pyg2p 2.0

Version	2.0	
Author	Domenico Nappo (domenico.nappo(at)gmail.com)	
Release Date	2016-01	

Table of Contents

pyg2p User Manual	1
Introduction	
Usage	3
Grabbing information from GRIB files	3
Input arguments	
Check output maps	
Interpolation modes	
GRIB API interpolation methods	6
Scipy interpolation methods	
OutMaps configuration	
Aggregation	
Average	
Accumulation	
Correction	
How to write formulas	
Which geopotential file will be read?	
Notes about formulas	14
Conversion	17
Logging	18
Comparing Test Suite	
Writing test.xml file	19
Writing the commands.txt file	
Sample Output	20
Appendix A - Execution XML files Examples	22
Appendix B – mapping between grib2pcraster and pyg2p	
Differences in glossary	
Mapping variables	
Known parameters mapping with conversions	28
Execution parameters mapping	
Difference in dealing with multiresolution	
Differences in Correction	
Differences in Aggregation	
Differences in Interpolation	
Appendix C – Glossary	31

Introduction

pyg2p is a tool to convert GRIB files into PCRASTER maps.

It reads georeferenced variables from GRIB version 1 and 2 files and it produces PCRaster maps, after some manipulation steps. Execution parameters are passed via command line arguments and JSON template files.

Manipulations are completely configurable in execution JSON files and are:

- conversion (formulas are to configure in parameters.json)
- correction
- aggregation (accumulation or average).

The interpolation mode is also configurable and it's needed to resample the input GRIB grid to the PCRaster target grid (which is determined by latitude and longitudes maps).

Installation

With pip tool:

1. In a python2.7 virtualenv, execute

pip install git+https://bitbucket.org/nappodo/pyg2p.git

Note: Your bitbucket account must have read access to repository.

Pip will download python package, install dependencies and will launch script to test environment.

Manual

- 1. Download last version from bitbucket.org (you have to grant permissions to repository): https://bitbucket.org/nappodo/pyg2p/get/<version>.tar.gz
- 2. Decompress file in <target directory> using tar xzvf pyg2p_v.<version>.tar.gz.
- 3. Install dependencies (see table below).

Some packages need additional requirements such as C libraries properly installed.

Software Requirement		
Python 2.7	Execution environment	
Python packages		
NumPy>=1.10.1	Scientific packages for array operations and reading/writing	
Numexpr	encoded files. They depends on several C libraries that must exist on your system before to proceed.	
SciPy>=0.16	— on your system before to proceed.	

gribapi* GDAL python**	<pre>\$pip install numpy>=1.10.1 -\$pip install numexpr \$pip install scipy>=0.16.0 \$pip install GDAL</pre>	
	* You may have troubles while installing right GDAL python package. If you have GDAL C/C++ libraries already installed on your system, you might need to install GDAL python matching the C GDAL version. ** For gribapi python package, please refer to GRIB API documentation.	
dask[array] dask[bag]	Light packages for parallelization (used in interpolation): \$ pip install toolz	
toolz	<pre>\$ pip install dask[array] \$ pip install dask[bag]</pre>	
xmljson (optional)	To convert old XML pyg2p v1 to v2 JSON format: \$pip install xmljson	
memory_profiler (optional)	Memory usage information for test suite. \$pip install memory_profiler	

Migrating from v 1.x

If you migrate from a pyg2p version 1.x, you can convert existing XML configuration files to json (execution command files, parameters.xml and geopotentials.xml) and also convert intertables (only grib api methods)

- 1. Copy default configuration (from sources) to your .pyg2p/ user folder, with command: \$pyg2p.py -P
- 2. Copy all your old XML configuration into a folder ./xml_backup. You can create different subfolders for convenience because conversion will be recursive.
- 3. Convert files (recursively) from XML to JSON with command: \$pyg2p.py -C ./xml_backup
- 4. Copy parameters.json and geopotentials.json from ./xml_backup into user folder ~/.pyg2p/
- 5. Copy your geopotential files to ~/.pyg2p/geopotentials/ folder.

You can also convert existing interpolation tables made with pyg2p version 1.x GRIB_API methods to be compliant with new format. New files will be copied under ~/.pyg2p/intertables/ folder and intertables.json configuration will be updated during conversion.

```
$pyg2p.py -z <path_to_convert>
```

Note: Intertables for rotated grids are not valid anymore so they won't be converted.

Configuration

There are several things to configure in pyg2p but default configuration is enough for most of conversion tasks. Since version 2.x, pyg2p use and store configuration into user folder ~/.pyg2p/. Intertables are read/write from ~/.pyg2p/intertables/ and geopotentials (for correction) are read from ~/.pyg2p/geopotentials/.

To copy default pyg2p configuration into user configuration folder execute this command: \$pyg2p -P

The command will copy main configuration and some existing geopotentials file for correction steps.

You have to download default intertables and copy into ~/.pyg2p/intertables from ECMWF ftp server or manually copy default intertables instead from <path_to_pyg2p>/configuration/intertables/ to ~/.pyg2p/intertables.

Advanced configuration

Adding a parameter to parameters.json

If you are extracting a parameter with shortName xy from a grib file that is not configured into ~/.pyg2p/parameters.json file, you need to add an element (a dictionary) to "Parameter" list:

You can add more than a conversion element with different ids and functions. You will use shortname and conversion id for the execution JSON template.

