2018-04-09 ISDA Team - Stand-up Meeting Notes

Look out - Shannon's Baaaaaack!

If you have the feeling you've experienced an event before in real life, call it déjà vu.

If you feel like you've previously experienced an event in a dream instead, there's a different term for it: déjà rêvé.

Who	Planned - Monday	Accomplished - Friday
Bing Zhang		
Benjamin Galewsky		
Chen Wang	Science code: wrap up Fuyu's code; pass 2nd round edits to Xian Pyincore: add data service test and fix pyincore according to failed cases Semantic: meet with Omar to start implementing Other tasks TBD SMM faculty trolling project: prettify the frontend and add UI components meetings Clowder related tasks (if have time)	Science code: wrap up Fuyu's code; pass 2nd round edits to Xian Pyincore: add data service test and fix pyincore according to failed cases Fragility browser on jupyterlab SMM faculty trolling project: prettify the frontend and add UI components
Christopher Navarro	Cover Crop Project meeting/web design review Update covercrop VM with new genotype for Rye Code review FarmDoc Document migration steps for moving covercrop postgres /geoserver to fd-geoserver IN-Core Continue working on moving building damage analyses to pyincore Work on utility to create result dataset Code review	
Craig Willis Gowtham Narahariset ty	INCORE Implement PUT test cases for data service Start moving Water Network Analysis code from v1 to python Farm doc Setup local dev environment. Run Soil Converter locally Migrate postgres tables to the new farmdoc VM	
Htut Khine Htay Win	Vacation Man Tun Sink Word Fri	Vacation Man. Tup. Sick Word. Eri
Indira Gutierrez Polo	Vacation Mon- Tue, Sick Wed - Fri	Vacation Mon- Tue, Sick Wed - Fri

Jing Ge	 KnowEng Various changes to imputation function in Data_Cleanup_Pipeline Allowing replace NA value with customized value when write to file Testing for output reproducebility for samples_clustering_pipeline and general_clustering_pipeline Add intersection check for Feature_Prioritization_Pipeline & refactor the common code between Gene_Prioritization_Pipeline and Feature_Prioritization_Pipeline Update Dockerfile in all repositories to sync on latest base_image and maintainer list Build docker image for data_cleanup_pipeline and signature_analysis_pipeline Investigate issue with input file cannot be correctly read by pandas In-Core review pull request 	 KnowEng Finished several changes to imputation function in Data_Cleanup_Pipeline Removed impute_na from Geneset_Characterization_Pipeline Added impute_na to Gene_Prioritization_Pipeline and Feature_Prioritization_Pipeline Added default option of imputation to be reject Added new tests Added option to allow replacing NA value with any customized characters. Tested samples_clustering_pipeline and general_clustering_pipeline for reproducebility issue and reported findings. Added intersection check for Feature_Prioritization_Pipeline & refactor the common code between Gene_Prioritization_Pipeline and Feature_Prioritization_Pipeline Update Dockerfile in all repositories to sync on latest base_image and maintainer lists Built docker image for data_cleanup_pipeline and signature_analysis_pipeline Investigated the issue that input data was altered without any changes and found that duplicate columns was causing this. Need more investigation. In-Core Reviewed new building_damage_analysis.py
Jong Lee		
Kenton McHenry	CSSI proposals	CSSI proposals
Luigi Marini	 CSSI proposal Clowder documentation Clowder 1.4 release and PR reviews Syngenta extractors Geodashboard PR reviews BD PR rewiews Vacation Friday 	CSSI proposal Clowder documentation Clowder 1.4 release and PR reviews BD PR rewiews Vacation Friday Created Cyprus Clowder instance Created IMLCZO parsers VM
Marcus Slavenas	 gltg ingesting data vbd parsing mosquito data running model 	
Maxwell Burnette	laser 3d format conversionpipeline health monitoring	
Michal Ondrejcek	MDF	MDF

Sara Lambert	NDS / TERRA Keep trying to scale up the TERRA Workbench Maintenance where needed KnowEnG Last week on the project Kubernetes Demo on Wednesday NIH Demo on Friday Crops in Silico Continue collecting feedback and iterating until satisfied	NDS / TERRA Failed attempts trying to scale up the TERRA Workbench Minor Workbench maintenance KnowEnG NIH Demo cancelled due to scheduling conflicts Wrap up final bits of Kubernetes for their stuff Crops in Silico Continue collecting feedback and iterating until satisfied
Michelle Pitcel	 GLTG	GLTG
Omar Elabd		
Pramod Rizal	 Industry - Pix4D Extractor (W.I.P.) k8s dev setup/test for KnowEnG (W.I.P.) 	Industry - Pix4D Extractor (W.I.P.) k8s dev setup/test for KnowEnG (W.I.P.)
Rob Kooper	 LSST operations requirement gathering parsl PEcAn htcondor meeting create RC VM Industry check on progress of new hardware extractors per space TERRA quarterly report NCSA/ISDA PEARC reviews 	
Sandeep Puthanveeti I Satheesan	BD Sprint tasks BD-CLI update Code review CCROP Tool to generate experiment file from template based on user input Other tasks JUDEL Apply Tesseract for OCR and newspaper article segmentation	BD Worked on BD-CLI update CCROP Completed development for tool to generate experiment file from template based on user input JUDEL Working on applying Tesseract for OCR and newspaper article segmentation.
Shannon Bradley	 Even tho I read a lot of email while I was out - I still have email to catch up on Sprint Updates Brown Dog Qtr 1 2018 report - kickoff Get back in a groove and catch up on everything else new tablet? 	

Yan Zhao		BD upgrade to pyclowder2 for meangrey check the silhouette for clusterman – failed to get an answer fix pecan shiny GLM ingest USGS 04024000 on dev. add helper text for USGS quality code CC new UI initial implement for user session service.
Yong Wook Kim	 Finish pycsw extractor by adding geotiff function Update docker for pyincore/opensees incore jupyterhub docker Update kubernetes with pyincore jupyterlab Prepare the COE demo 	 Finished pycsw extractor and tested it with docker and clowder Updated docker for pyincore, opensees, and other jupyterlab extension Updated kubernetes with new jupyter lab Tested web mapping on the jupyterlab