



Manaaki Whenua
Landcare Research

S-MAP Workflows on and off the Cluster

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Pascal Omondiagbe

Presenting an overview of technical work to
support scientists and data managers undertake
repeatable spatial soils research



OutLine

□ Intro

- Robbie and Pascal, Staff Diagram

□ **SMAP quick overview**

- General overview

□ **Digital Soil Mapping**

- Method Overview
- DSM Process
- S-map Ecosystem
- Covariate Process
- DSM modelling Process
- Segmentation Process
- Post processing/Upload

□ **Workflow Development**

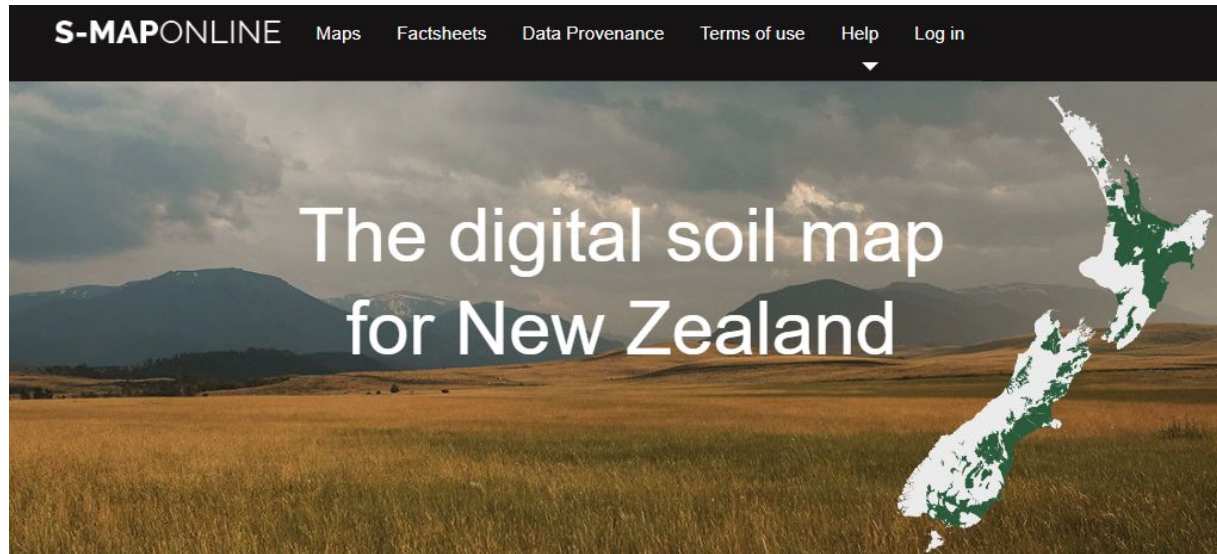
- Requirement
- Retrospective
- Workflow Principle
- Development Process
- DSM Workflow
- Performance/Re-evaluate
- S-Map Workflow Summary

□ **Conclusion**



SMAP

- <https://smap.landcareresearch.co.nz>



EXPLORE MAPS



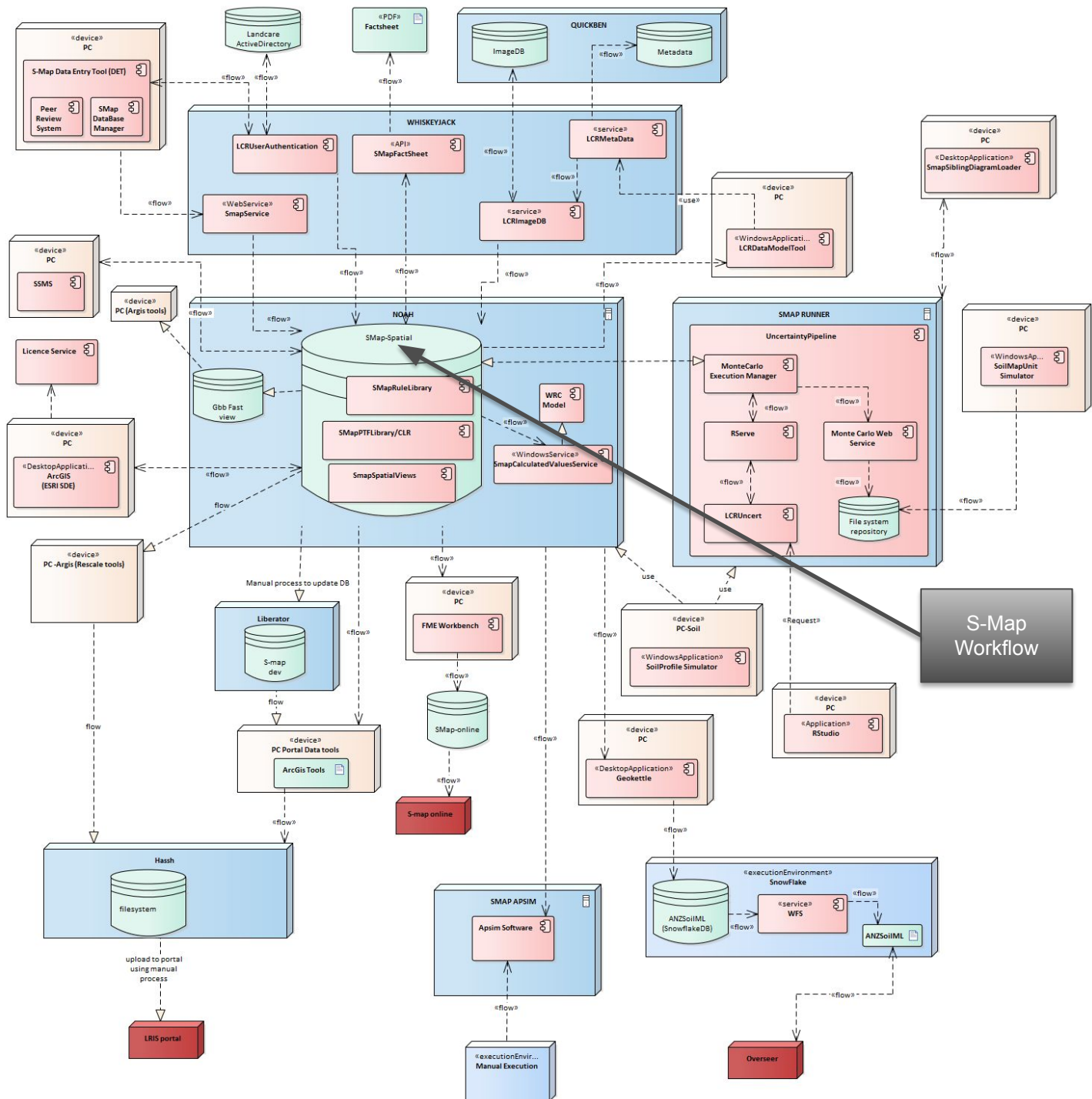
LOCATE ME



BROWSE
FACTSHEETS



FIRST TIME HERE?



