analyses on HPC and cloud resources
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PROTOTYPIC OSPREY ARCHITECTURE
The OSPREY architecture includes:
- Model exploration module
- FaaS task distribution
- EMEWS task DB
- Scalable worker pools
- Data sharing

These collectively provide the ability to robustly and securely coordinate (epidemic) workflows across a distributed ecosystem of heterogeneous remote resources.

We developed a task API, including remote worker and service monitoring and coordination.

Globus Compute provides:

- Ability to Start/stop EMEWS Service/DB on remote machine
- (with PSI/J) Ability to provision, monitor, tear down worker pools
- Secure, long running access to EMEWS Service/DB

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DATA STREAM AS A SERVICE (DSaaS)
DATA STREAM AS A SERVICE (DSaaS) OVERVIEW

Users specify data sources, provide validation and modification functions and a timer for regular updates from the data source.

A Globus timer event triggers a Globus flow at the specified interval to fetch, validate and modify the data and store it into a Globus Connect Server.

Analysis workflows (using e.g., Parsl, Swift/T, Globus Compute) can search and retrieve data and store analysis artefacts in DSaaS all via Globus Transfers.
USING DSAAS IN PARSL FUNCTIONS

- Source data should be added to the service outside of Parsl via the CLI

  dsaas -create_source -n 'covid19' -u
  https://data.cityofchicago.org/resource/naz8-j4nc.json -t 86400 -v
  'df2ab7fc-dad7-440f-a9cb-e02dbeac4141'

- Data can be accessed within a function using the DSaaS client API (WIP)

  @python_app
  def analysis(source_id=1):
      from dsaas.client import source_file

      df = source_file(source_id=1) # returns pandas dataframe. Transfer via Globus Transfer ...

- Future work: analysis outputs can be stored within DSaaS

  from dsaas.client import save_analysis

  save_analysis(df, name='parslfest-analysis.json', version=2)
THANK YOU FOR YOUR ATTENTION! QUESTIONS?

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