# **Biosciences Working Group**

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PRAGMA 31, Sept 9, 2016

#### **Bioscience Activities**

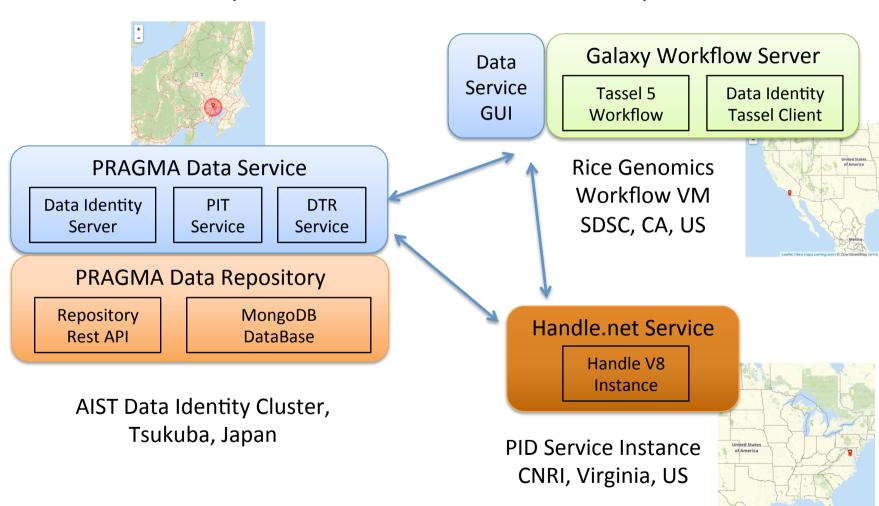
- Genomics Analysis Pipeline
  - Creating resources to enable genomics analysis
- PRAGMA Data Identity Service
  - Rice genomics analysis
- Both driving applications of PRAGMA Cloud
  - Cross-working group interaction
- DOCK in Docker
- Condor application for BLAST

#### Since PRAGMA30

- Genomics Analysis Pipeline
  - Installation of rice genomics analysis tools such as TASSEL5 (Venice, Ramil)
  - Additional testing of VM (Venice, Gabriel)
  - VM is now accessible via pragmaboot and scheduler (Shava, Nadya)
  - AIST PRAGMA Cloud resources prepared

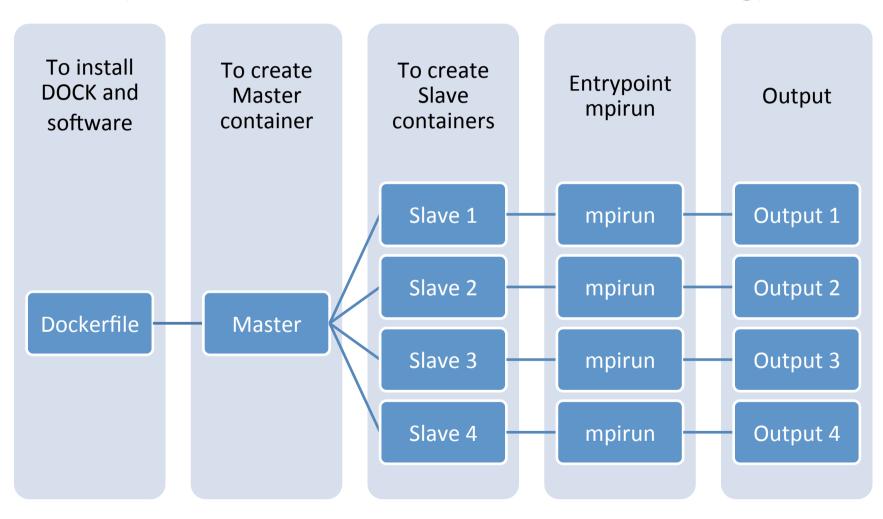
#### PRAGMA Data Service

(Gabriel, Ramil, Jason, Beth)



#### **DOCK** in Docker Containers

(Max, Thomas, Jason, Ichikawa, Weicheng)



### DOCK in Docker Containers (cont.)

- Dockerfile To install DOCK and software; generate a dedicated image for DOCK
- Entrypoint File that consists of all actions that will be executed on container
- Compose A tool for defining and running multi-container Docker applications

Performance Evaluation Total elapsed time (sec)

Number of containers						
	0	2	4	8	16	32
mpirun np 2	3010	3018				
mpirun np 4			985			
mpirun np 8			408	416		
mpirun np 16			211	216	215	

# Condor Computing Application (Tommy Lam, HKU)

- Utilize off-hour computers in library and computer labs in HKU campus
- Demonstration of Green Computing
- We are trying to adopt BLAST for condor system, e.g.
   BLASTer (http://funnelfiasco.com/research/gca13.pdf)
- BLAST is highly parallelizable
  - If you have many query sequences to search against a database, query sequences can be split into different condor nodes for individual BLAST search(easy; BLASTer way)
  - If you have only few query sequences but big database, splitting database into pieces for BLAST is a good option. But this will require MPI or endpoint summary computing of result

# Bioscience WG – Day 1

- Continue collaboration with IRRI
- Presentations by Biotec
  - Thailand rice genomics
  - Interest in PRAGMA data service
    - Will follow-up
  - Data is closed implications for data service
- DOCK in Docker
  - Work toward a stable service to demo at PRAGMA32

# Bioscience WG – Day 2

- Virtual screening
  - Data services for results
    - 10 million compound screen
      - 10 million DOs or single DO?
    - Harvesting input, output, VM specifications
- JHPCN International Grant (due 1/17)
  - Arry, Heru, Jason, Susumu
  - Centered around virtual screening activity
  - PyRx virtual screening tool
    - Enable the use of PRAGMA resources