

Digital Soi Mapping

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Seedling: A User-Friendly, Scalable, and Collaborative Digital Soil Mapping Workflow

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ISRIC – World Soil Information









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- Digital Soil Mapping (DSM) workflows have common features
- Modular structure to ease maintenance (functions and code library)
- Need-based tools and methods

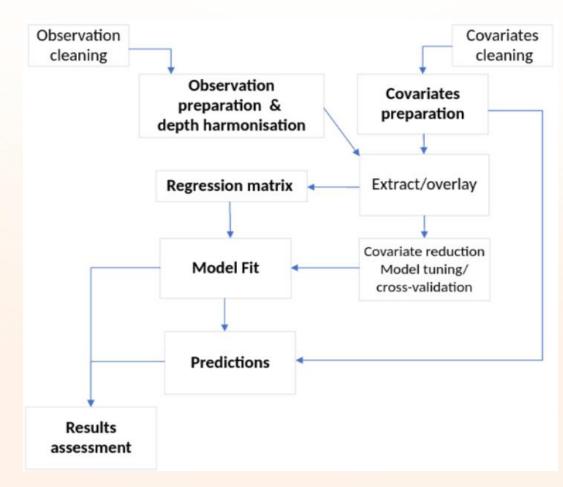
Operational tool that facilitates the process





DSM workflow - Motivation

DSM workflow Common features





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DSM workflow - Seedling

Main characteristics

- Completely R based
- Easy to Use for people with limited DSM expertise
- Multiplatform (Windows, Linux, Mac)
- Running on single laptop/workstation (and on HPC, cloud-computing, etc.)













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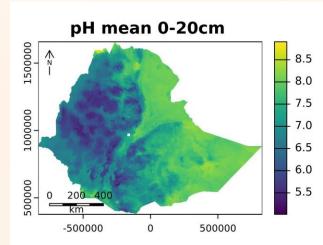
SoilMap

Use Planning

Seedling - What is it

- What is it
 - A modular software stack to perform DSM. R package and configuration files
 - An operational tool
 - Based on SoilGrids workflow (but not limited to)
 - Starts with standardised inputs
 - Produces: Maps, Accuracy stats, Plots, and more
- What is it **not**
 - A data preparation tool
 - A training tool











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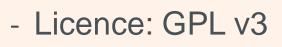
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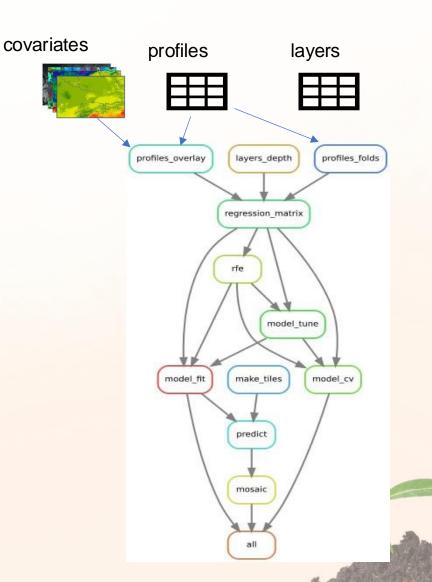
Soil Mapping

Use Planning

Seedling - Main features

- What does it do
 - 2D and 3D modelling
 - Depth standardisation: weighted average of layers (for 2D)
 - Recursive feature elimination
 - Hyperparameter tuning
 - Accuracy (Cross validation) Uncertainty maps
 - Parallel predictions with tiles







Vorking

- R package documentation {isric.dsm.base}
- The online documentation contains:
 - How to install the workflow and its dependencies
 - How to setup and run the workflow
 - Additional information on modelling and design choices
 - The Licence

