



Stimela and CARACal

Towards system agnostic, portable and and configurable data reduction pipelines



Stimela



Radio Astronomy Software*

- 01 | Often written by astronomers, not software developers
- 02 | Difficult to build/install (Time consuming)
- 03 | Difficult to install multiple packages on the same system
- 04 | CASA can do most things. But unnecessarily large, complex and rigid. Does not play well with others.





What's required of a *good* pipeline

1

Version control + issue tracking

github gitlab sourceforge svn

2

Documentation

readthedocs github gitlab

3

Packaging

pypi kernsuite well managed ppa

4

Continuous integration

travis jenkins integrity gitlab github

5

Isolation + portability

docker podman/buildah singularity udocker



Pipeline/Recipe

A set of tasks connected in a suitable way so as to produce a known result given a set of known input data and parameters. Furthermore, a pipeline must:

1. Be versioned
2. Exit with a non-zero exit status if it fails





Cab (pipeline task)

An application that processes or visualizes data.

1. A Cab takes inputs in a standardized form
2. Applications can be wrapped to conform to a Cab standard
3. Cabs are executed using container technology
 - a. Stimela supports **docker, singularity, and podman**





WSClean Cab

```
pipeline.add("cab/wsclean",
  "imager_example_robust_{:d}".format(i),
  {
    "msname": MS,
    "weight": "briggs {:d}".format(robust),
    "prefix": "{:s}_robust-{:d}".format(PREFIX, robust),
    "npix": 2048, # Image size in pixels
    "scale": 2, # Size of each square pixel
    # Perform 1000 iterations of clean (Deconvolution)
    "niter": 1000,
    "pol": "I",
    "multiscale": True,
    "multiscale-scales" : [0,2],
  },
  label="Imaging MS, robust={:d}".format(robust),
  cpus=2,
  memory_limit="2gb")
```

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Raw

```
1 {
2   "task": "wsclean",
3   "base": "stimela/wsclean",
4   "tag": ["1.6.0-1", "1.2.3", "1.6.3"],
5   "description": "WSClean imaging software",
6   "version": ["2.6", "2.8", "2.9"],
7   "prefix": "-",
8   "binary": "wsclean",
9   "junk": [],
10  "msdir": true,
11  "parameters": [
12    {
13      "info": "MS(s) to be imaged. If multiple mses are specified, they need to be phase-rotated to the same point on the sky",
14      "name": "msname",
15      "id": "msfile",
16      "default": null,
17      "dtype": "list:file",
18      "required": true
19    },
20    {
21      "info": "Prefix for output products. Default is prefix of MS",
22      "name": "prefix",
23      "id": "output",
24      "default": null,
25      "dtype": "file",
26      "mapping": "name"
27    },
28    {
29      "info": "Specify number of computing threads to use, i.e., number of cpu cores that will be used. None means use all cores",
30      "dtype": "int",
31      "default": null,
32      "name": "threads",
33      "mapping": "j"
34    },
35    {
36      "info": "Limit memory usage to the given fraction of the total system memory. This is an approximate value.",
37      "dtype": "float",
38      "default": 100,
39      "name": "mem"
40    }
41  ],
42 }
```

