2018-04-23 - ISDA Team - Stand-up Meeting Notes

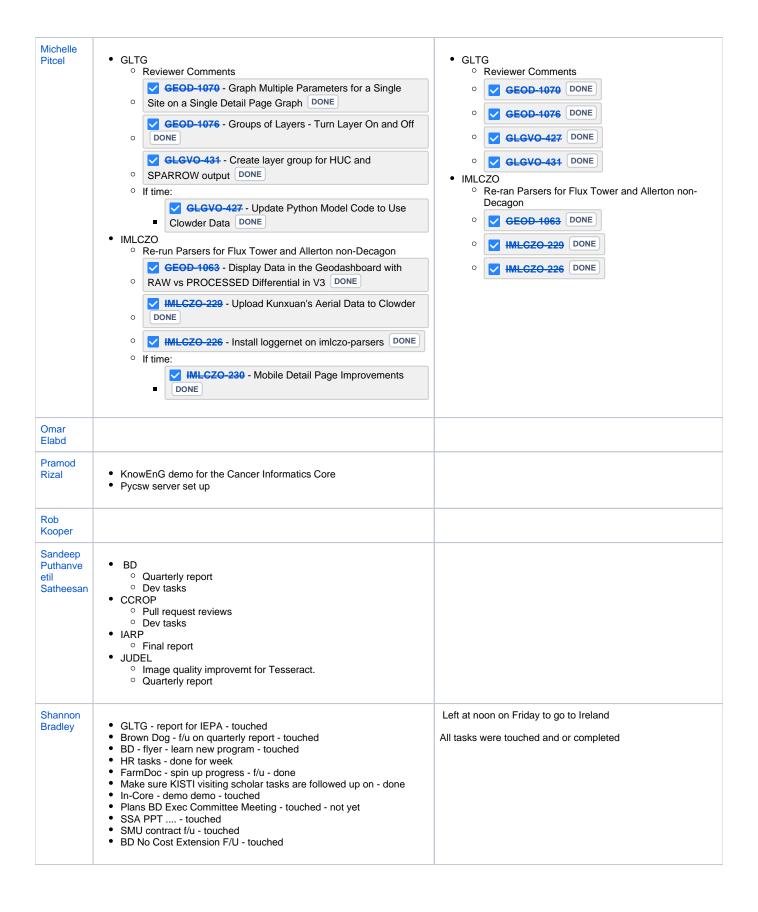
Did you know: The 100 folds in a chef's toque are said to represent 100 ways to cook an egg.

Have a great week -

Who	Planned - Monday	Accomplished - Friday
Bing Zhang		PR In Review design and implement SimpleExtractor in Pyclowder2. 1. wordcount-simepleextractor 2. meangrey-simpleextractor. Done update greenroute with feature to set step size in meter unit, discuss with students on the format of uploading test files. update greenroute on clowder dev. deploy clusterscale on dev swarm cluster with dockercompose file. Daily maintenance on Browndog.
Benjamin Galewsky	Prep for Globus World Collaborate on TACC Cyberinfrastructure for Archives proposal. Attend Globus World	
Chen Wang	Incore Data browser and Fragility Browser fix bug + styling SMM change citation and email address in the app hide boolean function add MIME type when uploading files to s3 bucket and clowder design clowder interface in SMM	Incore

