# Biosciences WG Updates

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### Since PRAGMA28

- Regular conference calls
- Two areas of activity: genomics and proteomics
  - Genomics Analysis Pipeline
    - Established a roadmap (Tommy)
    - Selected Biolinux (Jason/Tommy)
    - Created KVM-based VM with Biolinux (Kohei)
    - Preliminary testing of VM (Tommy)
  - Natural Products Discovery Portal
    - Database refinement/cleanup (Arry)
    - VietHerb database (Ly)

# Genomics Analysis Pipeline

- Infectious disease as domain
- Regular Skype conference calls
- Draft a "roadmap" (Tommy)
- Action items
  - Select and install genomics tools (Tommy, Jason)
    - Galaxy, iPlant
  - Influenza A virus and MRSA as small test for assembly and annotation (Ly)
    - Define file size, tools used, etc.
  - Hadoop support (Andrea)

#### **Genomics Analysis Pipeline in PRAGMA**



#### 16S Metagenomics

QC, taxonomic classification and composition analysis of NGS bacterial 16S amplicon samples.

Results feedback to the web-portal

#### Viral Metagenomics

QC, genome assembly, taxonomic classification and composition analysis of NGS viral samples.

### Pre-built workflows for common types of analysis

Molecular epidemiology

Sequence alignment, phylogeny inference, molecular clock dating of viral gene sequences.

#### Gene and protein structure

glycosylation site prediction, epitope prediction, electrostatic potential

More custom workflows...



Phase 1: Set up web-portal and computing infrastructure for Galaxy and genomics tools in PRAGMA resources.



Phase 2: Build workflows for common types of genomic analysis, initially for field of infectious disease research and then extend to other fields with the contributions of other PRAGMA participants.



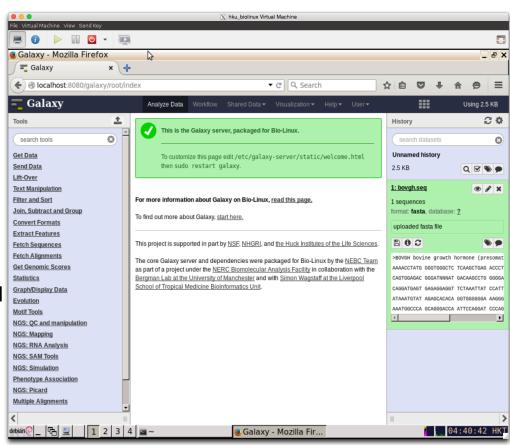
Phase 3: Testing using case studies contributed by PRAGMA participants.



Phase 4: Education via training workshops and invite wider contributions to building common analysis workflows and application with PRAGMA participants' data.

### PHASE 1: Installation of GALAXY in VM

- GALAXY was successfully installed in the PRAGMA-compatible VM
  - with Biolinux8
  - with window manager icewm
- But this GALAXY inside the VM is supposed to be isolated from the world outside the VM, so we
  - need to create a central web-portal to accept computational jobs
  - need to allow connection between VM-inside GALAXY and the web-portal GALAXY
- Other tasks in PHASE 1
  - Identify minimum package of Biolinux8 to reduce transfer load.
  - [Do you have any ideas?]



# PRAGMA Natural Products Discovery Platform

- Components
  - 1. Natural products database
  - 2. Virtual screening
  - Biodiversity mapping
- What are the infrastructure requirements for each component?
  - 1. Storage, data format
  - 2. Network, compute nodes
  - 3. Mapping tools, visualization



# Natural Products Discovery Platform

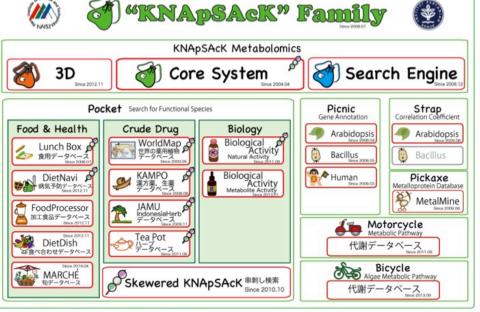
- Regular Skype conference calls
- Action Items
  - Define ideal computing environment, applications, databases, size of VM (Arry, Ly)
  - Continued discussion with resources WG to create web services (Nadya, Ari)
    - Based around NBCR efforts e.g. PyRx
  - Start combining the natural products databases i.e. remove duplicates and combine (Arry, Kanaya, Ly)
- Benefit Vietnam (limited computational resources), Indonesia (move to virtualized system)



## **KNApSAcK (Japan)**

## **HerbalDB** (Indonesia)

http://kanaya.naist.jp/KNApSAcK\_Family/

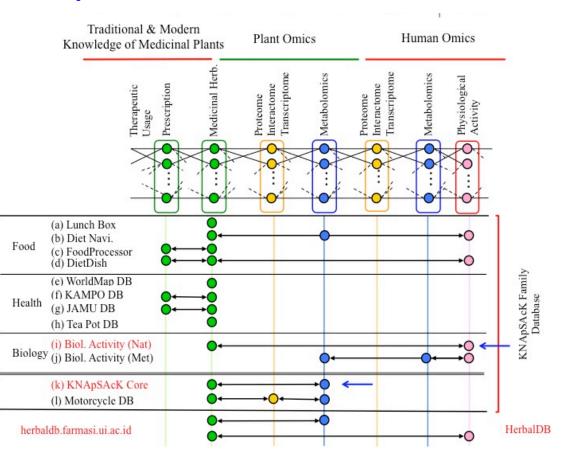


Instruction Manual(English)

Instruction Manual(Japanese)

## HerbalDB & KNApSAcK intersection

 To find possibility of data sharing between HerbalDB and KNApSAcK DB



### VietHerb Database

- Vietnam has an extremely rich sources of medicinal plant
- Detailed information about phytochemicals has not been standardized and digitized
- This database inspired by KNApSAcK database system (Japan) and Jamu database system (Indonesia)
- Create an interactive, electronic herbal database
  - Impartial, evidence-based information resource
  - Public site provides free access
  - Use of herbs for health