Note: Aware the syntax of conversion functions. They must start with x= followed by the actual conversion formula where x is the value to convert. Units are only used for logging.

Adding a geopotential for correction

If the input grib file has a geopotential message, pyg2p will use it for correction. Otherwise, it will read the file configured into \sim /.pyg2p/geopotentials.json.

To add a geopotential GRIB file to pyg2p configuration, use this command:

```
$pyg2p -g path_to_geopotential_grib
```

This will copy the file to ~/.pyg2p/geopotentials/ folder and will update geopotentials.json with the new item.

Interpolation tables

Interpolation tables are read from ~/.pyg2p/intertables folder.

If the table is missing, it will create it into same folder for future interpolations.

Depending on source and target grids size, and on interpolation method, table creation can take from minutes to days. To speed up interpolation table creation, use parallel option -X to have up to x6 speed gain.

Execution templates

Execution templates are JSON files that you will use to configure a conversion. You will pass path to the file to pyg2p with command line option'-c'.

Most of options can be both defined in this JSON file and from command line. Note that command line options overwrite JSON template.

If you have a large set of conversions for same parameters, where you change start step, end step and perturbation number parameter filters, you would use one JSON file to define parameter, interpolation and correction and you can change other parameters from command line.

Here some examples of JSON commands files:

```
{
  "Execution": {
    "@name": "Octahedral test 1",
    "Aggregation": {
        "@step": 24,
        "@type": "average"
    },
    "OutMaps": {
        "@cloneMap": "/dataset/maps/europe5km/dem.map",
        "@ext": 1,
```

There are four sections of configuration.

Aggregation

Defines the aggregation method and step. Method can be accumulation, average or instantaneous.

OutMaps

Here you define interpolation method and paths to coordinates PCRaster maps, output unit time, the clone map etc.

Interpolation

This is a subelement of OutMaps. Here you define interpolation method (see later for details), paths to coordinates maps.

Parameter

In this section, you configure the parameter to select by using its shortName, as stored in GRIB file. You also configure conversion with *applyConversion* property set to a conversion id. Parameter *shortName* must be already configured in ~/.pyg2p/parameters.json along with conversion ids.

If you need to apply correction based on DEM files and geopotentials, you can configure formulas and the path to DEM map.

Path configuration

You can use variables in JSON files to define paths. Variables can be configured in .conf files under \sim /.pyg2p/ folder.

/home/domenico/.pyg2p/myconf.conf

```
EUROPE_MAPS=/dataset/maps/europe5km

DEM_MAP=/dataset/maps/dem05.map

EUROPE_DEM=/dataset/maps/europe/dem.map

EUROPE_LAT=/dataset/maps/europe/lat.map

EUROPE_LON=/dataset/maps/europe/long.map
```

Usage in JSON command file:

```
"Aggregation": {
    "@step": 24,
    "@type": "average"
},
"OutMaps": {
    "@cloneMap": "{EUROPE MAPS}/lat.map",
    "@ext": 1,
    "@fmap": 1,
    "@namePrefix": "pT24",
    "Interpolation": {
        "@latMap": "{EUROPE MAPS}/lat.map",
        "@lonMap": "{EUROPE_MAPS}/long.map",
        "@mode": "grib nearest"
"Parameter": {
    "@applyConversion": "k2c",
    "@correctionFormula": "p+gem-dem*0.0065",
    "@demMap": "{DEM_MAP}",
    "@gem": "(z/9.81)*0.0065",
    "@shortName": "2t"
```

Full list of options

Attribute/Tag	Details
name	The short name of the parameter, as it is in the grib file. The application use this to select messages.
Parameter	See relative table
Aggregation	See relative table
OutMaps	See relative table

Parameter tag

Attribute/Tag	Details
shortName	The parameter short name, as it is in the grib file. Must be configured in the parameters.json file, otherwise the application exits with an error. Main grib selector.
tstart	Optional grib selectors perturbationNumber, tstart, tend, dataDate and dataTime can also be
tend	issued via command line arguments (-m, -s, -e, -D, -T), which overwrite the ones in the
perturbationNumber	execution JSON file.
dataDate	
dataTime	
level	
applyConversion	The conversion id to apply, as in the parameters.json file for the parameter to select. The combination parameter/conversion must be properly configured in parmaters.json file, otherwise the application exits with an error.
correctionFormula ¹	Formula to use for parameter correction with p, gem, dem variables, representing parameter value, converted geopotential to gem, and DEM map value. E.g.: p+gem*0.0065-dem*0.0065
demMap	The dem map used for correction.

¹ When configuring correction, all xml attributes *gem*, *correctionFormula*, and *demMap* must be present.

gem Formula for geopotential conversion for correction.	
---	--

OutMaps tag

Attribute/Tag	Details
cloneMap	The clone map with area (<u>must have a REAL cell</u> type and missing values for points outside area of interest. A dem map works fine. A typical area boolean map will not).
unitTime	Time unit in hours of output maps. Tipical value is 24 (daily maps).
namePrefix	Prefix for maps. Default is parameter shortName.
fmap	First map number. Default 1.
Interpolation	See relative table.
ext	Extension mode. It's the integer number defining the step numbers to skip when writing maps. Same as old grib2pcraster. Default 1. Refers to User Manual for further information.

