

# 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
<a href="#">Bing Zhang</a>		
<a href="#">Benjamin Galewsky</a>		
<a href="#">Chen Wang</a>	<p>INCROE</p> <ul style="list-style-type: none"><li>Science code: wrap up Fuyu's code; pass 2nd round edits to Xian</li><li>Pyincore: add data service test and fix pyincore according to failed cases</li><li>Semantic: meet with Omar to start implementing</li><li>Other tasks TBD</li></ul> <p>SMM</p> <ul style="list-style-type: none"><li>faculty trolling project: prettify the frontend and add UI components</li><li>meetings</li><li>Clowder related tasks (if have time)</li></ul>	<p>INCROE</p> <ul style="list-style-type: none"><li>Science code: wrap up Fuyu's code; pass 2nd round edits to Xian</li><li>Pyincore: add data service test and fix pyincore according to failed cases</li><li>Fragility browser on jupyterlab</li></ul> <p>SMM</p> <ul style="list-style-type: none"><li>faculty trolling project: prettify the frontend and add UI components</li></ul>
<a href="#">Christopher Navarro</a>	<ul style="list-style-type: none"><li>Cover Crop<ul style="list-style-type: none"><li>Project meeting/web design review</li><li>Update covercrop VM with new genotype for Rye</li><li>Code review</li></ul></li><li>FarmDoc<ul style="list-style-type: none"><li>Document migration steps for moving covercrop postgres/geoserver to fd-geoserver</li></ul></li><li>IN-Core<ul style="list-style-type: none"><li>Continue working on moving building damage analyses to pyincore</li><li>Work on utility to create result dataset</li><li>Code review</li></ul></li></ul>	
<a href="#">Craig Willis</a>		
<a href="#">Gowtham Naraharisetty</a>	<p>INCORE</p> <ul style="list-style-type: none"><li>Implement PUT test cases for data service</li><li>Start moving Water Network Analysis code from v1 to python</li></ul> <p>Farm doc</p> <ul style="list-style-type: none"><li>Setup local dev environment. Run Soil Converter locally</li><li>Migrate postgres tables to the new farmdoc VM</li></ul>	
<a href="#">Htut Khine Htay Win</a>		
<a href="#">Indira Gutierrez Polo</a>	Vacation Mon- Tue, Sick Wed - Fri	Vacation Mon- Tue, Sick Wed - Fri

Jing Ge	<ul style="list-style-type: none"> <li>KnowEng <ul style="list-style-type: none"> <li>Various changes to imputation function in Data_Cleanup_Pipeline</li> <li>Allowing replace NA value with customized value when write to file</li> <li>Testing for output reproducibility for samples_clustering_pipeline and general_clustering_pipeline</li> <li>Add intersection check for Feature_Prioritization_Pipeline &amp; refactor the common code between Gene_Prioritization_Pipeline and Feature_Prioritization_Pipeline</li> <li>Update Dockerfile in all repositories to sync on latest base_image and maintainer list</li> <li>Build docker image for data_cleanup_pipeline and signature_analysis_pipeline</li> <li>Investigate issue with input file cannot be correctly read by pandas</li> </ul> </li> <li>In-Core <ul style="list-style-type: none"> <li>review pull request</li> </ul> </li> </ul>	<ul style="list-style-type: none"> <li>KnowEng <ul style="list-style-type: none"> <li>Finished several changes to imputation function in Data_Cleanup_Pipeline <ul style="list-style-type: none"> <li>Removed impute_na from Geneset_Characterization_Pipeline</li> <li>Added impute_na to Gene_Prioritization_Pipeline and Feature_Prioritization_Pipeline</li> <li>Added default option of imputation to be reject</li> <li>Added new tests</li> </ul> </li> <li>Added option to allow replacing NA value with any customized characters.</li> <li>Tested samples_clustering_pipeline and general_clustering_pipeline for reproducibility issue and reported findings.</li> <li>Added intersection check for Feature_Prioritization_Pipeline &amp; refactor the common code between Gene_Prioritization_Pipeline and Feature_Prioritization_Pipeline</li> <li>Update Dockerfile in all repositories to sync on latest base_image and maintainer lists</li> <li>Built docker image for data_cleanup_pipeline and signature_analysis_pipeline</li> <li>Investigated the issue that input data was altered without any changes and found that duplicate columns was causing this. Need more investigation.</li> </ul> </li> <li>In-Core <ul style="list-style-type: none"> <li>Reviewed new building_damage_analysis.py</li> </ul> </li> </ul>
