

Introduction to PCR-GLOBWB code



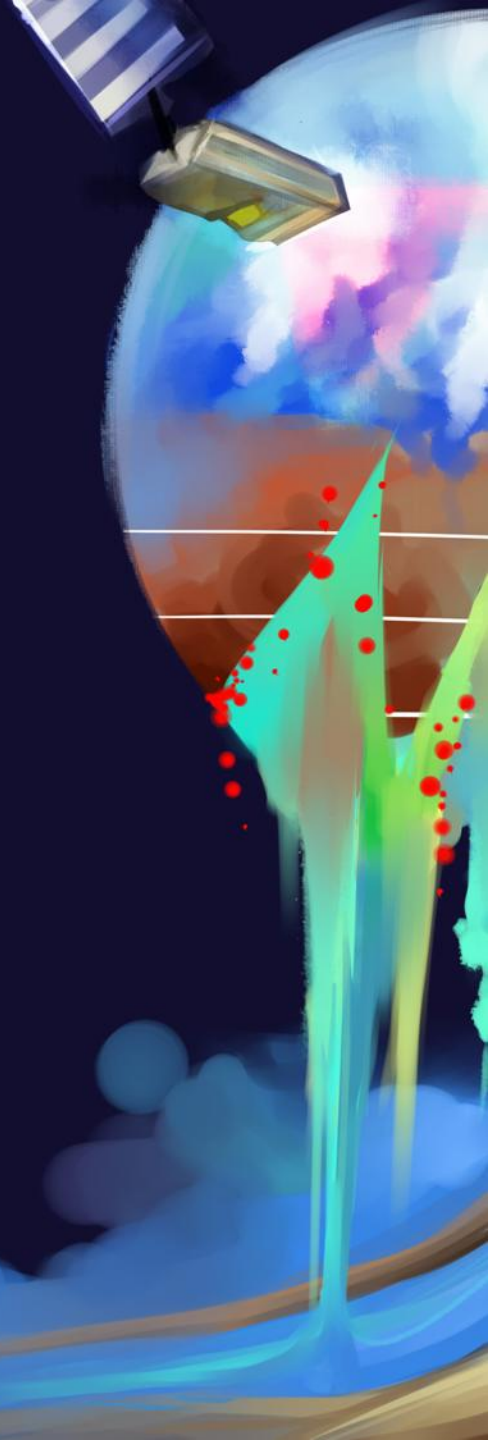
E.H.Sutanudjaja@uu.nl ; +31 30 253 2988 ; Room: Zon104



Universiteit Utrecht

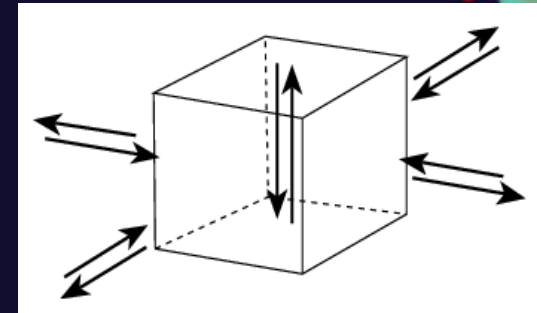
Outline

- PCRaster (some basic knowledge).
- PCR-GLOBWB
 - PCR-GLOBWB code/scripts
 - PCR-GLOBWB ini file (configuration/setting)



PCRaster

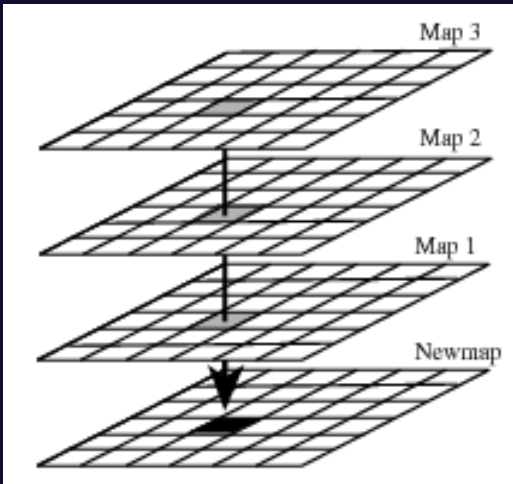
- Programming language used in PCR-GLOBWB.
 - PCR-GLOBWB also uses its Python framework.
- PCRaster:
 - a set of computer tools for storing, manipulating, analyzing and retrieving geographic information
 - RASTER based
 - POINT and NEIGHBOURHOOD operators



- PCRaster documentation:
<http://pcraster.geo.uu.nl/pcraster/4.1.0/doc/manual/index.html>

PCRaster: Local/point operation

- Point operation: A new map is generated on a cell-by-cell basis. No lateral/neighborhood relations between cells are included.
- Example: plus (+) operations



```
Expr1 = readmap("Expr1.map")  
Expr2 = readmap("Expr2.map")  
Result = Expr1 + Expr2
```

Result.map

MV	8	2
2	MV	-6
100	-7	16

Expr1.map

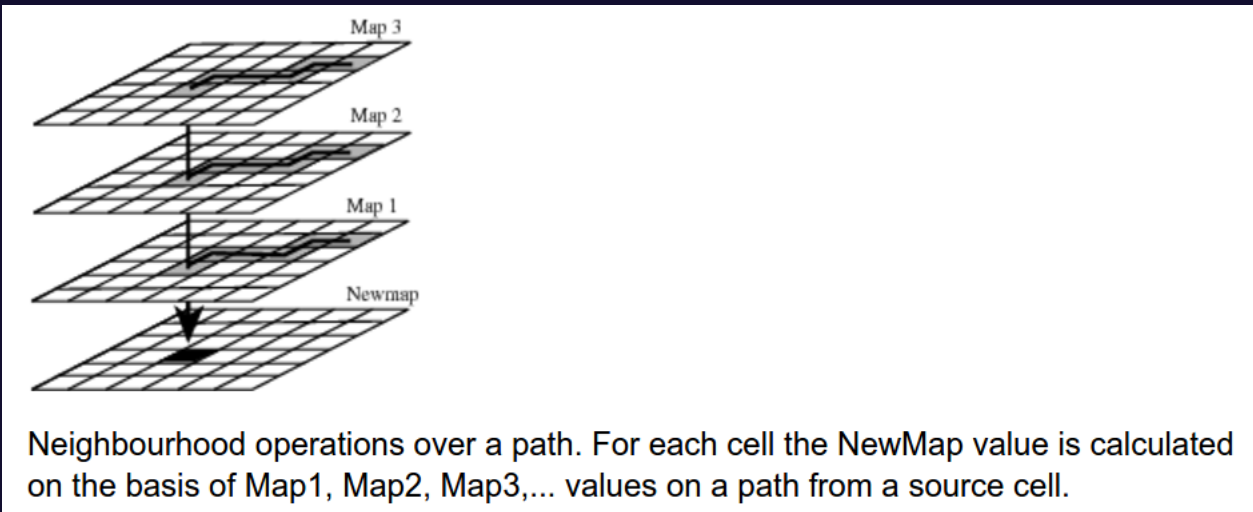
2	6.2	-3
1	MV	7
86	-1	12

Expr2.map

MV	1.8	5
1	3	-13
14	-6	4

PCRaster: Neighbourhood operations

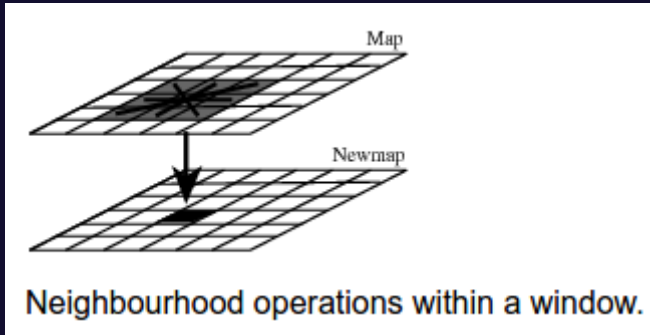
- Neighbourhood operations relate the cell to its neighbours.
- Example:



- Several categories:
 - Window operations (e.g. 3x3 averaging)
 - LDD operations (following a drainage network, upstream-downstream connection)
 - Area (zonal) operations (e.g. to calculate country statistics).
 - Map operations (e.g. statistics over an entire map).

PCRaster: Window operations

- Example: windowaverage



- Other functions:
windowtotal,
windowmaximum,
windowminimum, etc.

```
• python  
Expr = readmap("Expr.map")  
Result1 = windowaverage( Expr, 6)
```

Result1.map

0.333	0.6	-5.4	-4.83	-7.75
1.2	1.38	-2.25	-2.22	-4.17
1.4	1.62	6.75	5.89	7.83
0.667	1.44	6.67	7.22	9.33
-0.25	0.833	8.5	9.33	12.5

Expr.map

0	-1	1	-30	0
2	mv	1	2	-3
3	2	3	4	2
0	0	2	40	2
1	-2	4	7	1

PCRaster: LDD operations

- Example: upstream: sum of the cell values of its first upstream cells(s)

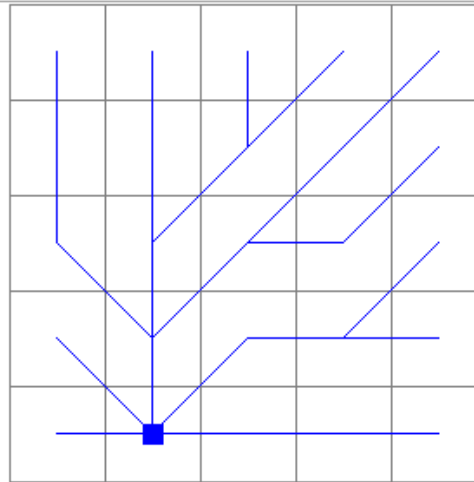
• python

```
Ldd = readmap("Ldd.map")  
Expr = readmap("Expr.map")  
Result = upstream(Ldd, Expr)
```

Result.map

0	0	0	0	0
1	1	4	4	0
1	3	MV	4	0
0	MV	4	8	0
0	14	5	5	0

Ldd.map



Expr.map

1	1	2	2	4
1	1	2	2	4
2	2	MV	4	4
2	2	2	4	4
3	7	5	5	5

- Some other functions:
 - accuflux: Accumulated material flowing into downstream cell
 - accutravelttime: Transport material downstream on a given velocity
 - kinematic, etc.