S:		Description This function generates a regression matrix from the inputs of soil layers, spatial overlay, folds and covariates list file. The function performs left joins on folds, soil layers and spatial overlay to generate a regression matrix. It also		
•		x based on the covariates list file if given.	rerace a regression macrix, it also	
1.16	Usage			
d its	layersDepthProcesse profilesFoldsFile = regressionMatrixFil covariatesListFile profilesStandardFil outputDir = NULL,	<pre>voi, profilesOverlayFile = "{outputDir}/points/profiles_overlay.csv", layersDepthProcessedFile = "{outputDir}/points/layers.depth_processed_{depth_name}.csv", profilesFoldsFile = "{outputDir}/points/forfiles_folds.csv", covariatesListFile = NULL, profilesStandardFile = NULL, outputDir = NULL, outputDir = NULL, ld name = "pid", lyrid name = "tyrid", fold name = "tyrid", easting_name = "x", depth_name = "%", depth_column_name = "depth",</pre>		
flow	lyrid name = "tyrid fold name = "fold", easting name = "x", northing name = "y" depth_name = "0-20" depth_column_name			
elling an	voi min value = 0, voi max value = Inf		_	
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		0 o P	a	
DSM Workflow D	ocumentation	Table of contents		
AUTHOR ISRIC - World Soil Information	PUBLISHED 2023-06-16	Welcome		
Welcome				
This is a Digital Soil Mapping workflow	developed at ISRIC.			
Starting from standardised inputs, it or modelling, performs modelling and pre- output.				
It is a file-based workflow, meaning that both inputs and outputs. Files are name according to the main variables/options	ed and organised	ng		
At this stage this workflow can only har properties	ndle continuous workflo			
Start with Chapter 1 and follow the inst			17 m	

Generate regression matrix

- - C 0 0

R Documentatio

③ R Help

regression_matrix {isric.dsm.base}



1 Getting started 2 Configuration file 3 Workflow script 4 Modeling Choices 5 Design Choices 6 LICENCE References







To use the workflow, you will need two files

- · Configuration file: where all the options are defined
- Workflow script: runs the steps of the workflow

```
# Input/Output files definition
outputDir: "./output/"
profilesStandardFile: "./input_example/points/geul_profiles_4326.csv"
layersStandardFile: "./input_example/points/geul_layers.csv"
covarsDir: "./input_example/covariates"
maskFile: "./input_example/cother/geul_mask.tif"
```

The variable of interest
voi: "pb"

voi_parameters: voi_min_value: 0 voi_max_value: Inf transformation: "notransform"

Dataset variables pid_name: "pid" lyrid_name: "lyrid" easting_name: "x" northing_name: "y" top_layer_name: "top" bottom_layer_name: "bottom" depth_column_name: "depth" fold_name: "fold" stratum_name: "stratum"

```
library(isric.dsm.base)
library(argparser)
config location <- "config.yaml"
p <- arg_parser("Run the entire workflow")</pre>
p <- add_argument(</pre>
     parser = p, arg = "--config_file",
     help = "configuration file location", default = config_location
config <- yaml::read_yaml(parse_args(p)Sconfig_file, eval.expr = TRUE)</pre>
# ## Prepare
profiles_overlay(
     profilesStandardFile = config$profilesStandardFile,
     covarsDir = config$covarsDir,
     profilesOverlayFile = configSprofilesOverlayFile,
     covariatesListFile = configScovariatesListFile,
     outputDir = configSoutputDir,
     easting_name = configSeasting_name,
     northing_name = config$northing_name,
     crs_profiles = configScrs_profiles,
     pid_name = config$pid_name,
     covariate pattern = configScovariate pattern, overwrite = configSoverwrite
layers_depth(
     laversStandardFile = configSlaversStandardFile.
     layersDepthProcessedFile = configSlayersDepthProcessedFile,
```

outputDir = config\$outputDir, pid_name = config\$pid_name,

lyrid_name = config\$lyrid_name





Users and impact

Several users in different countries

- ISRIC's (internal projects/ production pipelines)
- DSM trainings. Rwanda, Kenya, Ethiopia, Nepal
- EJP soil. All national partners (Europe)
- PhD projects
- Other research groups non necessarily involved in projects
- Growing community?



Land Soil Crop

Hubs



Rwanda – map comparison exercise





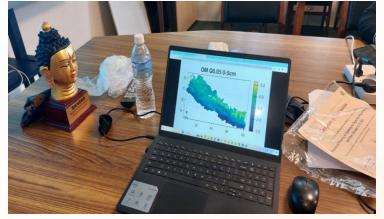




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Conclusions

- Easy to use and scalable tool to lower the bar to enter DSM
- Evolving and improving
- Open source, community involvement?
- Reach out if you want to know more and get involved

https://git.wur.nl/isric/dsm-general/dsm.workflows/seedling











Thank You

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Event Digital Partner





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