Aggregation tag

Attribute/Tag	Details
step	Step of aggregation in hours.
type	Type of aggregation (it was Manipulation in grib2pcraster). It can be <i>average</i> or <i>accumulation</i> .
forceZeroArray	Optional. In case of "accumulation", and only then, if this attribute is set to"y" (or any value different from "false", "False", "FALSE", "no", "NO", "No", "0"), the program will use a zero array as message at step 0 to compute the first map, even if the GRIB file has a step 0 message.

Interpolation tag

pyg2p 2.0

Attribute/Tag	Details
mode	Interpolation mode. Possible values are: "nearest", "invdist", "grib_nearest", "grib_invdist"
latMap	PCRaster map of target latitudes.
lonMap	PCRaster map of target longitudes.
intertableDir	Alternative home folder for interpolation lookup tables, where pyg2p will load/save intertables. Folder must be existing. If not set, pyg2p will use intertables from ~/.pyg2p/intertables/

<u>Usage</u>

To use the application, after the main configuration you need to configure a template JSON file for each type of extraction you need to perform.

Grabbing information from GRIB files.

To configure the application and compile your JSON templates, you might need to know the variable shortName as stored in the input GRIB file you're using or in the geopotential GRIB. Just execute the following GRIB tool command:

grib_get -p shortName /path/to/grib

Other keys you would to know for configuration or debugging purposes are:

- startStep
- endStep (for instantaneous messages, it can be the same of startStep)
- perturbationNumber (the EPS member number)
- stepType (type of field: instantaneous: 'instant', average: 'avg', cumulated: 'cumul')
- longitudeOfFirstGridPointInDegrees
- longitudeOfLastGridPointInDegrees
- latitudeOfFirstGridPointInDegrees
- latiitudeOfLastGridPointInDegrees
- Ni (it can be missing)
- Nj (it states the resolution: it's the number of points along the meridian)
- numberOfValues
- gridType (e.g.: regular ll, reduced gg, rotated ll)

See Mapping variables section in Appendix B – mapping between grib2pcraster and pyg2p for further information.

Input arguments

If you run ./pyg2p.py without arguments, it will show help of all input arguments.

```
usage: pyg2p.py [-h] [-c json file] [-o out dir] [-i input file]
          [-I input file 2nd] [-s tstart] [-e tend] [-m eps member]
          [-T data time] [-D data date] [-f fmap] [-x extension step]
          [-n outfiles prefix] [-l log level] [-N intertable dir] [-B]
          [-X] [-t cmds file] [-g geopotential] [-C path] [-z path] [-P]
Execute the grib to pcraster conversion using parameters from the input xml
configuration. Read user and configuration manuals
optional arguments:
 -h, --help
                  show this help message and exit
 -c json file, --commandsFile json file
               Path to ison command file
 -o out dir, --outDir out dir
               Path where output maps will be created.
 -i input file, --inputFile input file
               Path to input grib.
 -l input file 2nd, --inputFile2 input file 2nd
               Path to 2nd resolution input grib.
 -s tstart, --start tstart
               Grib timestep start. It overwrites the tstart in json
               execution file.
 -e tend, --end tend Grib timestep end. It overwrites the tend in json
               execution file.
 -m eps member, --perturbationNumber eps member
               eps member number
 -T data time, --dataTime data time
               To select messages by dataTime key value
 -D data date, --dataDate data date
               <YYYYMMDD> to select messages by dataDate key value
 -f fmap, --fmap fmap First map number
 -x extension step, --ext extension step
               Extension number step
 -n outfiles_prefix, --namePrefix outfiles_prefix
               Prefix name for maps
 -l log level, --loggerLevel log level
               Console logging level
 -N intertable dir, --intertableDir intertable dir
               interpolation tables dir
 -B, --createIntertable
               create intertable file
```

-X, --interpolationParallel Use parallelization tools to make interpolation faster. If -B option is not passed or intertable already exists it does not have any effect. -t cmds file, --test cmds file Path to a text file containing list of commands, defining a battery of tests. Then it will create diff pcraster maps and log alerts if differences are higher than a threshold (edit configuration in test.json) -g geopotential, --addGeopotential geopotential Add the file to geopotentials, json configuration file, to use for correction. The file will be copied into the right folder (configuration/geopotentials) Note: shortName of geopotential must be "fis" or "z" -C path, --convertConf path Convert old xml configuration to new json format -z path, --convertIntertables path Convert old pyg2p intertables to new version and copy to user folders Copy configuration from source to user folder (except -P, --copyConf intertables)

Example usages:

```
./pyg2p.py -c ./exec1.json -i ./input.grib -o /out/dir -s 12 -e 36

./pyg2p.py -c ./exec2.json -i ./input.grib -o /out/dir -m 10 -l INFO

./pyg2p.py -c ./exec3.json -i ./input.grib -l /input2ndres.grib -o /out/dir -m 10 -l DEBUG

./pyg2p.py -g /path/to/geopotential/grib/file # add geopotential to configuration

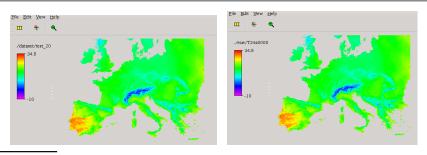
./pyg2p.py -t /path/to/test/commands.txt

./pyg2p.py -h
```

Check output maps

After the execution, you can check how output maps were written using the PCRaster² Aguila viewer. You can also compare maps from grib2pcraster with those from pyg2p you just created:

aguila ./eue/T24a0000.001+10 /dataset/test 2013330702/T24aEFAS.001+10



^{2 &}lt;a href="http://pcraster.geo.uu.nl/">http://pcraster.geo.uu.nl/

Maps will be written in the folder specified by **-o** input argument. If this is missing, you will find maps in the folder where you launched the application (./).