PCRaster: Area operation

- Example: areamaximum: maximum cell value within a class

- python

```
Class = readmap("Class.map")  
Expr = readmap("Expr.map")  
Result = areamaximum( Expr, Class)
```

Result.map

-6	8	-6	-6	MV
8	8	-6	-6	-6
8	8	8	8	8
8	8	8	8	8
8	MV	MV	2.5	2.5

Class.map

2	6	2	2	MV
6	6	2	2	2
6	6	0	0	0
6	6	0	0	0
6	3	3	4	4

Expr.map

-9	0	-6	-6	-6
1	1	-6	-6	MV
1	1	-1	7	2
1	1	3	5	8
8	MV	MV	2.5	1.4

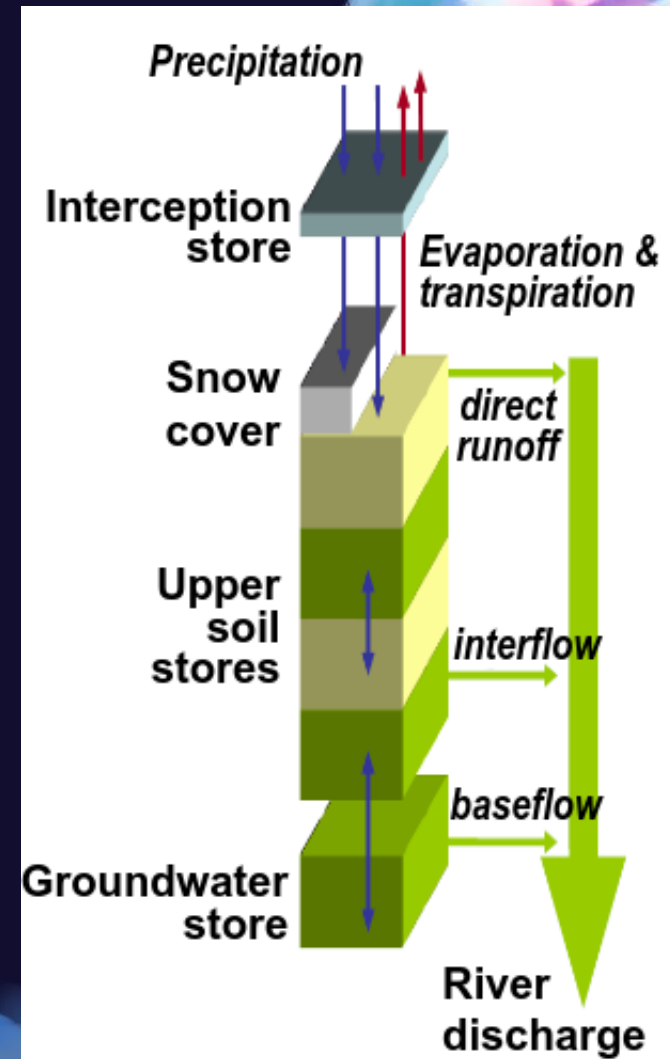
- Some other functions: areaaverage, areaminimum, etc.

PCR-GLOBWB code (a cooking book about how to run the model)



PCR-GLOBWB 1.0 (structure)

- Original PCR-GLOBWB 1.0 (Van Beek et al, 2011)
- Resolution: 30 arc min (~50 km)
- Daily time step
- **Vertical** flow through each cell (**POINT operations**).
 - Sub-grid variabilities within cells are considered: topography, soil and land cover variations.
- Interaction between cells:
 - Channel routing along drainage network (**LDD**).
 - Surface water bodies represent either streams or lakes/reservoirs that buffer stream-flow.
 - 600 largest reservoirs.
 - Lakes and wetlands (including evaporation).

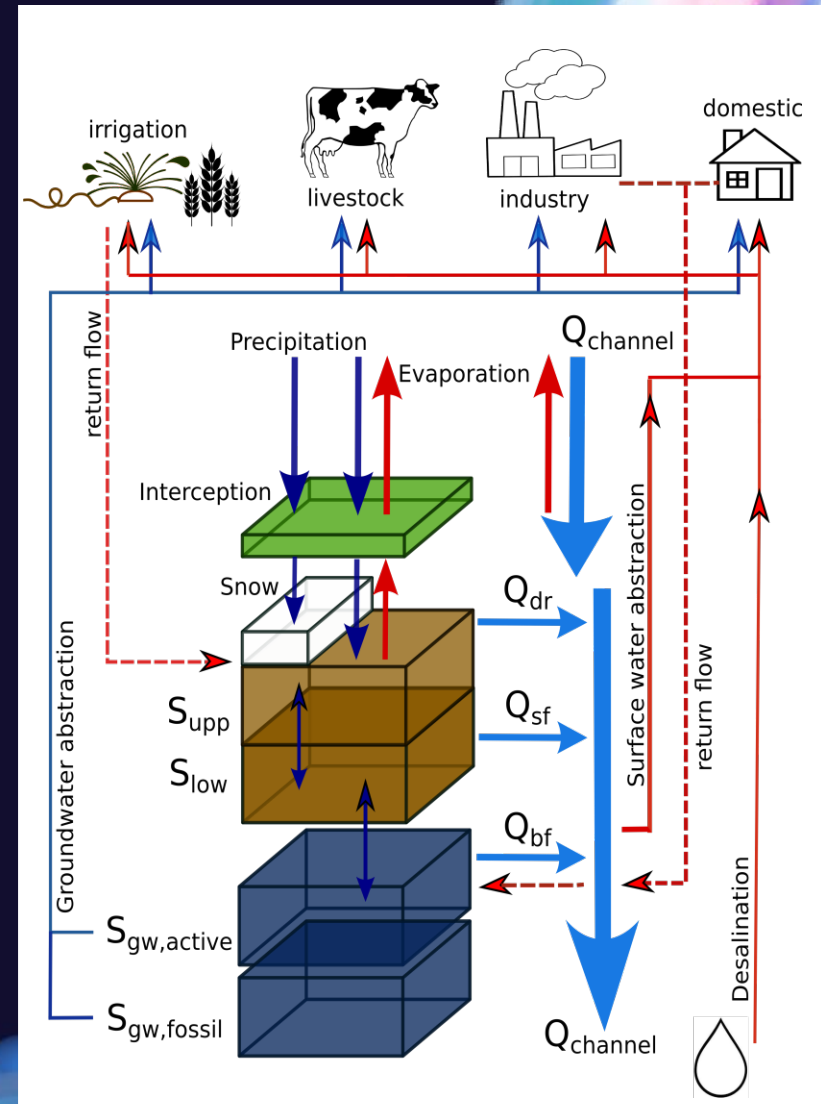
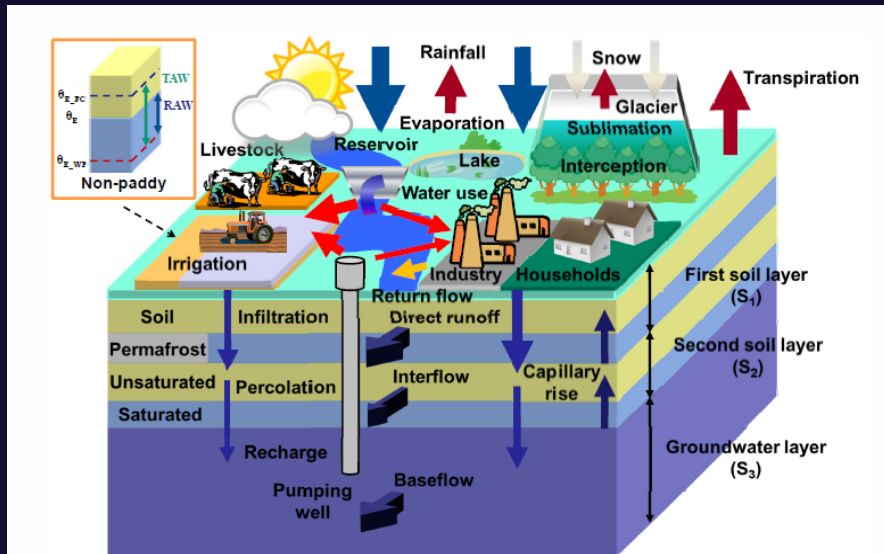


PCR-GLOBWB 2.0

- **Integrated hydrology & water resources model**, i.e. including online/interactive simulation of water demand and abstraction. Features added (examples):