Soils – multi scale response to geology, landform, climate and time



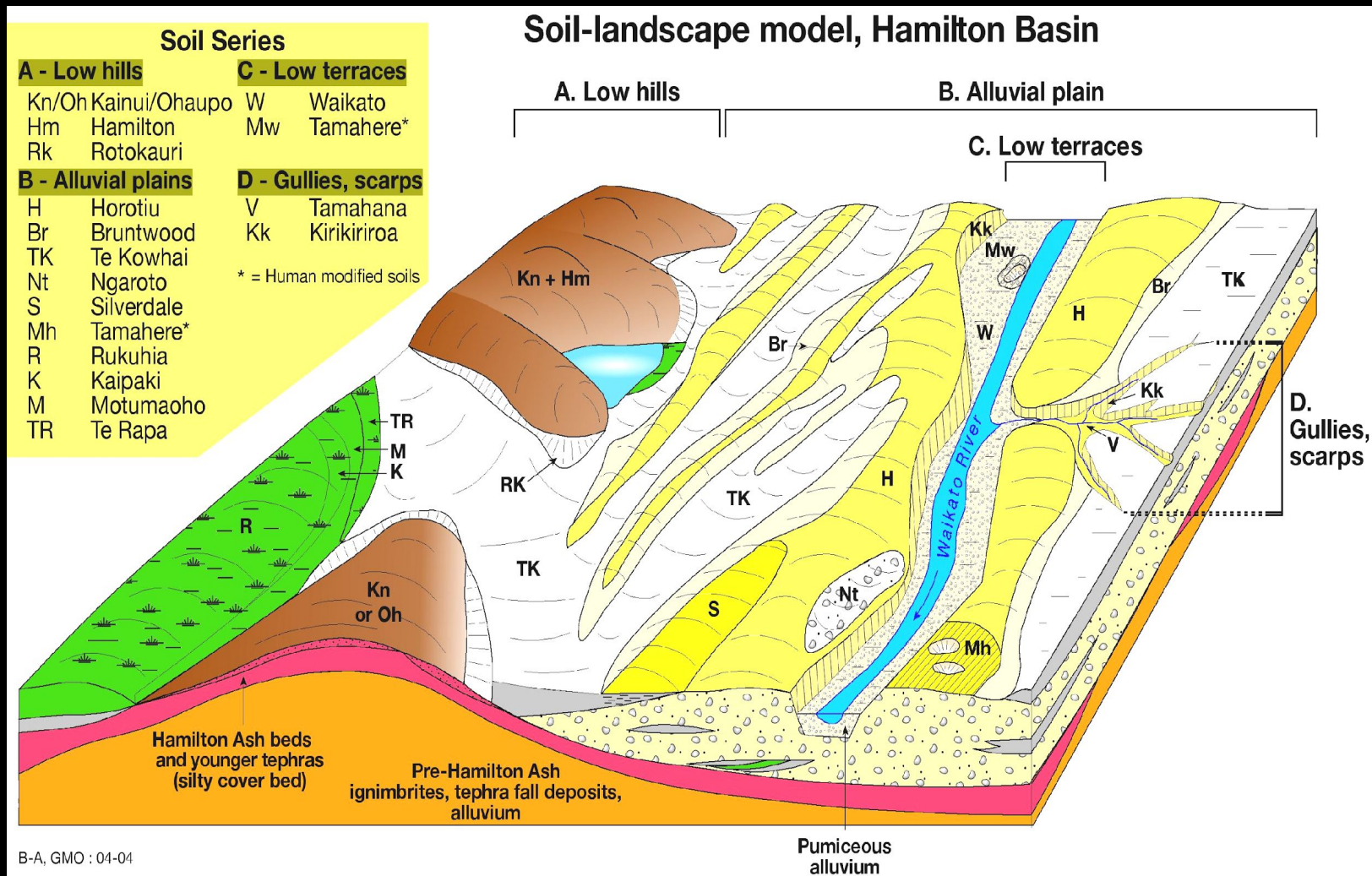
Pedologists







Soil-Landscape Model



B-A, GMO : 04-04



*Allophanic soils from composite
tephras over pumice alluvium*



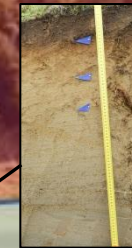
Otor_28 + Kinle_1.1 +
Aroha_3.1



Moes_2.1 + Ngah_9.1

Pumice soils from Taupo Pumice alluvium

Taup_82.1

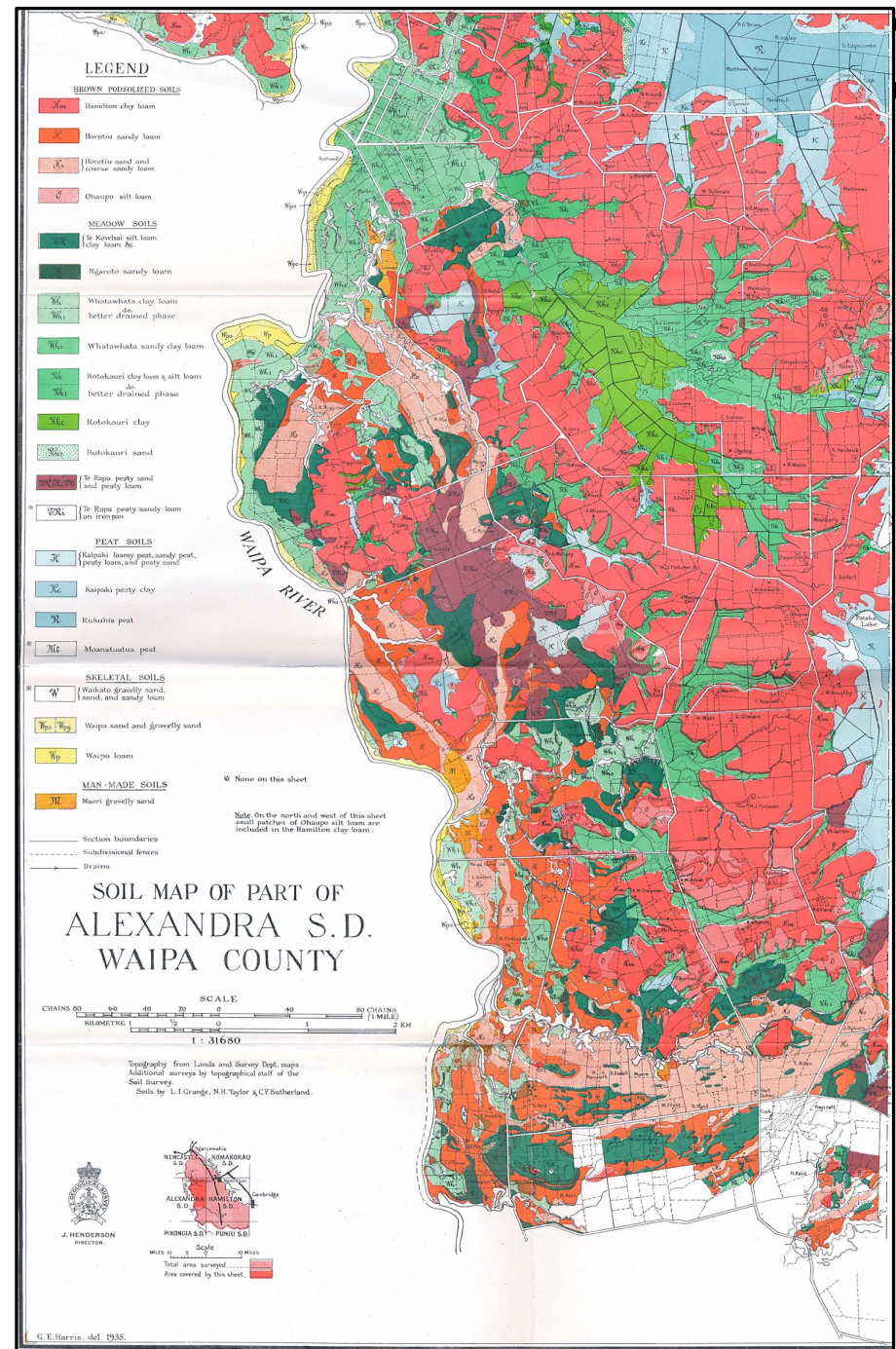


Allophanic soils from post 60ka composite tephras Otor_28.1

[illegible]

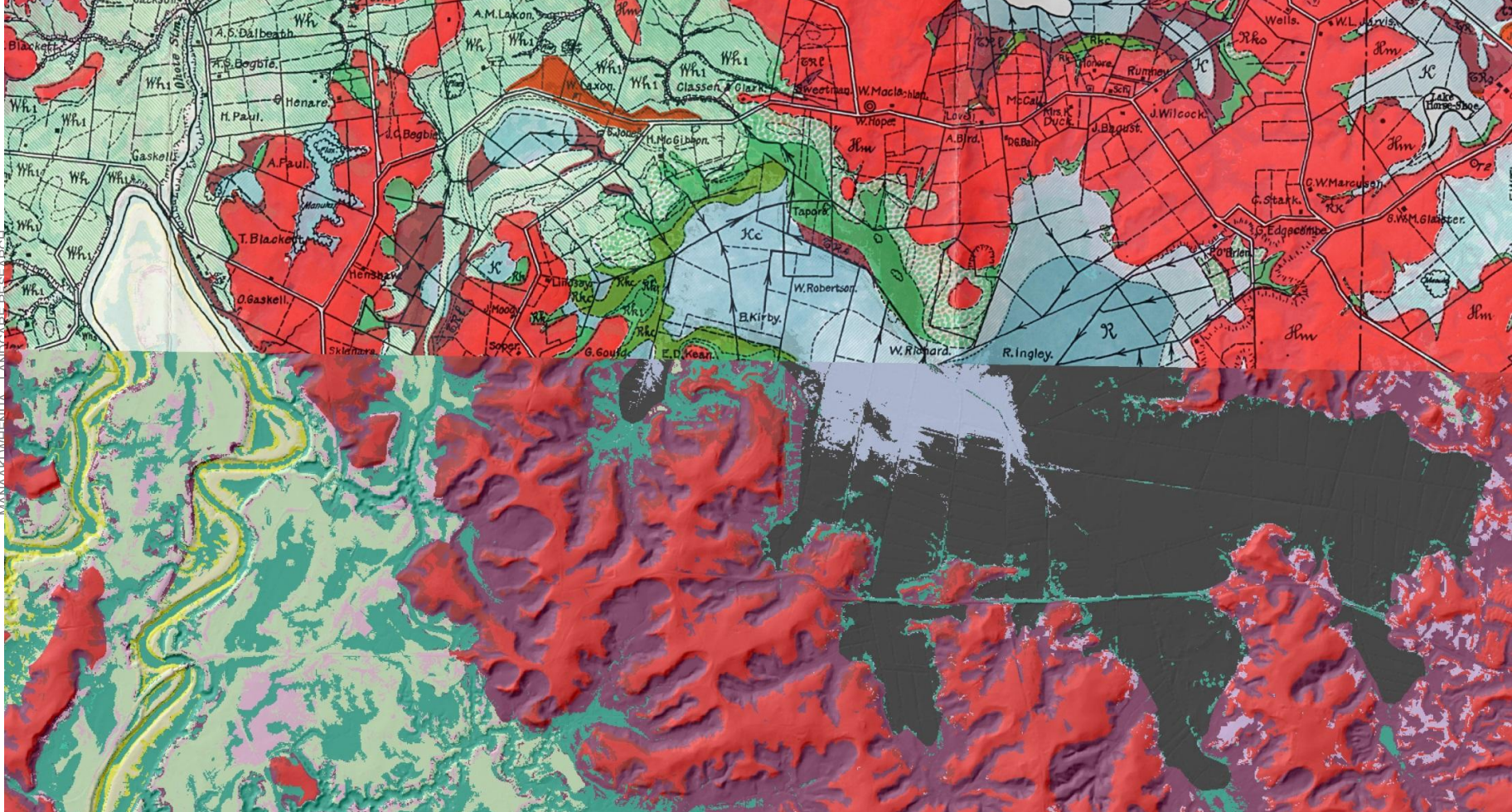
... Soil Map

- Art
- Not data driven
- Not suitable for modern demands
- Regional planning max
- Definitely not farm scale
- S-Map plenty not like