Refer to PCRaster documentation for further information about Aguila.

Interpolation modes

Interpolation is configured in JSON execution templates using the *Interpolation* attribute.

There are four interpolation methods available. Two are using GRIB_API nearest neighbours routines while the other two levereage on Scipy kd tree module.

Note: GRIB_API does not implement nearest neighbours routing for rotated grids. You have to use scipy methods and regular target grids (i.e.: latitudes and longitudes PCRaster maps).

Intertable creation

Interpolation will use precompiled intertables. They will be found in the .pyg2p/intertables folder under user home directory or a custom folder, using the attribute intertableDir within the *Interpolation* attribute or passing -N option on command line.

If interlookup table doesn't exist, the application will create it only if -B option is passed otherwise program will exit.

Be aware that for certain combination of grid and maps, the creation of the interlookup table (which is a numpy array saved in a binary file) could take several hours or even days for GRIB interpolation methods. To have better performances (up to x6 of gain) you can pass -X option to enable parallel processing.

Performances are much better with scipy based interpolation (minutes) but this option could not be viable for all GRIB inputs.

GRIB API interpolation methods

grib_nearest:

This method uses GRIB API to perform nearest neighbour query.

To configure this method:

```
<Interpolation mode="grib_nearest"
latMap="/dataset/maps/lat.map"
lonMap="/dataset/maps/lon.map" />
```

grib_invdist:

It uses GRIB_API to query for four neighbours and relative distances. It applies inverse distance calculation to compute the final value.

To configure this method:

```
<Interpolation mode="grib_invdist"
    latMap="/dataset/maps/lat.map"
    lonMap="/dataset/maps/lon.map" />
```

Scipy interpolation methods

nearest:

It's the same nearest neighbour algorithm of grib_nearest but it uses the *scipy kd_tree*³ module to obtain neighbours and distances.

```
<Interpolation mode="nearest"
latMap="/dataset/maps/lat.map"
lonMap="/dataset/maps/lon.map" />
```

invdist:

It's the inverse distance algorithm with *scipy.kd tree*, using 8 neighbours.

```
<Interpolation mode="invdist" p="1" leafsize="10" eps="0.1"
    latMap="/dataset/maps/lat.map"
    lonMap="/dataset/maps/lon.map" />
```

Attributes *p*, *leafsize* and *eps* for the kd tree algorithm are default in scipy library:

Attribute	Details
р	2 (Euclidean metric)
eps	0

³ http://docs.scipv.org/doc/scipv/reference/generated/scipv.spatial.KDTree.html

CÍZA	110
5120	110

OutMaps configuration

Interpolation is configured under the **outMaps** tag. With additional attributes, you also configure resulting PCRaster maps. Output dir is ./ by default or you can set it via command line using the option -o (--outDir).

Attribute	Details
namePrefix	Prefix name for output map files. Default is the value of shortName key.
unitTime	Unit time in hours for results. This is used during aggregation steps.
fmap	Extension number for the first map. Default 1.
ext	Same as grib2pcraster command line option. Default 1.
cloneMap	Path to a PCRaster clone map, needed by PCRaster libraries to write a new map on disk.

Aggregation

Values from grib files can be aggregated before to write the final PCRaster maps. There are two kind of aggregations available: average and accumulation. The JSON configuration in the execution file will look like:

To better understand what these two types of aggregations do, the DEBUG output of execution is presented later in the following paragraphs.

Average

Temperatures are often extracted as averages on 24 hours or 6 hours. Here's a typical execution configuration and the output of interest:

Configuration cosmo_t24.json:

```
{
    "Execution": {
        "@name": "cosmo_T24",
        "Aggregation": {
            "@step": 24,
            "@type": "average"
```

```
"OutMaps": {
    "@cloneMap": "/dataset/maps/europe/dem.map",
    "@ext": 4,
    "@fmap": 1,
    "@namePrefix": "pT24",
    "@unitTime": 24,
        "@latMap": "/dataset/maps/europe/lat.map",
        "@lonMap": "/dataset/maps/europe/lon.map",
        "@mode": "nearest"
},
"Parameter": {
    "@applyConversion": "k2c",
    "@correctionFormula": "p+gem-dem*0.0065",
    "@demMap": "/dataset/maps/europe/dem.map",
    "@gem": "(z/9.81)*0.0065",
    "@shortName": "2t"
```

Command:

```
.pyg2p.py -l DEBUG -c /execution_templates/cosmo_t24.json -i
/dataset/cosmo/2012111912_pf10_t2.grb -o ./cosmo -m 10
```

ext parameter

In configuration, ext value will affect numbering of output maps.

```
[2013-07-12 00:06:18,545] :./cosmo/T24a0000.001 written!
[2013-07-12 00:06:18,811] :./cosmo/T24a0000.005 written!
[2013-07-12 00:06:19,079] :./cosmo/T24a0000.009 written!
[2013-07-12 00:06:19,349] :./cosmo/T24a0000.013 written!
[2013-07-12 00:06:19,620] :./cosmo/T24a0000.017 written!
```

This is needed because we performed 24 hours average over 6 hourly steps.