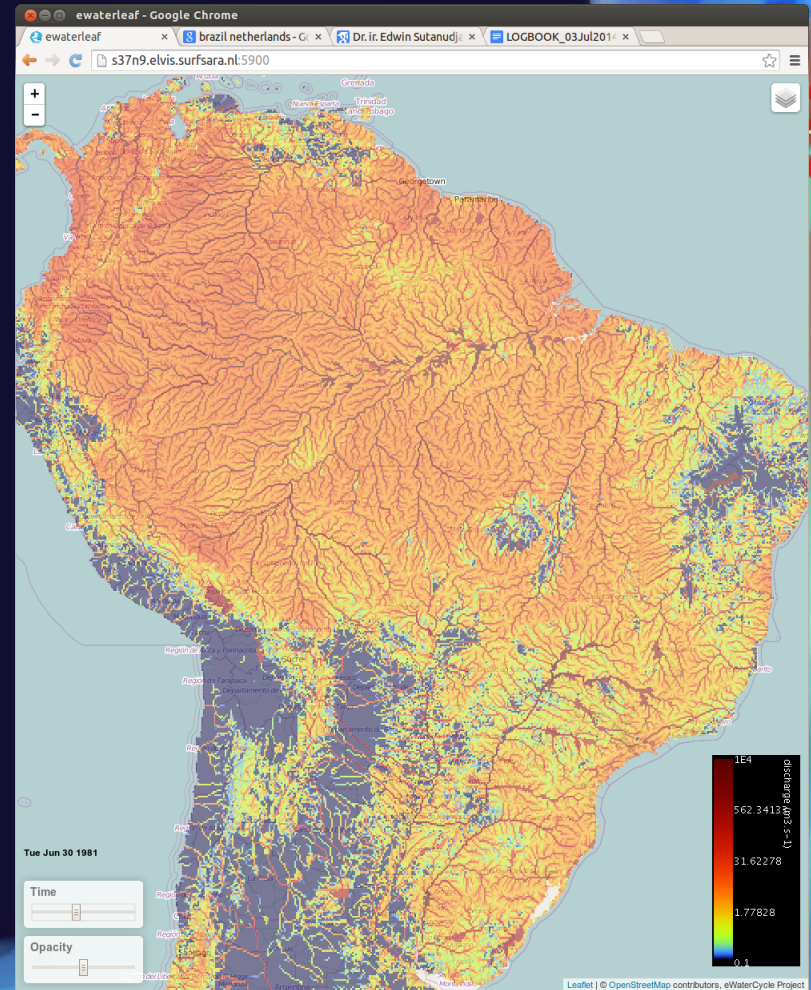
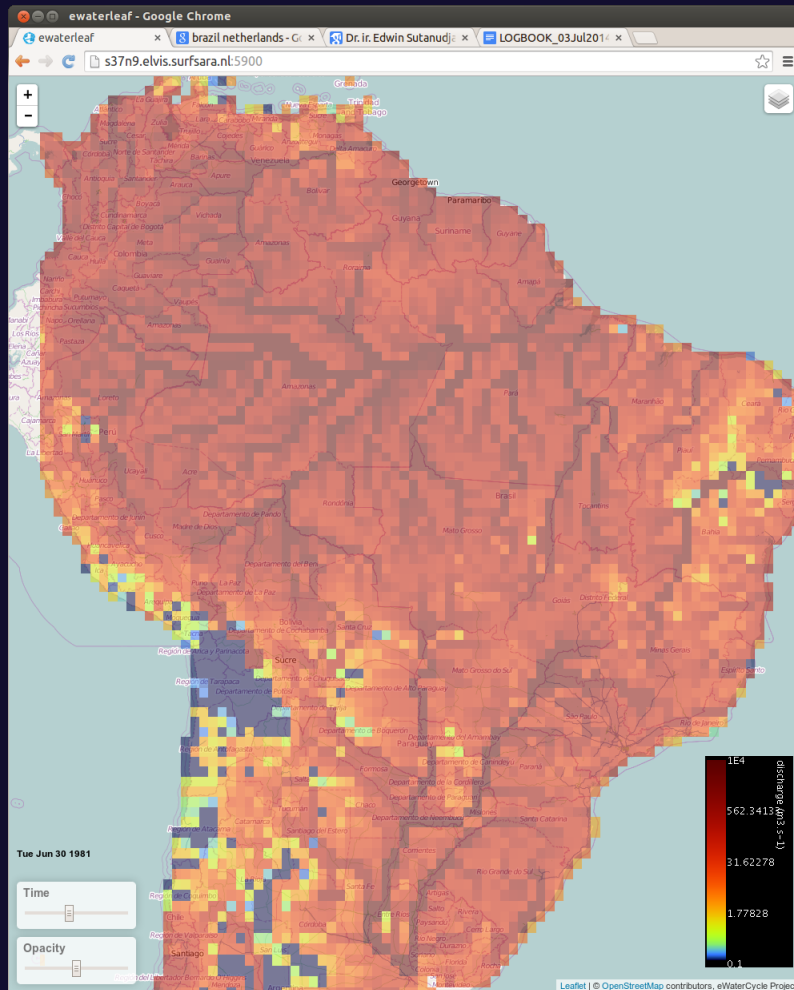
- Water demand & irrigation.
- > 6000 reservoirs (Grand).
- Integrated flood inundation module.

The model code is rewritten in Python and open source. Also, many refactoring actions have been done to speed up the computation (particularly for 5 arcmin ~10 km resolution)



PCR-GLOBWB 2.0

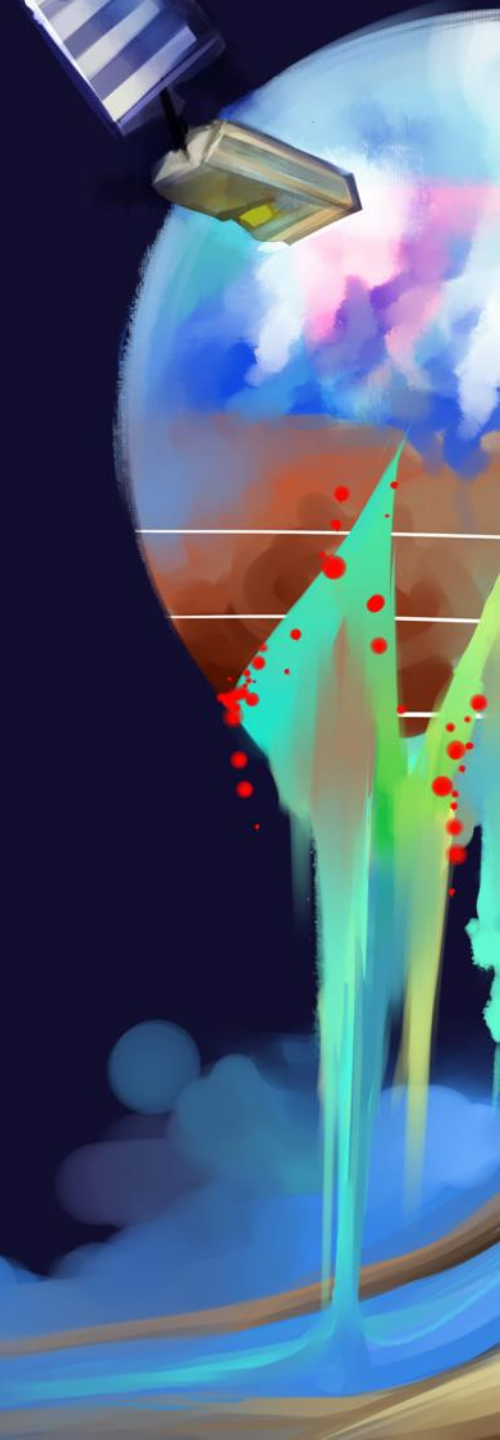
River discharge at the resolutions:
30 (~55km) vs. 5 arc minutes (~10km)



PCR-GLOBWB 2.0

A nice movie with output from PCR-GLOWB 2.0:
<https://goo.gl/3bPkwK>

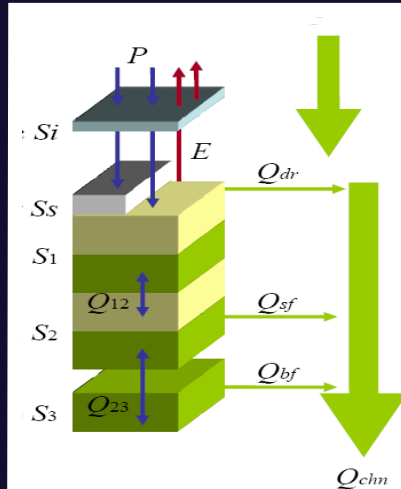
- 30 years of simulation at 5 arc minute resolution (~10 km) and daily time step.
- Shown are monthly averages of:
 - upper left: lower soil moisture (30-150 cm)
 - upper right: discharge (m³/s)
 - lower left: snow cover fraction
 - lower right: upper soil moisture (0-5 cm)
- This movie was rendered at the eScienceCenter in the project eWaterCycle.
 - <http://forecast.ewatercycle.org/>
 - <http://www.ewatercycle.nl/>



Refactoring model code

- PCR-GLOBWB 2.0 is re-written in Python (PCRaster framework).
 - Two possible resolutions: 5 arcmin and 30 arcmin
 - Suitable for regional models.

- A component based model.
- 4 main (hydrological) modules:
 - Meteo
 - Land surface
 - Groundwater
 - River / routing



- Example: Sub-module for land surface: **Land cover**
 - Loop through 2 or 4 land covers:
 - Natural: forest + grassland
 - Irrigation: irrPaddy + irrNonPaddy
 - Sub-modules for each land cover:
 - interception, snow, impArnoScheme, soilFluxes

```
class DeterministicRunner(DynamicModel):
    def __init__(self, configuration, model_time, initialState):
    def __init__(self):
    def dynamic(self):
        #current model timestep
        self.modelTime.update(self.currentTimeStep())

        #update model
        self.model.read_forcings()
        self.model.read_parameters()
        self.model.calculate_new_state()

        #do any needed reporting for this time step
        self.reporting.report()
```