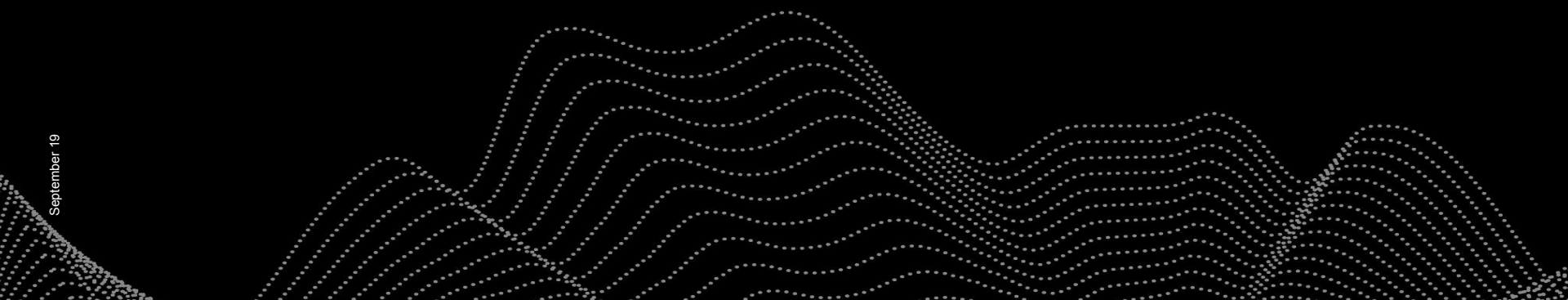


... And so digital soil mapping was born

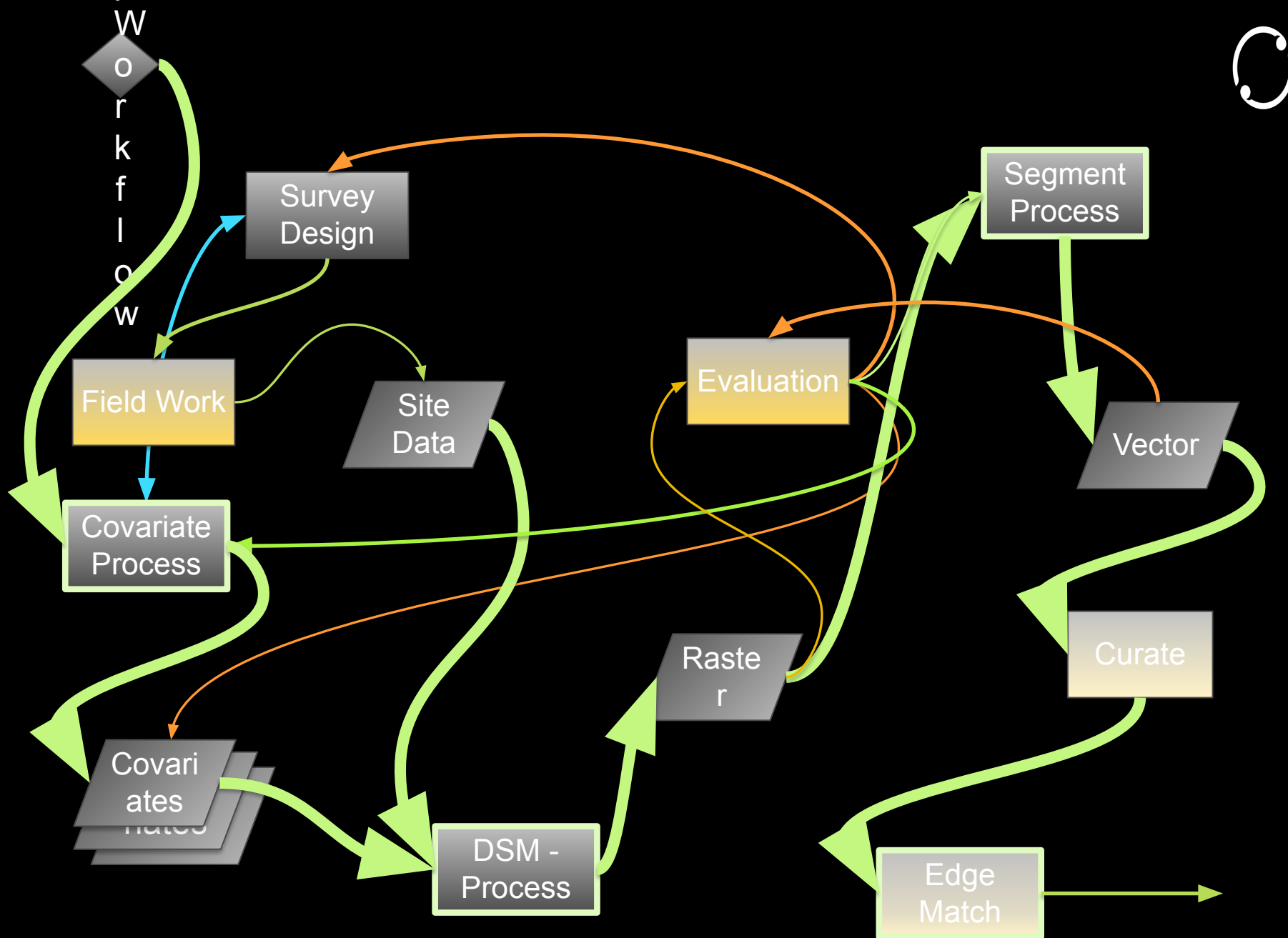




**DSM (Digital Soil Mapping) – developed by
Pedologists and slotted into the normal
manual way of doing things...**









Covariate Process

- LiDAR elevation model, Climate data, Geology layers
- Covariates derived using existing tools
 - ArcGIS
 - GRASS
 - SAGA
 - R
 - C/C++
 - Python
- New covariate algorithms
 - R – e.g Landscape Pattern (neighbourhood elevation range)

Workflow Progress

- Wrap existing non ESRI models in R covariate library
 - Standardised way of writing
 - Definitive Implementations as reusable functions
 - Metadata generation embedded into the workflow
- Assess alternative Implementations of ESRI models



DSM Modelling Process

- Expert feel for Area being mapped (Pedologists)
- Site Data to describe soils and locate geospatially
- Sites + Covariates + Model
 - Currently using an over-fitting model: Random Forest
- Iterative process to develop and test models and covariates
 - Choice of covariates
 - Field Sampling Strategy

Workflow Progress

- Site data “management” in Excel
- R based using Knitr to create evaluation documentation at run time
- Workflow to introduce type and play options
- Scope for parallel processing by tiling assessed but not implemented



Segmentation Process - Raster to Vector

- This area most considered an Art - tells a story
- In - Definitive Raster Soil model/s
- Out – Representative vector (linework) map
- Vector map is an inexact representation correct at the attribute level.
- Removal of small areas (minimum polygon size)
- Smoothing of boundaries

Workflow Progress

- Re-implemented ArcGIS concept in GRASS
- Workflow runs and works
- Segmentation needs complete rewrite for science-art reasons



Post Processing Issues

□ Curation

- Line work manually linked back to soils
- Yet to gain an understanding of the process

□ Edge Matching

- Joining different datasets along boundaries without obvious artefacts
- Traditionally done manually at end of mapping
- Attempting to put at beginning of process

□ Final Curation

- Scripts to prepare data for upload
- Burning in data that didn't model well
- ArcGIS based





Retrospective

□ King said I was daft to build a castle in a swamp...