Jong Lee		
Kenton McHenry	<ul style="list-style-type: none"> <li>CSSI proposals</li> </ul>	<ul style="list-style-type: none"> <li>CSSI proposals</li> </ul>
Luigi Marini	<ul style="list-style-type: none"> <li>CSSI proposal</li> <li>Clowder documentation</li> <li>Clowder 1.4 release and PR reviews</li> <li>Syngenta extractors</li> <li>Geodashboard PR reviews</li> <li>BD PR reviews</li> <li>Vacation Friday</li> </ul>	<ul style="list-style-type: none"> <li>CSSI proposal</li> <li>Clowder documentation</li> <li>Clowder 1.4 release and PR reviews</li> <li>BD PR reviews</li> <li>Vacation Friday</li> <li>Created Cyprus Clowder instance</li> <li>Created IMLCZO parsers VM</li> </ul>
Marcus Slavenas	<ul style="list-style-type: none"> <li>gltg <ul style="list-style-type: none"> <li>ingesting data</li> </ul> </li> <li>vbd <ul style="list-style-type: none"> <li>parsing mosquito data</li> <li>running model</li> </ul> </li> </ul>	
Maxwell Burnette	<ul style="list-style-type: none"> <li>laser 3d format conversion</li> <li>pipeline health monitoring</li> </ul>	
Michal Ondrejcek	<ul style="list-style-type: none"> <li>MDF <ul style="list-style-type: none"> <li>continue read the docs for the Sphinx</li> <li>NDS Workbench, run Forge examples</li> <li>script cron job discuss with Chad</li> <li>script for FF pipeline</li> </ul> </li> <li>Farmdoc <ul style="list-style-type: none"> <li><input checked="" type="checkbox"/> FD-10 - finish <span>IN PROGRESS</span></li> <li><input checked="" type="checkbox"/> FD-11 - convert arcCoPay <span>IN PROGRESS</span></li> </ul> </li> <li>setup VM for the presentation</li> </ul>	<ul style="list-style-type: none"> <li>MDF <ul style="list-style-type: none"> <li>not done</li> <li>yes</li> <li>e-mail to Chad, discussed, involve ben</li> <li>no big progress</li> </ul> </li> <li>Farmdoc <ul style="list-style-type: none"> <li>done</li> <li>done</li> </ul> </li> <li>e-mail to Amanda about resources, Friday meeting</li> <li>5 min presentation</li> </ul>

Sara Lambert	<ul style="list-style-type: none"> <li>NDS / TERRA <ul style="list-style-type: none"> <li>Keep trying to scale up the TERRA Workbench</li> <li>Maintenance where needed</li> </ul> </li> <li>KnowEnG <ul style="list-style-type: none"> <li>Last week on the project</li> <li>Kubernetes Demo on Wednesday</li> <li>NIH Demo on Friday</li> </ul> </li> <li>Crops in Silico <ul style="list-style-type: none"> <li>Continue collecting feedback and iterating until satisfied</li> </ul> </li> </ul>	<ul style="list-style-type: none"> <li>NDS / TERRA <ul style="list-style-type: none"> <li>Failed attempts trying to scale up the TERRA Workbench</li> <li>Minor Workbench maintenance</li> </ul> </li> <li>KnowEnG <ul style="list-style-type: none"> <li>NIH Demo cancelled due to scheduling conflicts</li> <li>Wrap up final bits of Kubernetes for their stuff</li> </ul> </li> <li>Crops in Silico <ul style="list-style-type: none"> <li>Continue collecting feedback and iterating until satisfied</li> </ul> </li> </ul>