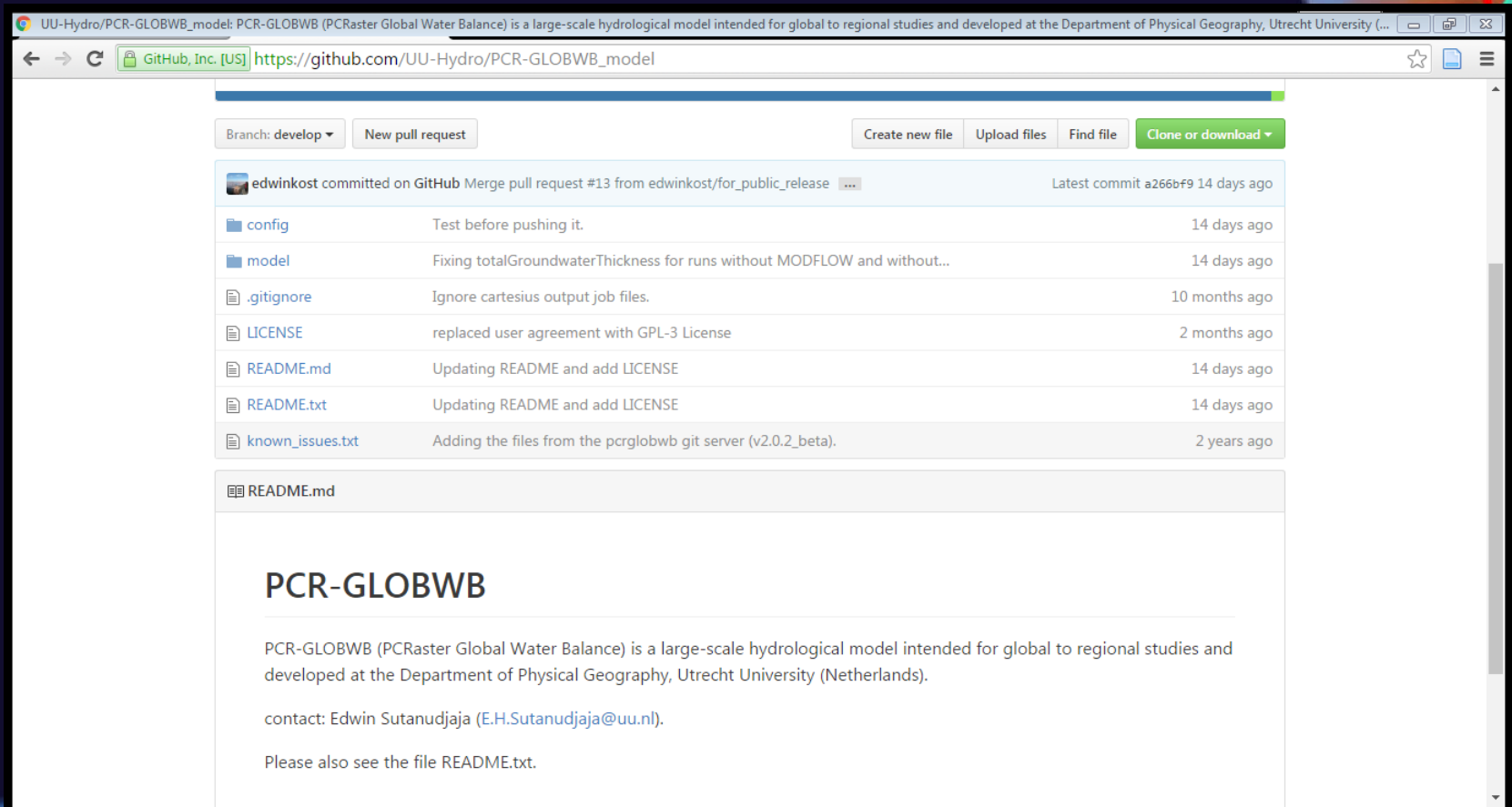
```
class PCRGlobWB(model_interface.ModelInterface):
    def __init__(self, configuration, model_time, \
    def calculate_new_state(self, report_water_balance):

        self.meteo.calculate_new_state(self._modelTime)
        self.landSurface.calculate_new_state(\
            self.meteo,\
            self.groundwater,\
            self.routing,\
            self._modelTime)
        self.groundwater.calculate_new_state(\
            self.landSurface,\
            self.routing,\
            self._modelTime)
        self.routing.calculate_new_state(\
            self.landSurface,\
            self.groundwater,\
            self._modelTime,\
            self.meteo)
```

Output in netcdf ; most input files also in netcdf (some still in pcraster maps).

PCR-GLOBWB 2.0 (open source)

- Model code of PCR-GLOBWB 2.0 is available from the following public repository: https://github.com/UU-Hydro/PCR-GLOBWB_model
- For example input and configuration files (Rhine catchment) we refer to: https://github.com/UU-Hydro/PCR-GLOBWB_input_example
- Our global input file (300 GB): <https://doi.org/10.5281/zenodo.1045338>



UU-Hydro/PCR-GLOBWB_model: PCR-GLOBWB (PCRaster Global Water Balance) is a large-scale hydrological model intended for global to regional studies and developed at the Department of Physical Geography, Utrecht University (...)

← → ↻ GitHub, Inc. [US] https://github.com/UU-Hydro/PCR-GLOBWB_model

Branch: develop ▾ New pull request

Create new file Upload files Find file Clone or download ▾

edwinkost committed on GitHub Merge pull request #13 from edwinkost/for_public_release Latest commit a266b9 14 days ago

config	Test before pushing it.	14 days ago
model	Fixing totalGroundwaterThickness for runs without MODFLOW and without...	14 days ago
.gitignore	Ignore cartesius output job files.	10 months ago
LICENSE	replaced user agreement with GPL-3 License	2 months ago
README.md	Updating README and add LICENSE	14 days ago
README.txt	Updating README and add LICENSE	14 days ago
known_issues.txt	Adding the files from the pcrglobwb git server (v2.0.2_beta).	2 years ago

README.md

PCR-GLOBWB

PCR-GLOBWB (PCRaster Global Water Balance) is a large-scale hydrological model intended for global to regional studies and developed at the Department of Physical Geography, Utrecht University (Netherlands).

contact: Edwin Sutanudjaja (E.H.Sutanudjaja@uu.nl).

Please also see the file README.txt.

script files and config files

- List of 'script' files and 'config' files:

```
edwinsut@intl.cartesius.surfsara.nl:/home/edwinsut/github/UU-Hydro/PCR-GLOBWB_model$  
ls model  
ETPFunctions.py          imagemean.pyx           pcrglobwb.py  
bmi.py                  landCover.py            pcrglobwb_v1  
bmiPcrglobwb.py         landSurface.py          reporting.py  
configuration.py        meteo.py                routing.py  
currTimeStep.py        modflow.py              setup.py  
debug_to_version_one.sh ncConverter.py           spinUp.py  
deterministic_runner.py oldcalc_framework.py    variable_list.py  
disclaimer.py           parallel_pcrglobwb_runner.py virtualOS.py  
groundwater.py          parameterSoilAndTopo.py  waterBodies.py  
edwinsut@intl.cartesius.surfsara.nl:/home/edwinsut/github/UU-Hydro/PCR-GLOBWB_model$  
edwinsut@intl.cartesius.surfsara.nl:/home/edwinsut/github/UU-Hydro/PCR-GLOBWB_model$  
ls config/  
05min_example  30min_example  
edwinsut@intl.cartesius.surfsara.nl:/home/edwinsut/github/UU-Hydro/PCR-GLOBWB_model$  
ls config/05min_example/  
setup_05min_global.ini  
edwinsut@intl.cartesius.surfsara.nl:/home/edwinsut/github/UU-Hydro/PCR-GLOBWB_model$
```

command to run the model:

Command to run the model:

python deterministic_runner.py *<configuration_file_name>*

```
sutanudjaja@cehs-hp:~/github/UU-Hydro/PCR-GLOBWB_model$ cd model
sutanudjaja@cehs-hp:~/github/UU-Hydro/PCR-GLOBWB_model/model$ python deterministic_runner.py
../config/05min_example/setup_05min_global.ini
```

PCR-GLOBWB (PCRaster Global Water Balance) Global Hydrological Model

Copyright (C) 2016, Ludovicus P. H. (Rens) van Beek, Edwin H. Sutanudjaja, Yoshihide Wada, Joyce H. C. Bosmans, Niels Drost, Inge E. M. de Graaf, Kor de Jong, Patricia Lopez Lopez, Stefanie Pessenteiner, Oliver Schmitz, Menno W. Straatsma, Niko Wanders, Dominik Wisser, and Marc F. P. Bierkens,
Faculty of Geosciences, Utrecht University, Utrecht, The Netherlands

This program comes with ABSOLUTELY NO WARRANTY

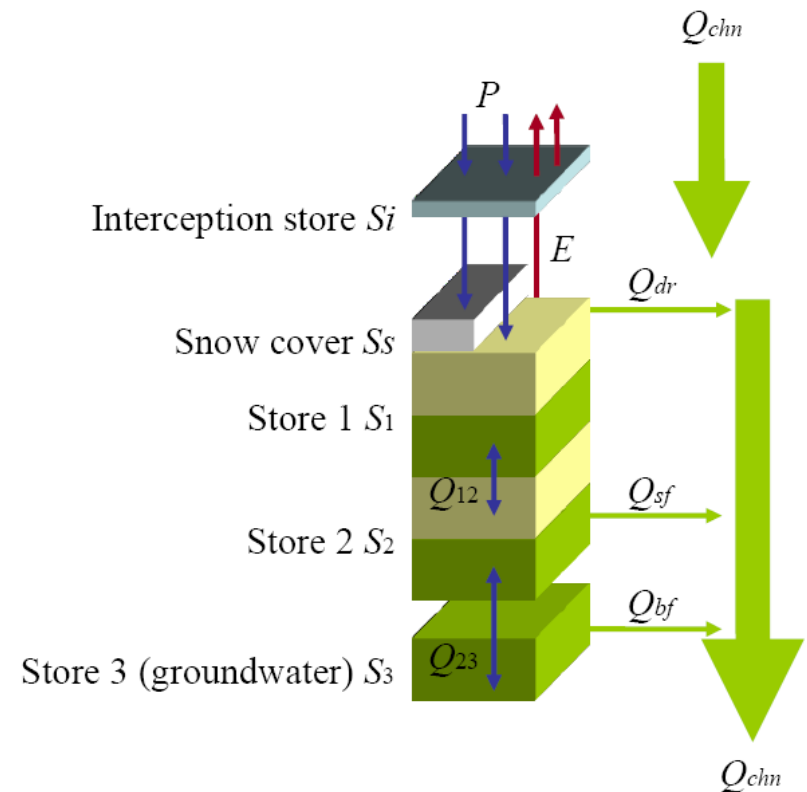
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See the LICENSE file for more details

script files

ETPFunctions.py	imagemean.pyx	pcrglobwb.py
bmi.py	landCover.py	pcrglobwb_v1
bmiPcrglobwb.py	landSurface.py	reporting.py
configuration.py	meteo.py	routing.py
currTimeStep.py	modflow.py	setup.py
debug_to_version_one.sh	ncConverter.py	spinUp.py
deterministic_runner.py	oldcalc_framework.py	variable_list.py
disclaimer.py	parallel_pcrglobwb_runner.py	virtualOS.py
groundwater.py	parameterSoilAndTopo.py	waterBodies.py