❖ <https://www.youtube.com/watch?v=aNaXdLWt17A>

□ DSM is Evolving Science

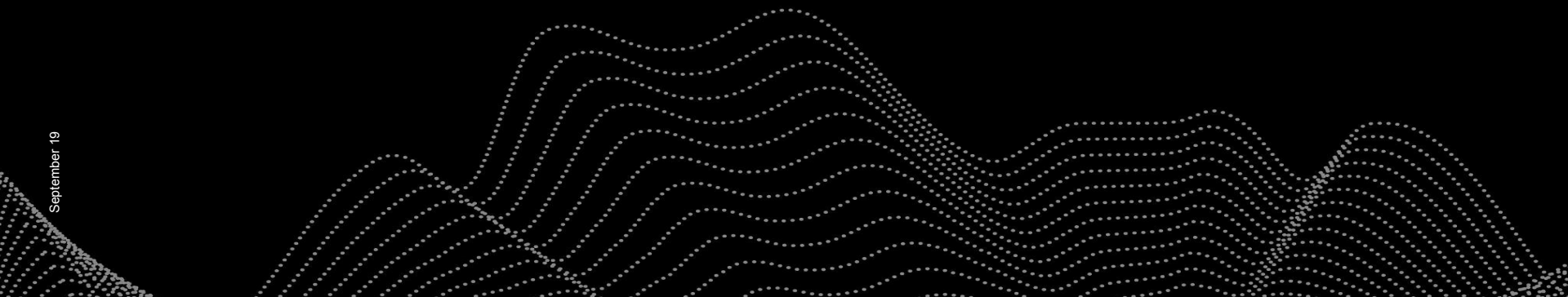
- No previous projects are significantly workflow compatible
- Spatial processing often quite different
- Highly dependent on the ESRI suite for GIS component
- Metadata or documentation for intermediate steps inadequate to replicate work
 - Even where it exists
- Some OS geospatial processing methods not particularly good

□ Pedologists require flexibility into the future

□ First attempt of Segmentation was done using the previous NeSI cluster (New hardware, software, rules... basically new system)



Someone wake Pascal up...





Requirement- The Brief as Given

- Turnkey end-to-end workflow for each DSM project
- Standardised Covariates with national coverage and reproducible

The Brief we wrote ourselves

- Standardise covariate algorithms
- Attempt to document prior DSM work
- Create workflow system for current and future work
- Identify non workflowable aspects of SMAP work
 - Are there other approaches?
- Find holes in the existing methodology
- Identify which steps of the workflow requires HPC
- Code management integrated into the system and documentation



Workflow Principles

- Single language entry-point
 - R – most of our analytical code
- One code base for all platforms
 - Same code to run on local workstation as NeSI
 - Tell it where to run
- R6 Classes for wrapping models
- Generate metadata for each output
 - ISO standard XML
- Home project location for all project data (File System)