Accumulation

For precipitation values, accumulation over 6 or 24 hours is often performed. Here's an example of configuration and execution output in DEBUG mode.

Configuration dwd_r06.json:

```
{
    "Execution": {
        "@name": "dwd_rain_gsp",
        "Aggregation": {
            "@step": 6,
            "@type": "accumulation"
},
    "OutMaps": {
            "@cloneMap": "/dataset/maps/europe/dem.map",
            "@fmap": 1,
            "@namePrefix": "pR06",
            "@unitTime": 24,
            "Interpolation": {
                 "@latMap": "/dataset/maps/europe/lat.map",
                 "@lonMap": "/dataset/maps/europe/lon.map",
                 "@mode": "nearest"
            }
        },
        "Parameter": {
                 "@shortName": "RAIN_GSP",
                 "@tend": 18,
                 "@tstart": 12
        }
    }
}
```

Command:

```
.pyg2p.py -l DEBUG -c /execution_templates/dwd_r06.json -i
/dataset/dwd/2012111912_pf10_tp.grb -o ./cosmo -m 10
```

Output:

```
[2013-07-11 23:33:20,299] : Accumulation at resolution: 657 [2013-07-11 23:33:20,300] : out[s:6 e:12 res:657 step-lenght:6] = grib:12 - grib:6 * (24/6)) [2013-07-11 23:33:20,316] : out[s:12 e:18 res:657 step-lenght:6] = grib:18 - grib:12 * (24/6))
```

Note: If you want to perform <u>accumulation</u> from Ts to Te with an aggregation step Ta, and **Ts-Ta=0** (e.g. Ts=6h, Te=48h, Ta=6h), the program will select the first message at step 0 if present in the GRIB file, while you would use a zero values message instead.

To use a zero values array, set the attribute forceZeroArray to "true" or "yes" in the Aggregation configuration element.

For some DWD⁴ and COSMO⁵ accumulated precipitation files, the first zero message is an instant precipitation and the decision at EFAS was to use a zero message, as it happens for UKMO⁶ extractions, where input GRIB files don't have a first zero step message.

```
$ grib get -p units,name,stepRange,shortName,stepType 2012111912 pf10 tp.grb
```

```
kg m**-2 Total Precipitation 0 tp instant kg m**-2 Total Precipitation 0-6 tp accum kg m**-2 Total Precipitation 0-12 tp accum kg m**-2 Total Precipitation 0-18 tp accum ... kg m**-2 Total Precipitation 0-48 tp accum
```

^{4 &}lt;a href="http://www.dwd.de/">http://www.dwd.de/

^{5 &}lt;a href="http://www.cosmo-model.org/">http://www.cosmo-model.org/

^{6 &}lt;a href="http://www.metoffice.gov.uk/">http://www.metoffice.gov.uk/

Correction

Values from grib files can be corrected with respect to their altitude coordinate (Lapse rate formulas). Formulas will use also a geopotential value (to read from a GRIB file, see later in this chapter for configuration).

Correction has to be configured in the Parameter tag, with three mandatory attributes.

- correctionFormula (the formula used for correction, with input variables parameter value (p), gem, and dem value.
- gem (the formula to obtain gem value from geopotential z value)
- demMap (path to the DEM PCRaster map)

Tested configurations are only for temperature and are specified as follows:

Temperature correction:

```
"Parameter": {
    "@applyConversion": "k2c",
    "@correctionFormula": "p+gem-dem*0.0065",
    "@demMap": "/dataset/maps/europe/dem.map",
    "@gem": "(z/9.81)*0.0065",
    "@shortName": "2t"
}
```

Evotranspiration correction:

```
"Parameter": {
        "@applyConversion": "k2c",
        "@correctionFormula": "p/gem*(10**((-0.159)*dem/1000))",
        "@demMap": "/dataset/maps/europe/dem.map",
        "@gem": "(10**((-0.159)*(z/9.81)/1000))",
        "@shortName": "2t"
}
```

How to write formulas

```
z is the geopotential value as read from the grib file gem is the value resulting from the formula specified in gem attribute I.e.: (gem="(10**((-0.159)*(z/9.81)/1000)))" dem is the dem value as read from the PCRaster map
```

Be aware that if your dem map has directions values, those will be replicated in the final map.

Which geopotential file is used?

The application will try to find a geopotential message in input GRIB file. If a geopotential message is not stored, pyg2p will select a geopotential file from ~/.pyg2p/geopotentials according the geodetic attributes of the input GRIB file. If it doesn't find any suitable grib file, application will exit with an error message.

Geodetic attributes compose the key *id* in the JSON configuration: Note the \$ delimiter.

longitudeOfFirstGridPointInDegrees\$longitudeOfLastGridPointInDegrees\$Ni\$Nj\$numberOfValues\$gridType

If you want to add another geopotential file to the configuration, just execute the command:

```
./pyg2p.py -g /path/to/geopotential/grib/file
```

The application will copy the geopotential GRIB file into .pyg2p/geopotentials folder (under user home directory) and will also add the proper JSON configuration to geopotentials.json file.