- **4 main hydrological modules:**

- meteo.py
 - ETPFunctions.py
- landSurface.py
 - parameterSoilAndTopo.py
 - landCover.py
- groundwater.py
- routing.py
 - waterBodies.py

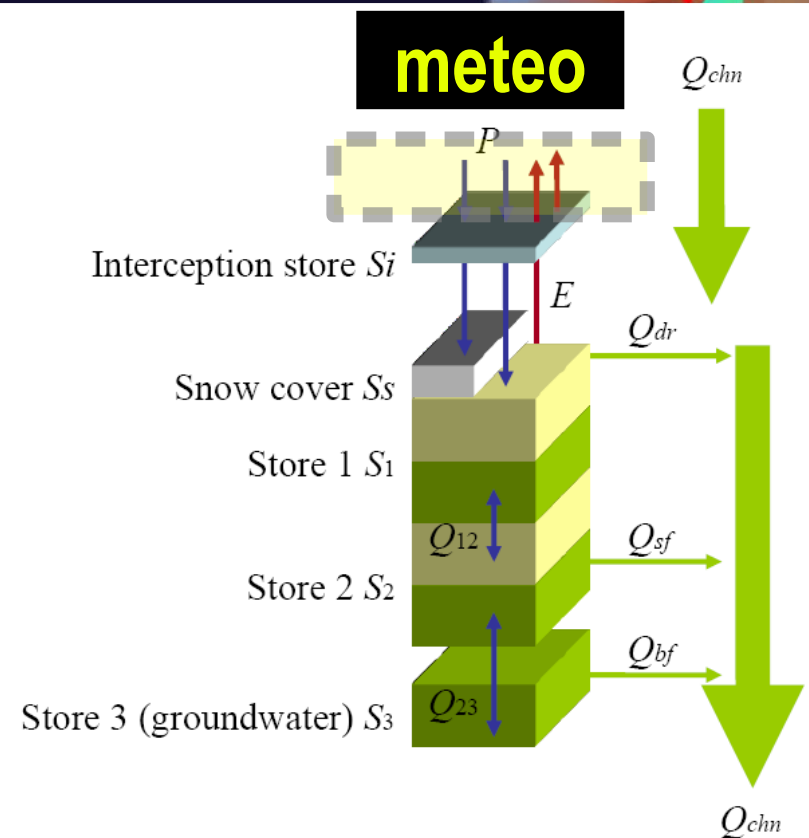


script files

ETPFunctions.py	imagemean.pyx	pcrglobwb.py
bmi.py	landCover.py	pcrglobwb_v1
bmiPcrglobwb.py	landSurface.py	reporting.py
configuration.py	meteo.py	routing.py
currTimeStep.py	modflow.py	setup.py
debug_to_version_one.sh	ncConverter.py	spinUp.py
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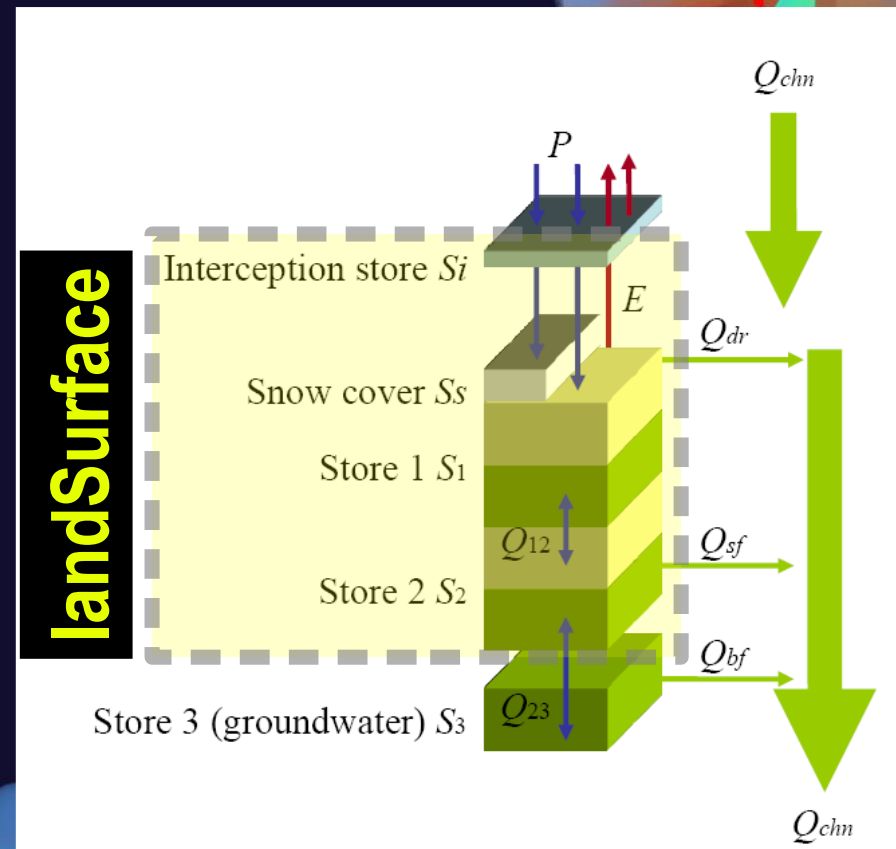
- **meteo.py**
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script files

ETPFunctions.py	imagemean.pyx	pcrglobwb.py
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groundwater.py	parameterSoilAndTopo.py	waterBodies.py

- **4 main hydrological modules:**
 - meteo.py
 - ETPFunctions.py
 - **landSurface.py**
 - parameterSoilAndTopo.py
 - **landCover.py** -called 4 times
 - groundwater.py
 - routing.py
 - waterBodies.py

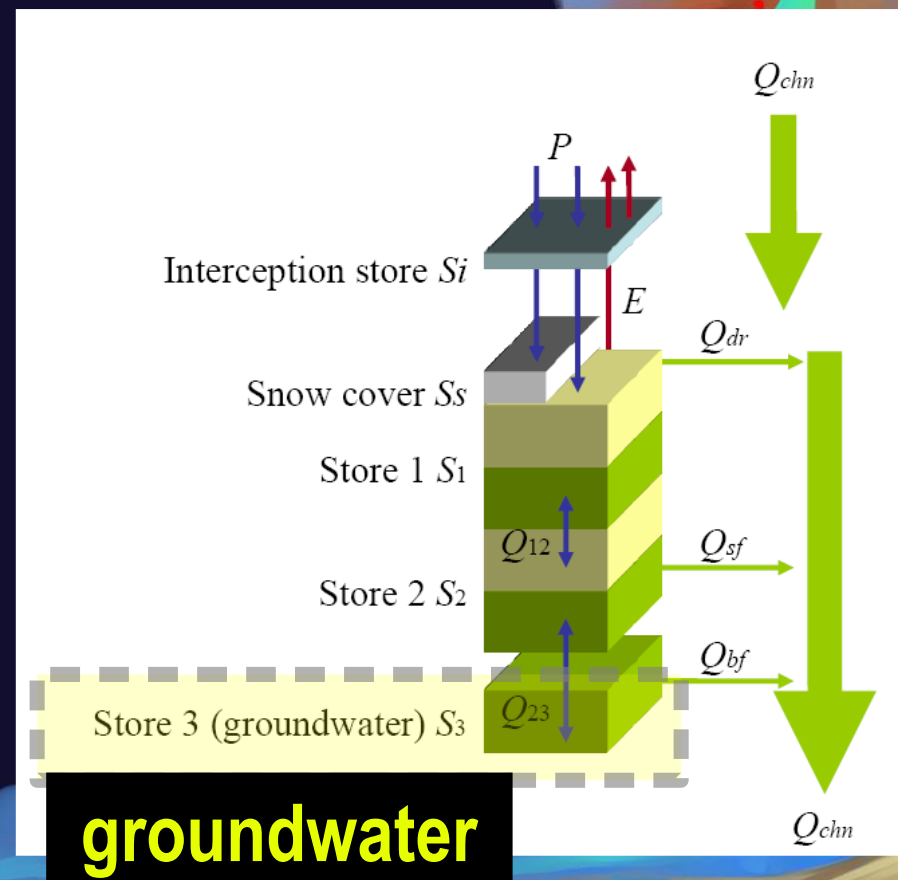


script files

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bmi.py	landCover.py	pcrglobwb_v1
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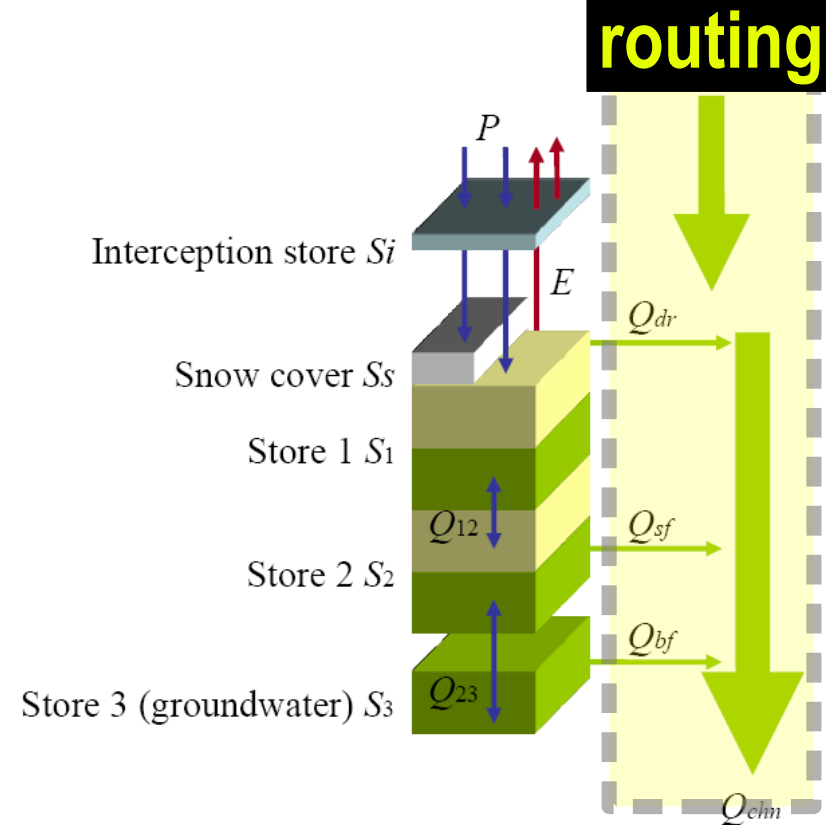