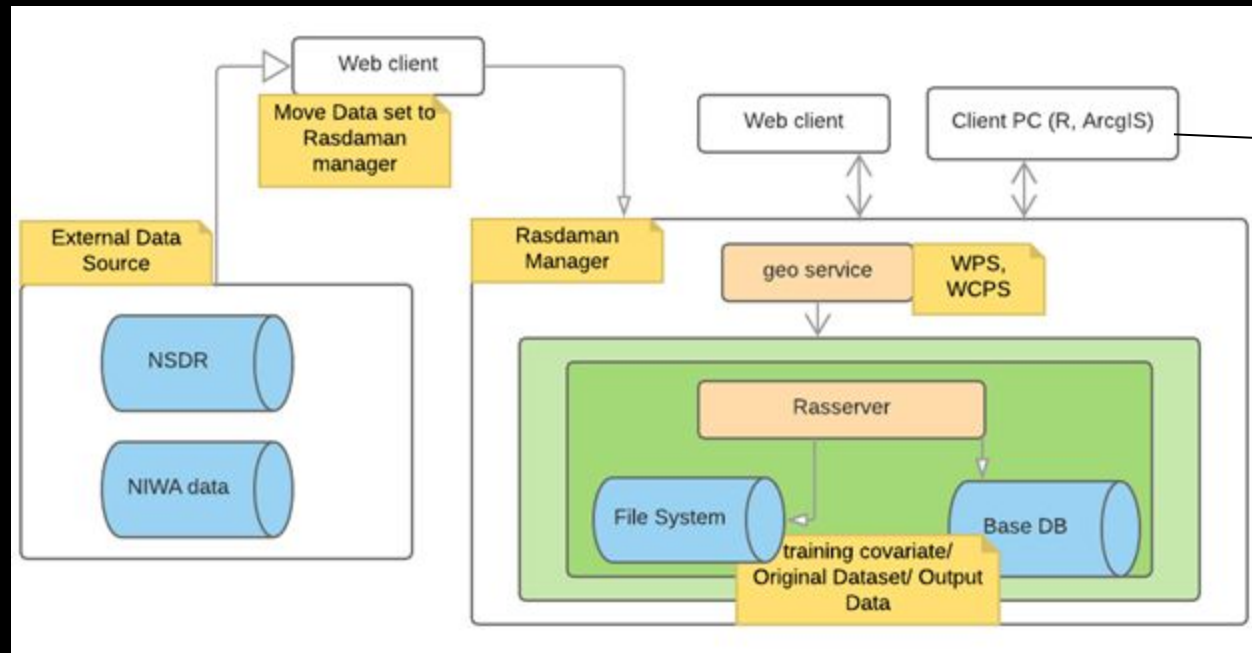


DEVELOPMENT PROCESS

- Coding Standardisation
- Metadata
- R6
- Workflow template

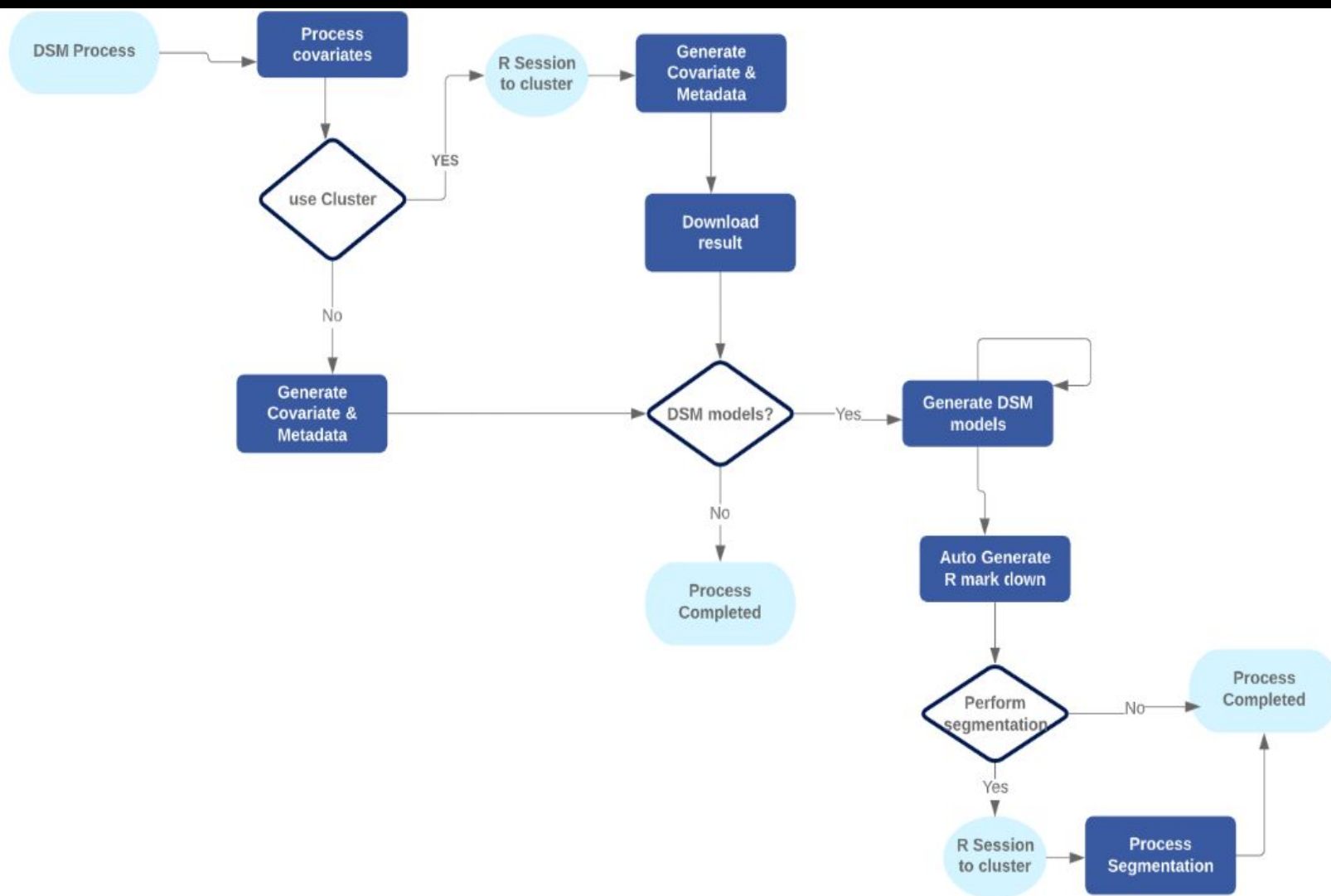


DSM Workflow –Initial plan



HPC

DSM Workflow –Design



- Workflow Via Drake Package





DSM Workflow – Covariate Process

- Wrap existing non ESRI models (e.g. Saga, Grass) in R library
- Previous Grass and Saga R library exists ?
- R6 Classes for wrapping models (easier to use by The DSM guys)
- Process Locally or on the cluster
- Provide easy template to DSM guys

DSM Workflow – Covariate Process (R6 Class)



```
##### STEP 1 Define saga algorithm to execute (e
## Topographic Wetness Index #####
sagaAlgorithm <-function(){
  sm=smapDSM::runSaga()
  sm$path = "C:/Program Files/saga-6.3.0/saga_cmd.exe"
  sm$projectDir = "D:/Projects/smap-dsm/data/"
  sm$rasterFile="test_DEM.tif"
  sm$rasterFolder="D:/Projects/smap-dsm/data/"
  sm$resultPath="result"
  sm$outputVector="final"
  sm$sagaInit()
  sm$metaDataFirstName="pascal"
  sm$metaDataSurName="pascal"
  sm$isCluster=F
  #specify saga algorithm name
  sm$sagaAlgorithm = sm$saga$ta_hydrology$saga_wetness_index
  #get paramters
  sm$sagaParameters()
  sm$parameters$`AREA` (Type: Grid (output):default value:NA)`="area"
  sm$parameters$`SLOPE` (Type: Grid (output):default value:NA)`="slope"
  sm$parameters$`SUCTION` (Type: Floating point:default value:10.000000)`=10.0
  sm$parameters$`AREA_TYPE` (Type: Choice:default value:1)`=1
  sm$parameters$`SLOPE_TYPE` (Type: Choice:default value:1)`=1
  sm$parameters$`SLOPE_OFF` (Type: Floating point:default value:0.100000)`=0.1