Developer notes about formulas

Previous correction formulas were derived directly from grib2pcraster C code. The need to set up two formulas – one for the gem value and one for the final corrected value (which uses the previously formulated gem value) – exists because the correction formulas are based on DEM and geopotential values. DEM values are in PCRaster format while geopotential data is a GRIB message, which needs to be interpolated before to put it into the correction formula. That means that the correction process is applied in two steps with a split formula as configured.

Grib2pcraster correction process:

In this paragraph, some C code is shown from the original grib2pcraster C application.

Geopotential value is read from a grib file as (note the correction):

```
for (i = 0; i < values\_len; i++) \{ \\ pgeopotential[i] = pgeopotential[i] / 9.81; \\ pgeopotential[i] = pow(10, ((-0.159) * (pgeopotential[i]) / 1000)); \\ \} \\ \}
```

Then, the actual correction is made during the interpolation process (some not relevant code was removed and comments added to the real code):

```
//interpolation
float tmpValue = (pGRIBValues[grid[i].indexInGRIBArrNode1]) *
(grid[i].CoefficientNode1) + (pGRIBValues[grid[i].indexInGRIBArrNode2]) *
(grid[i].CoefficientNode2) + (pGRIBValues[grid[i].indexInGRIBArrNode3]) *
(grid[i].CoefficientNode3)
                          + (pGRIBValues[grid[i].indexInGRIBArrNode4]) *
(grid[i].CoefficientNode4);
if (intCorrectionFlag != 0) {
  //interpolation of geopotential as read before
 float tmpGeo = (geopotential[grid[i].indexInGRIBArrNode1]) *
(grid[i].CoefficientNode1) +
                                 (geopotential[grid[i].indexInGRIBArrNode2]) *
(grid[i].CoefficientNode2) + (geopotential[grid[i].indexInGRIBArrNode3]) *
(grid[i].CoefficientNode3)
+ (geopotential[grid[i].indexInGRIBArrNode4]) * (grid[i].CoefficientNode4);
  //correction formula is applied here (the correctionFormula in pyg2p)
  if (intCorrectionFlag == 1)//temperature correction
            tmpValue = (tmpValue + tmpGeo) - (grid[i].demValue * 0.0065);
  if (intCorrectionFlag == 2)//evap correction
               tmpValue = (tmpValue / tmpGeo) * (pow(10, ((-0.159) * tmpValue ) ) *
(grid[i].demValue) / 1000)));
//store value in the output array
pPCRasterValues[grid[i].indexInPCRasterArr] = tmpValue;
```

pyg2p correction process

The process in pyg2p is almost the same, that means:

- read geopotential from grib, and apply the correction formula as configured in XML (gem="(z/9.81)*0.0065")
- interpolate the gem values so obtained to the PCRaster output grid
- read grib values and interpolate them
- apply the final correction formula as configured (correctionFormula="p+gem-dem*0.0065")

The only difference is that in pyg2p formulas are configurable, as explained in this chapter (read Configuration manual as well), while in grib2pcraster you can choose, with a binary flag in parameters.xml file, between two hard coded formulas (one for temperature and one for evaporation).

Conversion

Values from GRIB files can be converted before to write final output maps. Conversions are configured in the *parameters.json* file for each parameter (ie. shortName). The right conversion formula will be selected using the id specified in the *applyConversion* attribute, and the *shortName* attribute of the parameter that is going to be extracted and converted.

In the *Parameter* section of execution template you will configure only the id of the conversion:

The conversion formula is configured in *parameters.json* configuration file, under the relative parameter configuration, identified by *shortName*. Here the item relative to alhfl_s parameter:

```
{
    "@description": "latent heat flux",
    "@shortName": "alhfl_s",
    "@unit": "W/(m^2)",
    "Conversion to
    apply:
    {
        "@cutOffNegative": true,
        "@function": "x=x*(-0.0353)",
        "@id": "tommd",
        "@unit": "mm/d"
```

Note that formulas must be written in the form var = f(var).

The configuration file parameters.json is needed only for conversion functionality. Tags *description* and *units* serve for logging purposes.

Logging

Console logger level is INFO by default and can be optionally set by using -l (or -loggerLevel) input argument.

Possible logger level values are ERROR, WARN, INFO, DEBUG, in increasing order of verbosity.

Comparing Test Suite

Since version 1.1, pyg2p comes with a little test utility which helps to compare a new version release with the grib2pcraster C application. This functionality will produce and check PCRaster *diff* maps, and print aguila commands on screen; it is being activated with the command:

./pyg2p.py -t test.xml

The program will warn if the number of produced maps differs for a certain test.

For convenience, the output will present the aguila commands for visualizing grib2pcraster, pyg2p and diff maps.