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- groundwater.py
- **routing.py**
 - waterBodies.py



ini files

```
edwinsut@int1.cartesius.surfsara.nl:/home/edwinsut/github/UU-Hydro/PCR-GLOBWB_model$  
edwinsut@int1.cartesius.surfsara.nl:/home/edwinsut/github/UU-Hydro/PCR-GLOBWB_model$  
ls config/  
05min_example  30min_example  
edwinsut@int1.cartesius.surfsara.nl:/home/edwinsut/github/UU-Hydro/PCR-GLOBWB_model$  
ls config/05min_example/  
setup_05min_global.ini  
edwinsut@int1.cartesius.surfsara.nl:/home/edwinsut/github/UU-Hydro/PCR-GLOBWB_model$
```

- What should be defined in a configuration file (*.ini)?
 - Input file locations or input values
 - Forcing: precipitation, temperature, etc.
 - Water demand files
 - Model parameters
 - Initial conditions.
 - Other configuration: startTime, endTime, cloneMap, landmask, etc.
 - Which output variables do you want to report?

ini files

- Structure of a configuration file

```
1
2  [+][globalOptions]
36  L
37  [+][meteoOptions]
49  [+][meteoDownscalingOptions] # only for 5 arcmin runs
71  L
72  [+][landSurfaceOptions]
119 [+][forestOptions]
163 [+][grasslandOptions]
207 [+][irrPaddyOptions]
250 [+][irrNonPaddyOptions]
293 L
294 [+][groundwaterOptions]
337 L
338 [+][routingOptions]
407 L
408 [+][reportingOptions]
432 L
433 [-][mergingOutputOptions] # only for parallel runs
434 L
```

Location of PCR-GLOBWB input data

- All input data are provided in the global coverage.
- On speedy/rapid (our servers): /data/hydroworld/
- On Cartesius: /projects/0/dfguu/data/hydroworld/
- The fixed/stable/release version is available on <https://doi.org/10.5281/zenodo.1045338> (300 GB)
- A small input example (Rhine-Meuse) is also provided on https://github.com/UU-Hydro/PCR-GLOBWB_input_example

```
sutan101@speedy:/data/hydroworld$  
sutan101@speedy:/data/hydroworld$ ls  
basedata  forcing  others  parameterizationscripts  PCRGLOBWB10  PCRGLOBWB20  synchronize.sh  
sutan101@speedy:/data/hydroworld$  
sutan101@speedy:/data/hydroworld$ cd PCRGLOBWB20/  
sutan101@speedy:/data/hydroworld/PCRGLOBWB20$  
sutan101@speedy:/data/hydroworld/PCRGLOBWB20$ ls -l  
total 12  
drwxrwsr-x 8 sutan101 hydroworld 4096 Mar 12 14:28 input30min  
drwxrwsr-x 9 straa005 hydroworld 4096 Feb  4 14:24 input5min  
drwxrwsr-x 4 straa005 hydroworld 4096 Oct  4 2013 scripts  
sutan101@speedy:/data/hydroworld/PCRGLOBWB20$ cd input5min/  
sutan101@speedy:/data/hydroworld/PCRGLOBWB20/input5min$ ls -l  
total 28  
drwxrwsr-x 2 straa005 hydroworld 4096 Aug 27 2013 global  
drwxrwsr-x 3 straa005 hydroworld 4096 Mar  5 17:32 groundwater  
drwxr-sr-x 3 beek0120 hydroworld 4096 Dec 24 16:54 hydeLandCover  
drwxrwsr-x 6 straa005 hydroworld 4096 Mar 11 11:14 landCover  
drwxrwsr-x 5 straa005 hydroworld 4096 Mar 27 00:38 landSurface  
drwxrwsr-x 3 straa005 hydroworld 4096 Mar 11 10:18 meteo  
drwxrwsr-x 4 straa005 hydroworld 4096 Mar  7 14:26 routing  
sutan101@speedy:/data/hydroworld/PCRGLOBWB20/input5min$
```

globalOptions in the ini file

```
[globalOptions]

# Set the input directory map in an absolute path.
# - The input forcing and parameter directories will be relative to this.
inputDir = /data/hydroworld/

# Set the output directory in an absolute path.
outputDir = /scratch/edwin/test_3_layers/

# Map of clone (must be provided in PCRaster maps)
# - Spatial resolution and coverage are based on this map:
cloneMap = others/Columbia/Columbia05min.clone.map

# The area/landmask of interest:
landmask = others/Columbia/Columbia05min.landmask.map
# If None, area/landmask is limited for cells with ldd value.

# start and end simulation period
startTime = 1979-01-01
endTime   = 2010-12-31
# Format: YYYY-MM-DD
timeStep = 1.0
timeStepUnit = day
# PS: The current model still runs of the daily time step.

# spinning up options:
# Note: for the purpose of DA, there should be no SpinUp
maxSpinUpsInYears = 5
minConvForTotlSto = 0.5
minConvForSoilSto = 0.5
minConvForGwatSto = 0.5
minConvForChanSto = 0.5
```

- cloneMap: all cells have TRUE values
 - Model area and model resolution.
 - Border coordinates must be in integer.
- landmask: only TRUE at your area of interest

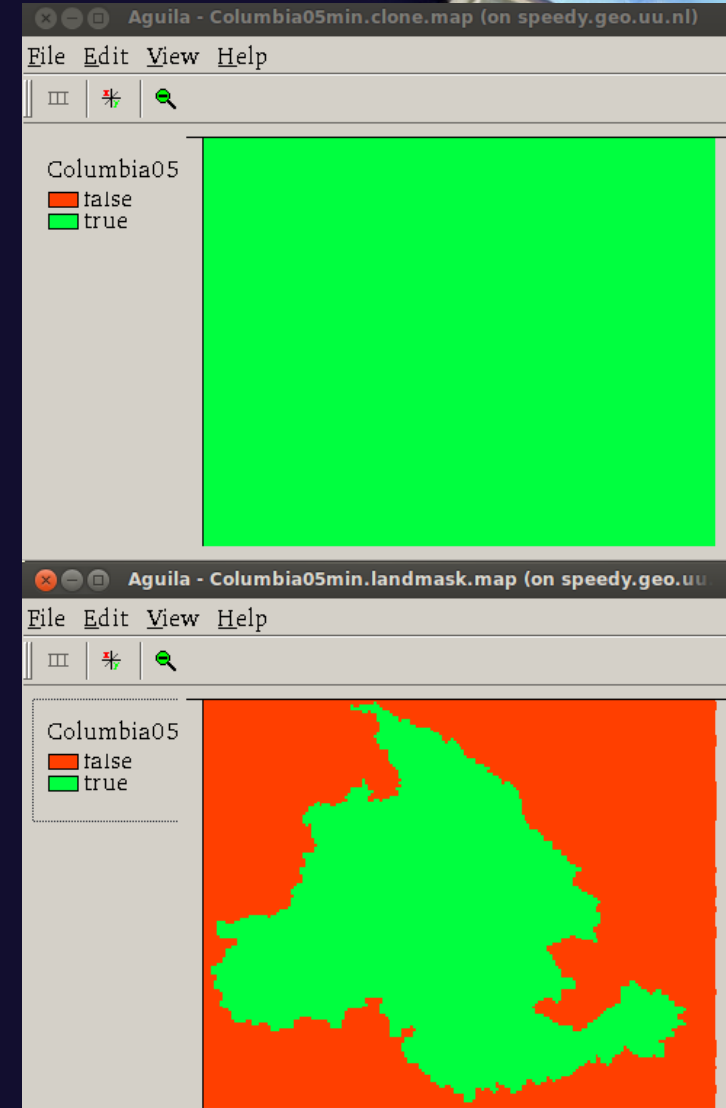


Fig. CloneMap & landmask maps at 5 arc-min for Columbia river basin.

meteoOptions in the ini file

[meteoOptions]

```
# Set the forcing temperature and precipitation files (relative to inputDir)
temperatureNC = forcing/ERA-Interim-GPCPCorrected/temperature_ERA_Interim_1979to2010.nc
precipitationNC = forcing/ERA-Interim-GPCPCorrected/precipitation_ERA_Interim_GPCPCorrected_1979to2010.nc

# Method to calculate referencePotETP (reference potential evaporation+transpiration)
referenceETPotMethod = Hamon
# options are "Hamon" and "Input" ; If "Input", the netcdf input file must be given:
refETPotFileNC = None

...
```