  sm$parameters$`SLOPE_WEIGHT` (Type: Floating point:default value:1.000000)`=1.0
  sm$process()
}
```



DSM Workflow – Covariates Process

```
##### workflow Plan when processing a single raster file  
singlePlan <- drake::drake_plan()
```

```
#wetness  
covariate=sagaAlgorithm(),
```

```
#geomorphons  
covariate2=grassAlgorithm()
```

```
)
```

```
clean(destroy = TRUE)|  
make(singlePlan, parallelism = "future", jobs = 2,force = TRUE)  
config<- drake_config(singlePlan,verbose = 3)  
outdated(config)  
vis_drake_graph(config)  
..
```



Up to dat



Importec



Object



Function



DSM Modelling Using Random Forest (Previous Approach)

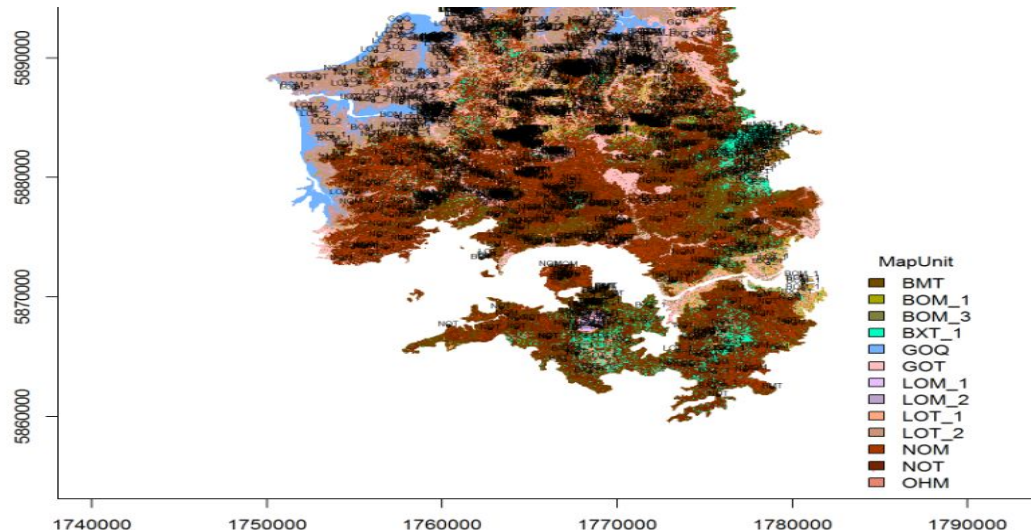


```
#-- -DEM derived
1stGrds[04] <- "P:\\Projects\\SL1705_Franklin_SMap\\Data\\GIS\\Grids\\DJP_backup30_3_2017\\F5ranklin DEM\\TerrainAtts\\mrvbf
5m.img"

1stGrds[05] <- "P:\\Projects\\SL1705_Franklin_SMap\\Data\\GIS\\Grids\\DJP_backup30_3_2017\\F5ranklin DEM\\TerrainAtts\\img
\\eudist.img"
1stGrds[06] <- "P:\\Projects\\SL1705_Franklin_SMap\\Data\\GIS\\Grids\\DJP_backup30_3_2017\\F5ranklin DEM\\TerrainAtts\\img
\\lelementx.img"
1stGrds[07] <- "P:\\Projects\\SL1705_Franklin_SMap\\Data\\GIS\\Grids\\DJP_backup30_3_2017\\F5ranklin DEM\\TerrainAtts\\img
\\lengthx.img"
1stGrds[08] <- "P:\\Projects\\SL1705_Franklin_SMap\\Data\\GIS\\Grids\\DJP_backup30_3_2017\\F5ranklin DEM\\TerrainAtts\\img
\\planc.img"

#--- run the random forest
#rfResult <- randomForest(frmFields, data = tblInput[tblTrainingData,], ntree=500, mtry= 5)
rfResult <- randomForest(frmFields, data = tblTrainingData, ntree=500, mtry= 5)

plot(rfResult)
```