Sample output:

```
===> Producing pcraster diff maps. Copy and paste aguila commands to compare them.
aguila /dataset/testdiffmaps/cosmoT24/diff.013
/dataset/testdiffmaps/cosmoT24/gT240000.013
/dataset/testdiffmaps/cosmoT24/pT240000.013
[GOOD] values are good!
...
```

Writing test.xml file

In test.xml file, you configure:

- path to **grib2pcraster** executable
- path to a *commands.txt* file containing the list of input arguments for both **grib2pcraster** and **pyg2p**
- path to the **pcrcalc** executable, for producing *diff* maps
- the error tolerance when checking diff maps

An example of *text.xml*:

```
<TestConfiguration atol="1e-02"
commands="/home/dominik/ECMWF_grib2pcraster/pyg2p_workingcopy/configuration/tes
ts/commands.txt">
    <g2p exec="/home/dominik/ECMWF_grib2pcraster/grib2pcraster/grib2pcraster" />
    <!--pcrcalc diff.001 = mapEFAS.001 - mappyg2p.001-->
    <PcRasterDiff exec="/home/dominik/devapps/PCRaster-3.0/bin/pcrcalc"/>
</TestConfiguration></test
```

An example of *commands.txt* file:

```
q1@ -p 228.128 -n qR24 -t 24 --fmap=1 -i EpsN320-2013063000.grb -o ./eueR24 -m 10
p1@ -I ERROR -c eue r24.xml -i /EpsN320-2013063000.grb -o ./eueR24 -m 10
g2@ -p 182.128 -n gE24 -t 24 --fmap=1 -m 10 -i EpsN320-2013063000.grb -o ./eueE24
p2@ -| ERROR -c eue e24.xml -i EpsN320-2013063000.grb -o ./eueE24 -m 10
#glofas commands. grib2pcraster needs two executions for each pyg2p command
q3@ -i 20130325 en0to10.grib -o ./multiresSRO -p 8.128 -m 50 -n gsro --date=20130325
-s 0 -e 240 --unit=mm -t 24 --interlookup=intertable 2011 medium.txt
g3@ -i 20130325 en11to15.grib -o ./multiresSRO -p 8.128 -m 50 -n gsro
--date=20130325 -s 264 --fmap=11 -e 360 --unit=mm -t 24
--interlookup=intertable 2011 low.txt
p3@ -c glofas sro.xml -i 20130325 en0to10.grib -l 20130325 en11to15.grib -o
./multiresSRO -m 50
#only pyg2p tests (these are not reproducible with grib2pcraster 1.07)
p4@ -i UKMO 20121218.grib -o ./UKMO/ -c UKMO t24 LA.xml -m 0
p5@ -i EUE 2013073100 cv.grb -o ./UKMO/ -c EUE RainAnim CV.xml -m 0
#scipy comparison tests
p6@ -i UKMO 20121218.grib -o ./UKMO/ -c UKMO t24 LA.xml -m 0
z6@ -i EUE 2013073100 cv.grb -o ./UKMO/ -c EUE RainAnim CV scipy.xml -m 0
```

See the following paragraph or the Configuration manual for further details.

Writing the commands.txt file

When writing the commands.txt file, please follow these rules:

- Each test is composed by both the list of arguments for **grib2pcraster** and **pyg2p**. Prefix each list of arguments with 'g' (for **grib2pcraster**) and 'p' (for **pyg2p**) (or 'p' for **pyg2p with grib api interpolation** and 'z' for **pyg2p with scipy interpolation** or any alternative execution configuration to compare the two)
- and use the same id to tie them.
 - If in a test case you need to compare two executions of grib2pcraster with one of pyg2p (i.e. for multiresolution grib files), just write the two list of grib2pcraster arguments using the same id.
 - If in a test case you want to execute only the pyg2p command, just write the respective list of argument as pN@.
- Output directories of each test must be the same for the two executions. This limitation happens because of how the **pcrcalc** command works.
- Maps prefixes of each test must be 'g' for **grib2pcraster** output maps (or pyg2p alternative

- <u>output maps e.g. with scipy interpolation) and 'p' for **pyg2p** output maps (to configure in xml).</u>
- Don't use -report=2 option for grib2pcraster commands and use -l ERROR for pyg2p commands to avoid pollution of output console.

Sample Output

```
======== Running Test 5
     - pyg2p comm -l ERROR -c dwd_e06.xml -i dwd_grib1_ispra_LME_2012111900 -o
./dwdE06
     - g2p comms [u'/grib2pcraster/grib2pcraster -p 121.2 -n gE06 -t 06 --fmap=1 -s 12
-e 18 -i dwd_grib1_ispra_LME_2012111900 -o ./dwdE06']
     - out dir ./dwdE06/
Running grib2pcraster...
Running pyg2p...
====> Producing peraster diff maps. Copy and paste aguila commands to compare
them.
aguila ./dwdE06/diff.002 ./dwdE06/gE060000.002 ./dwdE06/pE060000.002
[GOOD] values are good!
aguila ./dwdE06/diff.001./dwdE06/gE060000.001 ./dwdE06/pE060000.001
[GOOD] values are good!
========= SUMMARY ==========
pyg2p test 5 executed in 0:00:01.005321
pyg2p memory usage: max 205.64MB, avg 117.13MB
grib2pcraster test 5 executed in 0:00:00.586653
Difference: 0:00:00.418668
```

pyg2p API

From version 1.3, pyg2p comes with a simple API to import and use from other python programs (e.g. pyEfas).

The pyg2p API is intended to mimic the pyg2p.py script execution from command line so it provides a Command class with methods to set input parameters and a *run_command(cmd)* module level function to execute pyg2p.