FarmDoc Review matible code Review Review			
### Planning TERRA WholeTale	er	 Review matlab code Create VMs for shared architecture with Cover Crop CoverCrop Code review Help updateworkflow with template parsing tool IN-Core v2 demo Code review General Finish zenodo entry and initial password reset page for 	
1. Start converting Water Damage Analysis to pyincore 2. Create initial farmdox web application FD-15		PlanningTERRAWholeTaleGodaddy API integration	TERRA
InCore	Naraharis	Create initial farmdoc web application FD-15	converting Water Facility Damage. 2. Made progress on farmdoc webapp setup. Setup flask and folder structure.
## Help in preparation for Demo GLM # Start working on visualizations for Fish Data SMU # Follow up on extractors not running after update # KnowEng Add logic for supporting mixed data type in phenotype expander SAP: Add column/row duplication check Add gene order check Add remove NA index to signature data and user spreadsheet data Update README Explore solution for jupyterhub user data backup in AWS InCore Finish adding new multiprocessing library to buildingdamage.py Benchmark testing on parallelization of buildingdamage.py Moved mapping_id from constructor to class function Started 2.6.0 release # KnowEng Added logic for supporting mixed data type in phenotype expander SAP: Added column/row duplication check Added gene order check Added gene order check Added gene order check Added gene order check Added related unit tests Update README Looking into solution for jupyterhub user data backup in AWS InCore Finished adding new multiprocessing library to buildingdamage.py Moved mapping_id from constructor to class function Started benchmarking testing on parallelization of buildingdamage.py, need clarification on incore2-test machine resource.	Khine		
 KnowEng Add logic for supporting mixed data type in phenotype expander SAP:	Gutierrez	 Help in preparation for Demo GLM Start working on visualizations for Fish Data SMU 	 Wrapped up the development of JupyterLab_Incore. Updates in the data viewer, added a creator filter, previews for xml and text files, added a dataset metadata component GLM Updates to pull request on filtering by date on the trends by region detail page Started 2.6.0 release
Jong Lee	Jing Ge	 Add logic for supporting mixed data type in phenotype expander SAP: Add column/row duplication check Add gene order check Add remove NA index to signature data and user spreadsheet data Update README Explore solution for jupyterhub user data backup in AWS InCore Finish adding new multiprocessing library to buildingdamage.py 	 Added logic for supporting mixed data type in phenotype expander SAP: Added column/row duplication check Added gene order check Added remove NA index to signature data and user spreadsheet data Added related unit tests Updated README Looking into solution for jupyterhub user data backup in AWS InCore Finished adding new multiprocessing library to buildingdamage.py Moved mapping_id from constructor to class function Started benchmarking testing on parallelization of buildingdamage.py, need clarification on incore2-
	Jong Lee		

Kenton McHenry	NASA meetings	NASA meetingsCisHR
Luigi Marini	BD Clowder 1.4 release and 2.0 development Report section Industry Documentation and planning GLM, GLTG, IMLCZO Pull requests and release HR	 Last minute meetings and HR. Cyprus Clowder instance support. Clowder landing page.
Marcus Slavenas	 gltg script for iepa sensor name usgs parser deployed release? iepa -review comments vbd changes to app for model year selection 	gltg iepa -review comments usgs parsers deployed updated iepa sensors on dev vbd fix graphs for new 2017 data created data endpoints for dates, weather, and forecaste
Maxwell Burnette		
Michal Ondrejcek	vacation We-Fri MDF script for FF pipeline Farmdoc finish FD-13 - Jira project doesn't exist or you don't have permission to view it. , start FD-14 - Jira project doesn't exist or you don't have permission to view it.	done MDF Script for FF pipeline Farmdoc initial PR before departure
Sara Lambert	 NDS Attempt to reprovision the public Workbench beta (again) Planning next steps (upcoming demos, releases, etc) Crops in Silico Add the real models to UI to prepare for spring demo Update UI to support any new features/quirks in the real models Plan the spring demo of the UI, write slides and a script Industry Get the proxy endpoint working Ask about next steps / backlog? 	NDS Provisioning process is once again working, but I will not have cycles to perform the necessary user migration for another week or so Crops in Silico Added real models to the UI Added new editable fields on inport/outports: read_meth, write_meth Added new editable fields on edges: field_names, field_units Implemented submitting a new GitHub issue via Google Forms Industry Basic proxy is functioning, but it cannot yet handle large responses



Yan Zhao		Monday off BD MRI extractor, some problem with memory, not finish BD report GLM test the time for binning, seems a lot of work, maybe has no time to do that. CC fix private route with new router lib. – not working.
Yong Wook Kim	 Update dataType in data serivce with new mapping Update jupyterlab mapping preparation Update jupyterlab docker Update extractors-geo to use pyclowder 2 	 Worked on updating pycsw extract to use pyclowder 2 Tested pycsw clowder dockers Updated incore jupyterlab docker Updated incore kubernetes