DSM Workflow –Modelling Via SmapDSM Package

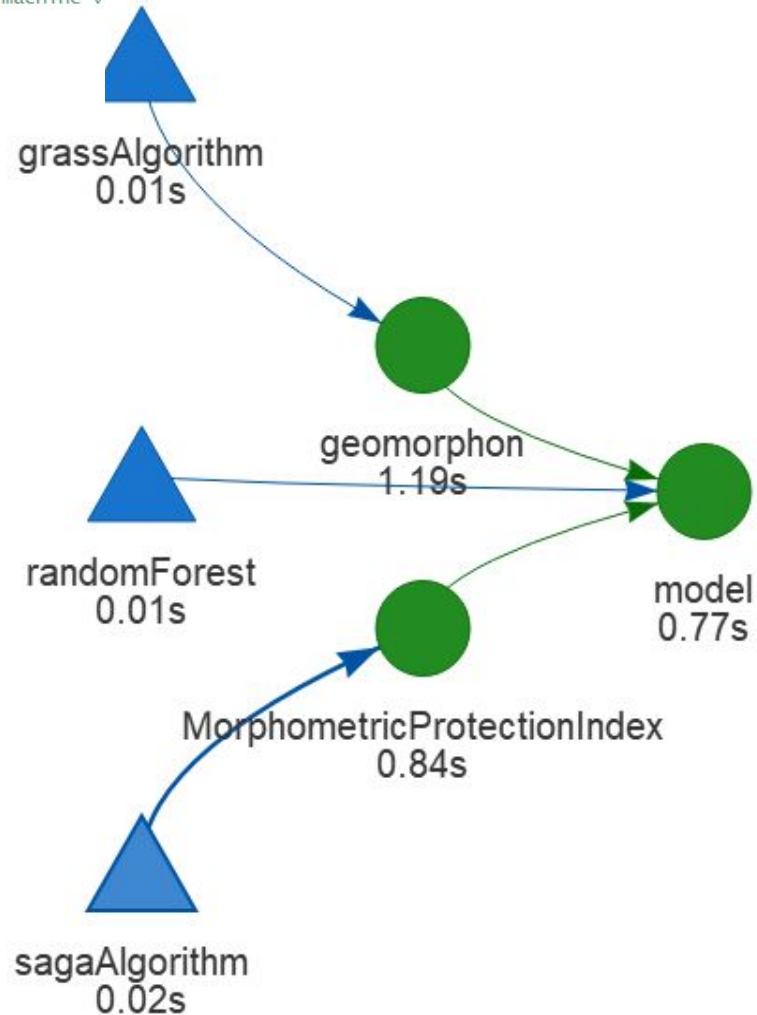
```
##### workflow Plan when processing a single raster file .  
singlePlan <- drake::drake_plan()
```

```
#Morphometric Protection Index  
MorphometricProtectionIndex=sagaAlgorithm(),
```

```
#Calculates geomorphons (terrain forms) and associated geometry using machine v  
geomorphon=grassAlgorithm(),
```

```
model=randomForest(MorphometricProtectionIndex,geomorphon)
```

```
)
```