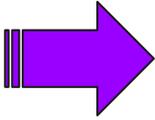


Python UI

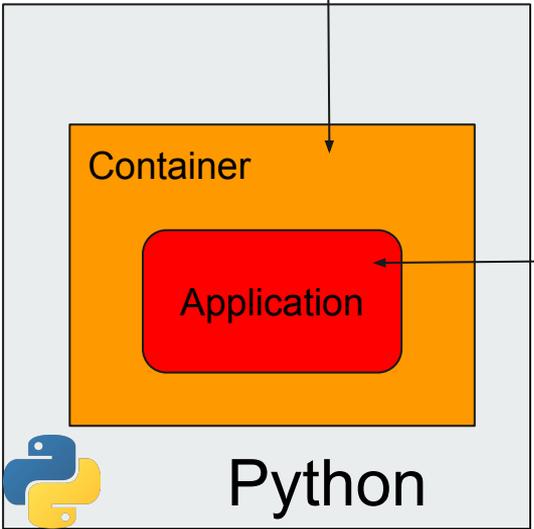
User has access a diverse set of applications
via a simple python interface.

```
1 import stimela
2
3 MS = "sim-meerkat.ms"
4
5 pipeline = stimela.Recipe("Simulate point sky", msdir="msdir",
6                           indir="inputs", outdir="outputs")
7
8 pipeline.add("cab/simms", "makems", {
9     "msname": MS,
10    "tel": "meerkat",
11    "direction": "J2000,0deg,-30deg",
12    "freq0": "1.326GHz",
13    "dfreq": "1MHz",
14    "nchan": 32,
15    "dtime": 1, # in seconds
16    "synthesis": 1, # in hours
17 }, label="Create an empty meerkat MS")
18
19 # the simulator cab is wrapper of the MeqTrees turbo-sim.py script
20 pipeline.add("cab/simulator", "simsky", {
21     "msname": MS,
22     "skymodel": "points_sky.lsm.html",
23     "addnoise": True,
24     "sefd": 541,
25     "column": "DATA",
26 }, label="Simulate point sources into MS")
27
28 for i, robust in enumerate([-1, 0, 1]):
29     pipeline.add("cab/wsclean", f"image_{i}", {
30         "msname": MS,
31         "column": "DATA",
32         "scale": "lsec",
33         "size": 1024,
34         "niter": 5000,
35         "name": MS[:-3],
36         "weight": f"briggs {robust:.2f}",
37     }, label=f"Imaging simulation with Briggs robust={robust}")
38
39 pipeline.run()
```

stimela



Docker *singularity* podman/buildah *udocker*



CASA	Kernsuite/ pypi
WSClean	
ragavi	
Cubical	
simms	
MeqTrees	
DDFacet	
SoFiA	
+++	



Key Takeaways

1

Task definition is standardized to allow a simple and unified UI. Radio interferometry tools essentially become Python functions (even CASA!)

2

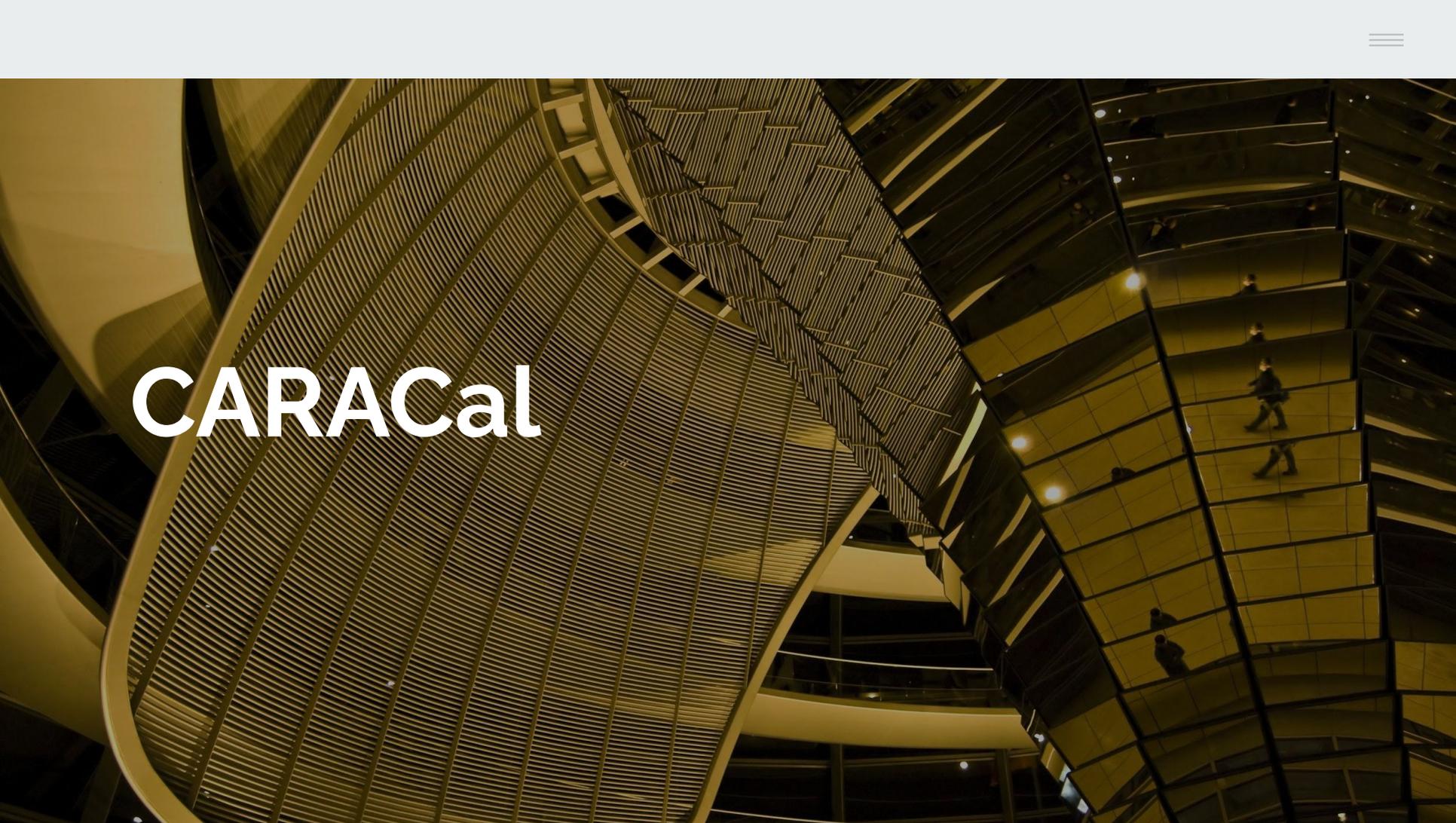
User does not need to install any of the applications supported by stimela. The applications are containerized and can be downloaded from an online registry (hub.docker.com).