Michelle Pitcel	<ul style="list-style-type: none"> <li>GLTG <ul style="list-style-type: none"> <li><input checked="" type="checkbox"/> <del>GLGVO-426</del> - Groups of Layers - Visually Group Associated Layers <span>DONE</span></li> </ul> </li> <li>IMLCZO <ul style="list-style-type: none"> <li><input checked="" type="checkbox"/> <del>IMLCZO-225</del> - Correct Mahomet Ingestion <span>DONE</span></li> <li>Re-run Parsers for Flux Tower and Allerton non-Decagon</li> </ul> </li> </ul>	<ul style="list-style-type: none"> <li>GLTG <ul style="list-style-type: none"> <li><input checked="" type="checkbox"/> <del>GEOD-4076</del> <span>DONE</span></li> <li><input checked="" type="checkbox"/> <del>GLGVO-422</del> <span>DONE</span></li> <li><input checked="" type="checkbox"/> <del>GLGVO-426</del> <span>DONE</span></li> </ul> </li> <li>IMLCZO <ul style="list-style-type: none"> <li>Re-ran Parsers for Flux Tower and Allerton non-Decagon</li> <li><input checked="" type="checkbox"/> <del>IMLCZO-225</del> <span>DONE</span></li> <li><input checked="" type="checkbox"/> <del>IMLCZO-226</del> <span>DONE</span></li> </ul> </li> </ul>
Omar Elabd		
Pramod Rizal	<ul style="list-style-type: none"> <li>Industry - Pix4D Extractor (W.I.P.)</li> <li>k8s dev setup/test for KnowEnG (W.I.P.)</li> </ul>	<ul style="list-style-type: none"> <li>Industry - Pix4D Extractor (W.I.P.)</li> <li>k8s dev setup/test for KnowEnG (W.I.P.)</li> </ul>
Rob Kooper	<ul style="list-style-type: none"> <li>LSST <ul style="list-style-type: none"> <li>operations requirement gathering</li> <li>parsl</li> </ul> </li> <li>PEcAn <ul style="list-style-type: none"> <li>htcondor meeting</li> <li>create RC VM</li> </ul> </li> <li>Industry <ul style="list-style-type: none"> <li>check on progress of new hardware</li> <li>extractors per space</li> </ul> </li> <li>TERRA <ul style="list-style-type: none"> <li>quarterly report</li> </ul> </li> <li>NCSA/ISDA <ul style="list-style-type: none"> <li>PEARC reviews</li> </ul> </li> </ul>	
Sandeep Puthanveeti I Satheesan	<ul style="list-style-type: none"> <li>BD <ul style="list-style-type: none"> <li>Sprint tasks <ul style="list-style-type: none"> <li>BD-CLI update</li> <li>Code review</li> </ul> </li> </ul> </li> <li>CCROP <ul style="list-style-type: none"> <li>Tool to generate experiment file from template based on user input</li> <li>Other tasks</li> </ul> </li> <li>JUDEL <ul style="list-style-type: none"> <li>Apply Tesseract for OCR and newspaper article segmentation</li> </ul> </li> </ul>	<ul style="list-style-type: none"> <li>BD <ul style="list-style-type: none"> <li>Worked on BD-CLI update</li> </ul> </li> <li>CCROP <ul style="list-style-type: none"> <li>Completed development for tool to generate experiment file from template based on user input</li> </ul> </li> <li>JUDEL <ul style="list-style-type: none"> <li>Working on applying Tesseract for OCR and newspaper article segmentation.</li> </ul> </li> </ul>
Shannon Bradley	<ul style="list-style-type: none"> <li>Even tho I read a lot of email while I was out - I still have email to catch up on</li> <li>Sprint Updates</li> <li>Brown Dog Qtr 1 2018 report - kickoff</li> <li>Get back in a groove 😊 and catch up on everything else</li> <li>new tablet?</li> </ul>	

Yan Zhao		<ul style="list-style-type: none"> <li>• BD <ul style="list-style-type: none"> <li>◦ upgrade to pycldowder2 for meangrey</li> <li>◦ check the silhouette for clusterman – failed to get an answer</li> <li>◦ fix pecan shiny</li> </ul> </li> <li>• GLM <ul style="list-style-type: none"> <li>◦ ingest USGS 04024000 on dev.</li> <li>◦ add helper text for USGS quality code</li> </ul> </li> <li>• CC <ul style="list-style-type: none"> <li>◦ new UI</li> <li>◦ initial implement for user session service.</li> </ul> </li> </ul>
Yong Wook Kim	<ul style="list-style-type: none"> <li>• Finish pycsw extractor by adding geotiff function</li> <li>• Update docker for pyincore/opensees incore jupyterhub docker</li> <li>• Update kubernetes with pyincore jupyterlab</li> <li>• Prepare the COE demo</li> </ul>	<ul style="list-style-type: none"> <li>• Finished pycsw extractor and tested it with docker and clowder</li> <li>• Updated docker for pyincore, opensees, and other jupyterlab extension</li> <li>• Updated kubernetes with new jupyter lab</li> <li>• Tested web mapping on the jupyterlab</li> </ul>