[meteoDownscalingOptions]

```
# These options are only used and tested for 5 min runs which are using 30 min forcing

downscalePrecipitation = True
downscaleTemperature = True
downscaleReferenceETPot = True
...

# lapse rates:
temperLapseRateNC = PCRGLOBWB20/input5min/meteo/downscalingFrom30ArcMin/temperature_slope.nc
precipLapseRateNC = PCRGLOBWB20/input5min/meteo/downscalingFrom30ArcMin/precipitation_slope.nc
```

- Precipitation and temperature files must be in the netcdf format.
- You can also use “absolute path” (while defining your own forcing).
- Downscaling options are only used (and tested) for 5 min runs.

landSurfaceOptions in the ini file

```
[landSurfaceOptions]
...

numberOfUpperSoilLayers = 2
# 2 soil layers: 30 cm and 120 cm; 3 soil layers: 5 cm, 25 cm and 120 cm

includeIrrigation = True
# if True, there are four land cover types defined: forest,grassland,irrPaddy,irrNonPaddy
# if False, two (natural) land cover types defined: forest,grassland
historicalIrrigationArea = PCRGLOBWB20/input30min/landSurface/waterDemand/irrigated_areas/irrigationArea30ArcMin.nc

includeDomesticWaterDemand = True
includeIndustryWaterDemand = True
domesticWaterDemandFile = PCRGLOBWB20/input30min/landSurface/waterDemand/domesticWaterDemand30ArcMin.nc
industryWaterDemandFile = PCRGLOBWB20/input30min/landSurface/waterDemand/industryWaterDemand30ArcMin.nc

limitAbstraction = False
```

- 3 soil layers are mainly used for the eWaterCycle project.
- IncludeIrrigation = True → 4 land covers = forest, grassland, irrPaddy & irrNonPaddy
- limitAbstraction = False → allowing other abstraction sources (e.g. fossil water)

<landCover>Options (2 layer model)

```
[forestOptions]
...
# snow module properties
snowModuleType      = Simple
freezingT           = -0.0
degreeDayFactor      = 0.0025
...

# land cover properties:
landCoverMapsNC      = PCRGLOBWB20/input30min/landCover/forest/forestProperties.nc
cropCoefficientNC    = PCRGLOBWB20/input30min/landCover/forest/Global_CropCoefficientKc-Forest_30min.nc
interceptCapNC       = PCRGLOBWB20/input30min/landCover/forest/interceptCapInputForest366days.nc
coverFractionNC      = PCRGLOBWB20/input30min/landCover/forest/coverFractionInputForest366days.nc

# initial conditions (for two layer model):
interceptStorIni     = 0.0
snowCoverSWEIni      = 0.0
snowFreeWaterIni     = 0.0
topWaterLayerIni     = 0.0
storUpIni            = 0.0
storLowIni           = 0.0
interflowIni         = 0.0
...

[grasslandOptions]
[irrPaddyOptions]
[irrNonPaddyOptions]
```

- Each landCover has each own field.
- Initial Conditions must be defined in each land cover.
- The example above is for a **2-layer model**.
- For a 3-layer model, initial conditions are different (i.e. there are more states).

<landCover>Options (3 layer model)

```
[forestOptions]
...
# snow module properties
snowModuleType      = Simple
freezingT           = -0.0
degreeDayFactor      = 0.0025
...

# land cover properties:
landCoverMapsNC      = PCRGLOBWB20/input30min/landCover/forest/forestProperties.nc
cropCoefficientNC    = PCRGLOBWB20/input30min/landCover/forest/Global_CropCoefficientKc-Forest_30min.nc
interceptCapNC       = PCRGLOBWB20/input30min/landCover/forest/interceptCapInputForest366days.nc
coverFractionNC      = PCRGLOBWB20/input30min/landCover/forest/coverFractionInputForest366days.nc

# initial conditions (for three layer model):
interceptStorIni     = 0.0
snowCoverSWEIni      = 0.0
snowFreeWaterIni     = 0.0
topWaterLayerIni     = 0.0
storUpp000005Ini     = 0.0
storUpp005030Ini     = 0.0
storLow030150Ini     = 0.0
interflowIni         = 0.0
...

[grasslandOptions]
[irrPaddyOptions]
[irrNonPaddyOptions]
```

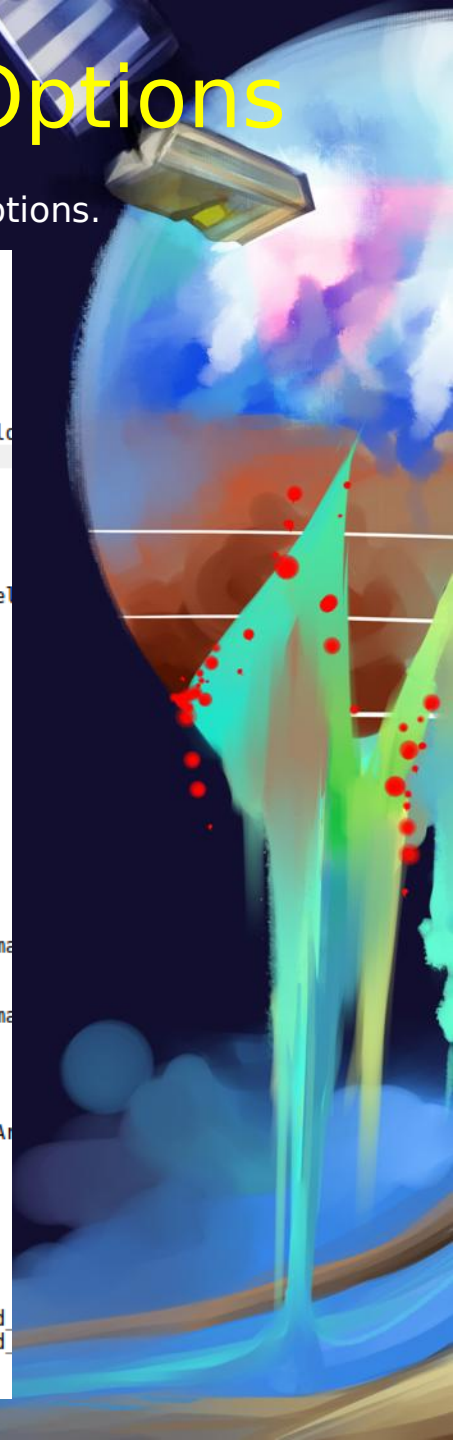
- Each landCover has each own field.
- Initial Conditions must be defined in each land cover.
- The example above is for a **3-layer model**.
- For a 2-layer model, initial conditions are different (i.e. there are fewer states).

groundwaterOptions & routingOptions

- Both 2-layer & 3-layer models have the same groundwaterOptions & routingOptions.

```
[groundwaterOptions]
groundwaterPropertiesNC = PCRGLOBWB20/input5min/groundwater/groundwaterProperties5ArcMin.nc
...
# initial conditions:
storGroundwaterIni      = /projects/0/aqueduct/users/edwinsut/pcrglobwb_runs_2016_oct_nov/pcrglobwb_4_land
...

[routingOptions]
lddMap      = PCRGLOBWB20/input5min/routing/lddsound_05min.map
cellAreaMap = PCRGLOBWB20/input5min/routing/cellsize05min.correct.map
gradient     = /projects/0/dfguu/users/edwin/data/floodplain_05arcmin_world_final/based_on_daily_runoff/map/channel
...
# manning coefficient
manningsN   = 0.04
...
# manning coefficient for floodplain
floodplainManningsN = 0.07
...
# routing method: (options are kinematicWave, simplifiedKinematicWave and accuTravelTime)
routingMethod = accuTravelTime
...
# Option for flood plain simulation
dynamicFloodPlain = True
...
# constant channel width (optional)
constantChannelWidth = /projects/0/dfguu/users/edwin/data/floodplain_05arcmin_world_final/based_on_daily_runoff/ma
...
# constant channel depth
constantChannelDepth = /projects/0/dfguu/users/edwin/data/floodplain_05arcmin_world_final/based_on_daily_runoff/ma
...
# lake and reservoir parameters
waterBodyInputNC = PCRGLOBWB20/input5min/routing/reservoirs/waterBodiesFinal_version15Sept2013/maps/waterBodies5A
onlyNaturalWaterBodies = False
...
# composite crop factors for WaterBodies:
cropCoefficientWaterNC = PCRGLOBWB20/input30min/routing/cropCoefficientForOpenWater.nc
...
# initial conditions:
waterBodyStorageIni      = /projects/0/aqueduct/users/edwinsut/pcrglobwb_runs_2016_oct_nov/pcrglobwb_4_land
channelStorageIni        = /projects/0/aqueduct/users/edwinsut/pcrglobwb_runs_2016_oct_nov/pcrglobwb_4_land
...
```



reportingOptions

[reportingOptions]

```
# output files that will be written in the disk in netcdf files:
# - daily resolution
outDailyTotNC = discharge
# - monthly resolution
outMonthAvgNC =
discharge,dynamicFracWat,surfaceWaterStorage,interceptStor,snowFreeWater,snowCoverSWE,topWaterLayer,storUppTotal,storLowTotal,storG
roundwater,storGroundwaterFossil,totalActiveStorageThickness,totalWaterStorageThickness,satDegUpp,satDegLow,channelStorage,waterBod
yStorage
outMonthEndNC =
storGroundwater,storGroundwaterFossil,waterBodyStorage,channelStorage,totalWaterStorageThickness,totalActiveStorageThickness
outMonthTotNC =
totalEvaporation,waterBodyActEvaporation,actualET,irrPaddyWaterWithdrawal,irrNonPaddyWaterWithdrawal,domesticWaterWithdrawal,indust
ryWaterWithdrawal,livestockWaterWithdrawal,precipitation,gwRecharge,runoff,totalRunoff,baseflow,directRunoff,interflowTotal,totalGr
oundwaterAbstraction,desalinationAbstraction,surfaceWaterAbstraction,nonFossilGroundwaterAbstraction,fossilGroundwaterAbstraction,i
rrGrossDemand,nonIrrGrossDemand,totalGrossDemand,nonIrrWaterConsumption,nonIrrReturnFlow
# - annual resolution
outAnnuaAvgNC = None
outAnnuaEndNC = None
outAnnuaTotNC = None
outMonthMaxNC = None
outAnnuaMaxNC = None

# netcdf format and zlib setup
formatNetCDF = NETCDF4
zlib = True
```