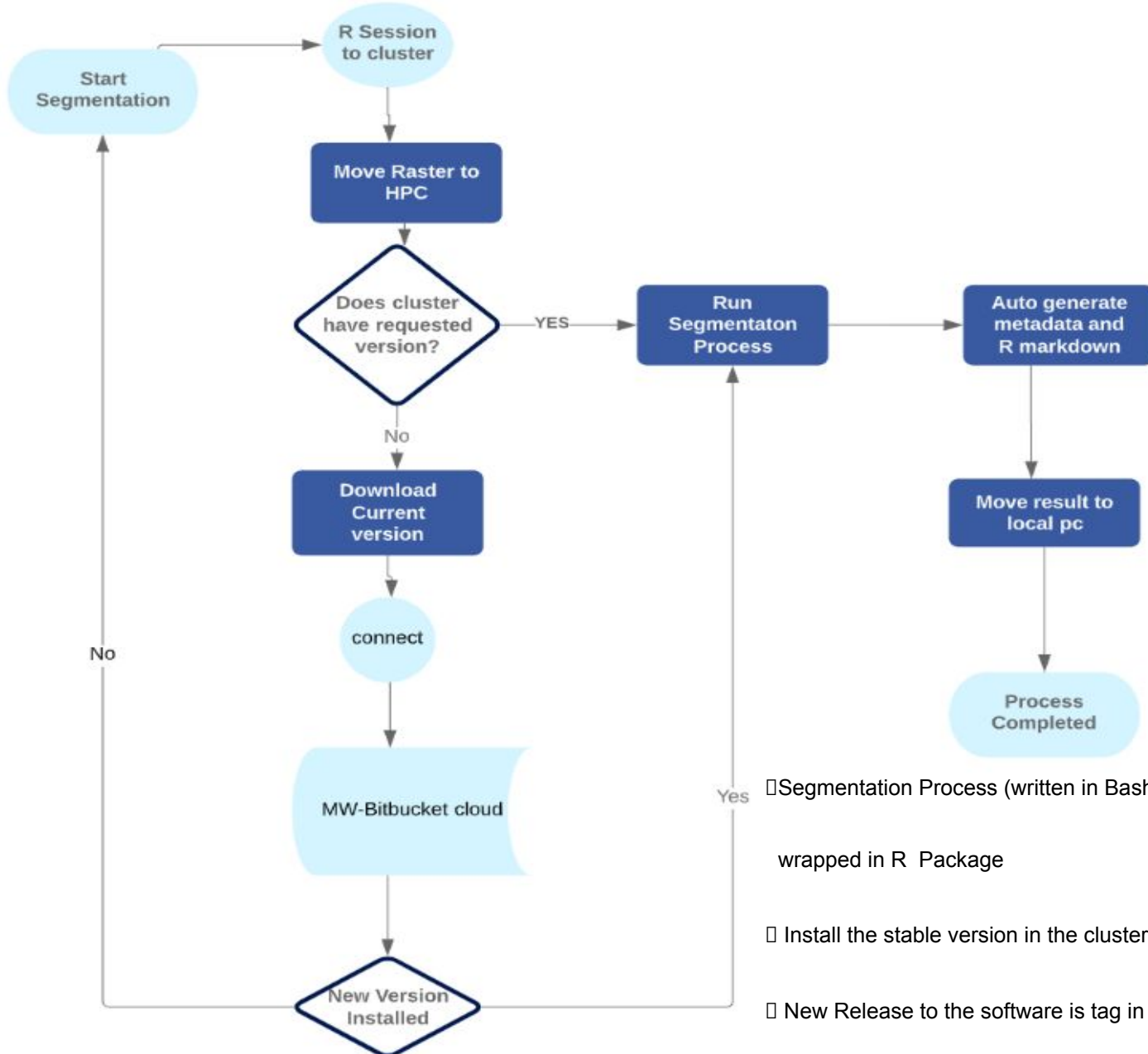
Segmentation Workflow - Initial Design

- Manually Upload segmentation script to cluster
- Manually Run segmentation process
- (you get the picture) Download Result back

□ Not Efficient ?

□ Lots of passwords and keys

Segmentation Workflow Re-design



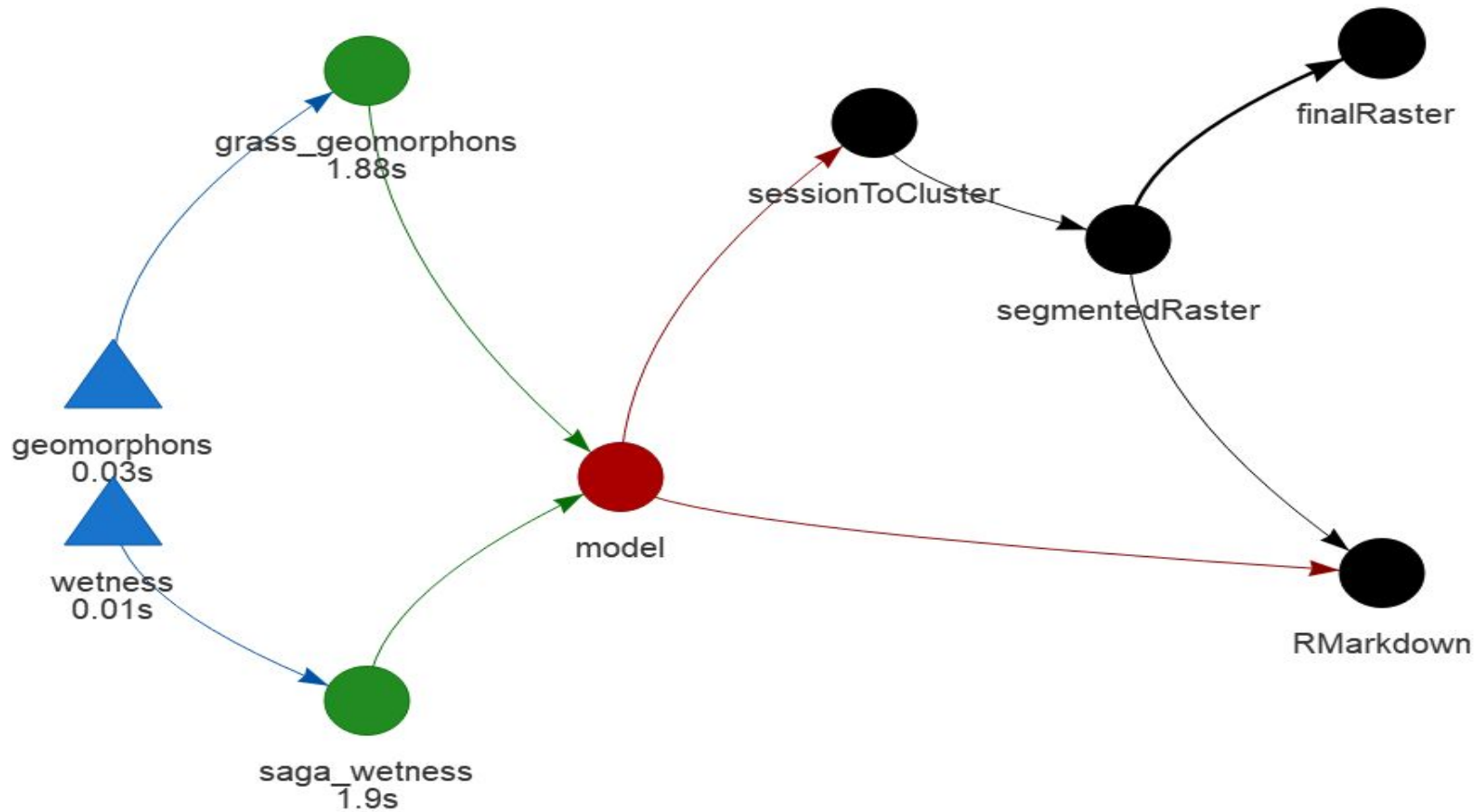
□ Segmentation Process (written in Bash and uses Grass Gis)

wrapped in R Package

□ Install the stable version in the cluster inside the project directory

□ New Release to the software is tag in MWLR Repository

DSM Workflow : End-to-End





Performance/Re-evaluate - Workflow

- Ben Roberts (NeSI) provides report on usage
- Evaluate the performance
- Redesign segmentation script
 - Auto-generate individual tiles
 - Auto-generate SLURM script on the fly
 - Process each tile job on different node
 - R SLURM job to stitch job together



S-Map Workflow - Summary

- Dream of an end-to-end workflow not fully realised (almost there, (not)
- Chose a single language of entry
- An over arching code protocol
- Designing a flexible coding system for evolving science
- Individual components of Workflow being built
- Code management crucial



Acknowledgements

- Linda Lilburne – funding and driving
- James Barringer – preliminary segmentation, feedback
- Scott Fraser, Sharn Hainsworth – DSM work guinea pigs (photos we stole from Scott's presentations)
- Ben Roberts (and other NeSI support)