3

Stimela is not a pipeline, but a framework that allows the creation of pipelines.

4

Container technology guarantees reproducibility and stability, while the simple UI facilitates transparency and debugging.



CARACal

Graduate students

Feb 2019
Busy week

Hogsback,
Eastern Cape,
South Africa



Rhodes University | SARAO | INAF | University of Pretoria | Wits University



What is it?

01



Stimela-based end-to-end data reduction pipeline for RI data. Not limited to MeerKAT data

Configurable, portable and system agnostic. Easy to install

03



Can do both continuum and spectral line reductions. Includes full direction-dependent treatment and mosaicing.

02



How does it work



```
3 general:
4   prefix: mypipelineun
5
6 getdata:
7   dataid: ['']
8   extension: ms
9
10 transform:
11   enable: true
12   field: calibrators
13
14 flag:
15   enable: true
16   field: calibrators
17   label_in: cal
18   flag_autocorr:
19     enable: true
20   flag_spw:
21     enable: true
22     chans: '*:856~880MHz , *:1658~1800MHz, *:1419.8~1421.3MHz'
23     ensure_valid: false
24   flag_mask:
25     enable: true
26     mask: meerkat.rfimask.npy
27     uvrange: '0~1000'
28   flag_rfi:
29     enable: true
30     flagger: aoflagger
31     aoflagger:
32       strategy: firstpass_QUV.rfis
33
34 crosscal:
35   enable: true
36   label_in: 'cal'
37   set_model:
38     enable: true
```

User Interface (YAML)

Backend

- crosscal_worker.py
- ddcal_worker.py
- flag_worker.py
- getdata_worker.py
- inspect_worker.py
- line_worker.py
- mask_worker.py
- mosaic_worker.py
- obsconf_worker.py
- prep_worker.py
- selfcal_worker.py
- transform_worker.py

workers = stimela recipes



Data Products

Gain tables and plots

RAW visibility plots

Continuum images

Spectral-line cubes

Detailed and summary log files

Jupyter notebook (or HTML) reports

Calibrated visibilities and plots

Source catalogs

A horizontal bar with a teal segment on the left and an orange segment on the right, positioned above the main heading.

Over to you, Josh.

<https://caracal.readthedocs.io> <https://github.com/caracal-pipeline/caracal>

<https://github.com/ratt-ru/Stimela>

<https://github.com/ratt-ru/scabha>