- Output variables can be reported in **Daily**, **Monthly** and **Annual** resolution; in cumulative (**Tot**), average (**Avg**), end of a period (**End**), or maximum (**Max**) values.

list of output variables

- See the model script file “variable_list.py” (inside the “model” folder).

```
# totLandSurfaceActuaET
pcrglobwb_variable_name = 'totLandSurfaceActuaET'
netcdf_short_name[pcrglobwb_variable_name] = 'land_surface_actual_evaporation'
netcdf_unit[pcrglobwb_variable_name] = 'm.day-1'
netcdf_monthly_total_unit[pcrglobwb_variable_name] = 'm.month-1'
netcdf_yearly_total_unit[pcrglobwb_variable_name] = 'm.year-1'
netcdf_long_name[pcrglobwb_variable_name] = 'total_actual_evaporation_and_transpiration_at_land_surface'
description[pcrglobwb_variable_name] = None
comment[pcrglobwb_variable_name] = 'Not including water bodies. Values given are over the entire cell area.'
latex_symbol[pcrglobwb_variable_name] = None

# fractionLandSurfaceET
pcrglobwb_variable_name = 'fractionLandSurfaceET'
netcdf_short_name[pcrglobwb_variable_name] = 'land_surface_evaporation_fraction'
netcdf_unit[pcrglobwb_variable_name] = '1'
netcdf_monthly_total_unit[pcrglobwb_variable_name] = None
netcdf_yearly_total_unit[pcrglobwb_variable_name] = None
netcdf_long_name[pcrglobwb_variable_name] = 'ratio_between_actual_and_potential_values_of_evaporation_and_transpiration_at_land_surface'
description[pcrglobwb_variable_name] = None
comment[pcrglobwb_variable_name] = 'Not including water bodies.'
latex_symbol[pcrglobwb_variable_name] = None

# interceptStor
pcrglobwb_variable_name = 'interceptStor'
netcdf_short_name[pcrglobwb_variable_name] = 'interception_storage'
netcdf_unit[pcrglobwb_variable_name] = 'm'
netcdf_monthly_total_unit[pcrglobwb_variable_name] = None
```


mergingOutputOptions

- Only used for global 5 arcmin runs that are parallelized.
- List of output variables that are reported and will be merged.

```
[reportingOptions]
# output files that will be written in the disk in netcdf files:
# - daily resolution
outDailyTotNC = discharge
# - monthly resolution
outMonthAvgNC = totalEvaporation,gwRecharge,totalRunoff,totalGroundwaterAbstraction
outMonthEndNC =
storGroundwater,storGroundwaterFossil,waterBodyStorage,channelStorage,totalWaterStorageThickness,totalActiveStorageThickness
outMonthTotNC = None
# - annual resolution
outAnnuaAvgNC = None
outAnnuaEndNC = None
outAnnuaTotNC = None
outMonthMaxNC = None
outAnnuaMaxNC = None

[mergingOutputOptions]
# output variables/files that will be merged:
outDailyTotNC = None
outMonthAvgNC = totalEvaporation,gwRecharge
outMonthEndNC = None
outMonthTotNC = None
outAnnuaAvgNC = None
outAnnuaEndNC = None
outAnnuaTotNC = None
outMonthMaxNC = None
outAnnuaMaxNC = None
```

check your output / model results

```
[screen 0: bash]
File Edit View Search Terminal Help

Left      File      Command      Options      Right
<- /scratch/edwin/test_3_layers .[^]> <- /scratch/edwin/test_3_layers/netcdf .[^]>
'n        Name      Size      Modify time
/..        UP--DIR May 15 14:21
/log       4096 May 12 18:46
/netcdf    4096 May 12 18:46
/scripts   4096 May 12 18:46
/states    4096 May 12 18:46
/tmp       4096 May 12 18:46

/tmp 934G/666G (140%)

disChanWaterBody_dailyTot.nc 15089K May 12 18:50
disChanWaterBody_monthAvg.nc 416928 May 12 18:49
discharge_annuaAvg.nc 2180 May 12 18:46
discharge_dailyTot.nc 15089K May 12 18:50
discharge_monthAvg.nc 416916 May 12 18:49
fracNonFossilG~r_monthAvg.nc 416952 May 12 18:49
fracSurfaceWater_monthAvg.nc 416928 May 12 18:49
fracUnmetDemand_monthAvg.nc 416928 May 12 18:49
gwRecharge_annuaTot.nc 2180 May 12 18:46
gwRecharge_monthTot.nc 416916 May 12 18:49
infiltration_annuaTot.nc 2180 May 12 18:46
infiltration_monthTot.nc 416916 May 12 18:49
interceptStor_monthAvg.nc 416928 May 12 18:49
interflowTotal_annuaTot.nc 2192 May 12 18:46

interceptStor_monthAvg.nc 934G/666G (140%)

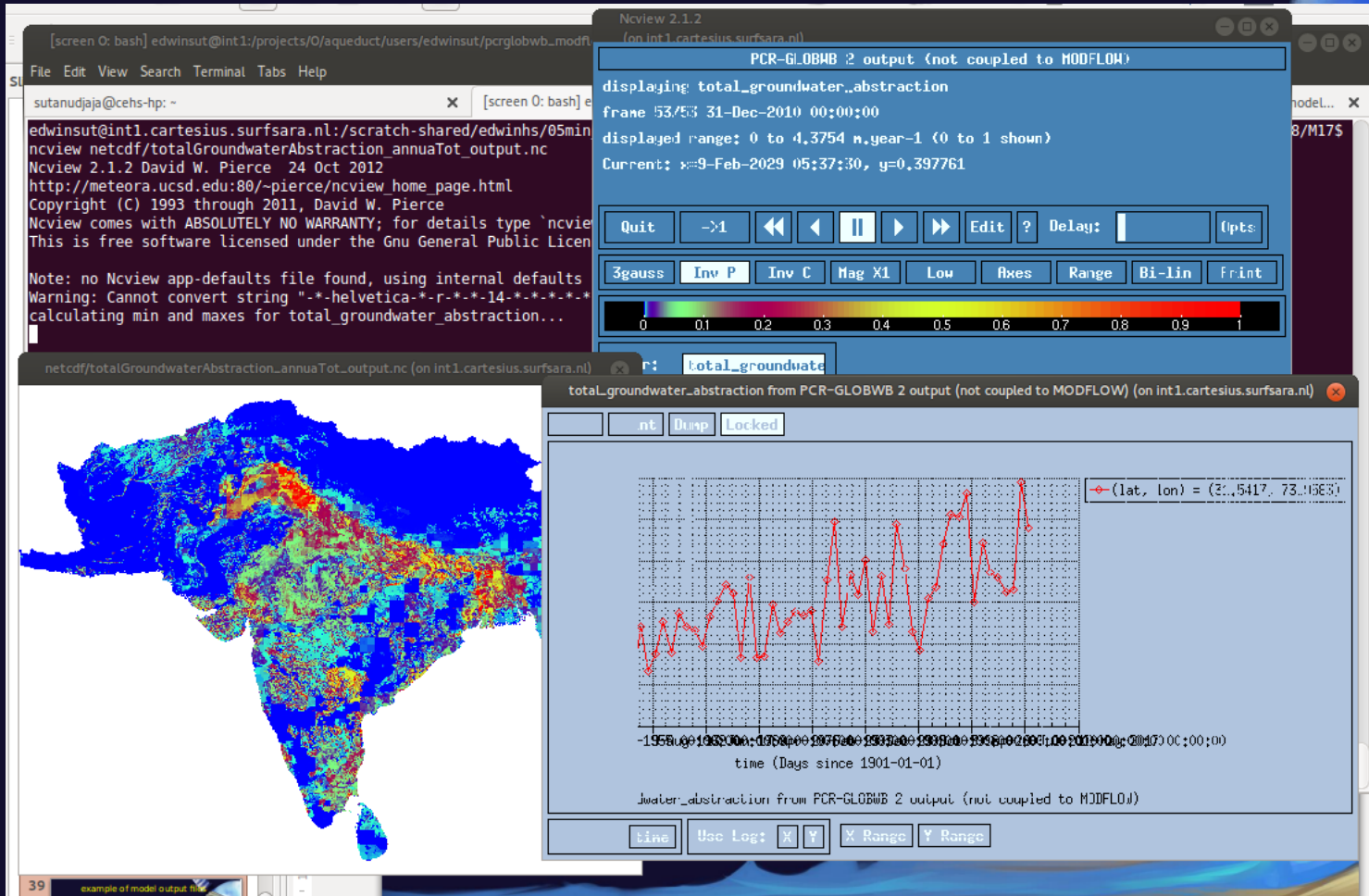
Hint: % macros work even on the command line.
sutan101@speedy:/scratch/edwin/test_3_layers/netcdf$
```

1Help 2Menu 3View 4Edit 5Copy 6RenMov 7Mkdir 8Delete 9PullDn 10Quit

- Contents of the outputDir: **log** (including backup of the ini file); **netcdf** output ; backup **scripts** used to run the model ; model **states** at the end of each year ; **tmp** directory (used during the resampling process)
- We will explore these during the exercises.
- To visualize a certain netcdf output variable: **ncview <name_of_netcdf_file>**

example of model output files

- Output files are mainly in netcdf files (some are in pcraster maps)
- To visualize a netcdf file: **ncview <name_of_netcdf_file>**



Questions?



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