Setting execution parameters

First, create a pyg2p command:

```
>>> import pyg2p
>>> command = pyg2p.command()
```

Then, set up the execution parameters using a chain of methods or single calls:

```
>>>command.with_cmdpath('a.xml')
>>>command.with_inputfile('0.grb')
>>>command.with_log_level('ERROR')
>>>command.with_outdir('/dataout/').with_tstart('6').with_tend('24').with_eps('10').with_fmap('1')
>>>command..with_ext('4')
>>> print(str(command))
pyg2p.py -c a.xml -e 240 -f 1 -i 0.grb -l ERROR -m 10 -o /dataout/test -s 6 -x 4
```

You can also create a command object using the input arguments as you would do when execute pyg2p from shell:

```
>>>args_string = '-I ERROR -c /pyg2p_git/execution_templates_devel/eue_t24.xml -i /dataset/test_2013330702/EpsN320-2013063000.grb -o /dataset/testdiffmaps/eueT24 -m 10' >>>command2 = pyg2p.command(args_string)
```

Execute

Use the *run_command* function from pyg2p module. This will delegate the main method, without shell execution.

>>>ret = pyg2p.run_command(command)

The function returns the same value pyg2p returns if executed from shell (0 for correct executions, included those for which messages are not found).

Adding geopotential file to configuration

You can add a geopotential file to configuration from pyg2p API as well, using the *addGeo* function:

>>>pyg2p.addGeo('/dataset/maps/fredrik/T3999.gph.grb')

The result will be the same of ./pyg2p -g /dataset/maps/fredrik/T3999.gph.grb

Appendix A - Execution XML files Examples

This paragraph will explain typical execution xml configurations.

Example 1: Correction with dem and geopotentials

./pyg2p.py -c example1.xml -i /dataset/cosmo/2012111912_pf2_t2.grb -o ./out_1

This configuration, will select the 2t parameter from time step 0 to 12, out of a cosmo t2 file. Values will be corrected using the dem map and a geopotential file as in geopotentials.xml configuration.

You can use another well known correction whose formulas are configured as below:

```
gem="(z/9.81)*0.0065"
correctionFormula="p+gem-dem*0.0065"
```

Maps will be written under ./out_1 folder (the folder will be created if not existing yet). The clone map is set as same as dem.map.

The original values will be converted using the conversion "k2c". This conversion must be configured in the parameters.xml file for the variable which is being extracted (2t). See Parameter XML tag configuration paragraph in Configuration Manual or in the Conversion chapter in this manual.

The interpolation method is <code>grib_invdist</code>. For grib interpolation methods (<code>grib_nearest</code> and <code>grib_invdist</code>), no additional parameters are needed. Latitudes and longitudes values will be used only if the interpolation lookup table (intertable) hasn't be created yet but it's mandatory to set latMap and lonMap because the application uses their metadata raster attributes to select the right intertable.

The table filename to be read and used for interpolation is automatically found by the application, so there is no need to specify it in configuration. However, lat and lon maps are mandatory configuration attributes.

Example 2: Dealing with multiresolution files

```
./pyg2p.py -c example1.xml -i 20130325_en0to10.grib -I 20130325_en11to15.grib -o ./out_2
```

Performs accumulation 24 hours out of sro values of two input grib files having different vertical resolutions. You can also feed pyg2p with a single multiresolution file.

./pyg2p.py -c example1.xml -i 20130325 sro 0to15.grib o ./out 2 -m 0

This execution configuration will extract global overlapping messages sro (perturbation number 0) from two files at different resolution.

Values will be converted using "tomm" conversion and maps (interpolation used here is grib_nearest) will be written under ./out 6 folder.

Example 3

```
./pyg2p.py -c example1.xml -i dwd_grib_ispra_gme_2006_2_168_i384_2012111500.grib2 -o ./out 13
```

Performing this execution will extract 10u messages (from timestep 12 to 15) from a dwd GRIB2 file.

For resampling, dwd gmi maps file are used.

Again, if there is a precompiled interlookup table, this will be used instead of performing an expensive resampling using GRIB API nearest functions.

Example 4: Average 24 hours

```
./pyg2p.py -c execution_file_examples/execution_6.xml -o ./out_6 -i /dataset/dwd/grib/dwd grib1 ispra LME 2012111900
```

Note the intertableDir specifing an alternative folder where to read and store interlookup tables.

Example 5: Accumulation 24 hours

```
./pyg2p.py -i /dataset/eue/EpsN320-2012112000.grb -o ./out_eue -c execution_file_examples/execution_9.xml
```

Appendix B - mapping between grib2pcraster and pyg2p

This appendix aims to help to migrate from grib2pcraster to pyg2p.

Differences in glossary

- The Manipulation as intended in grib2pcraster becomes Aggregation in pyg2c; if you read the term manipulation in the pyg2c documentation, that means a more generic term, including correction and conversion. Anyway, if you read the python code, the python class which performs aggregation is called Manipulator itself.
- What it's called TypeOfField in grib2pcraster becomes stepType in pyg2p (as for the GRIB standard). It defines the aggregated nature of the variable which can be instant, averaged, or cumulated.
- In grib2pcraster, the variable to extract is called parameter and it's referenced by a couple of numbers: **x.y**, where:
 - **x** is the value of the *indicatorOfParameter* key as found in the GRIB message
 - **y** is an internal number identifying the TableVersion as found in the parameters.xml file of grib2pcraster.
- In pyg2p, while the Parameter term was retained, it is identified by the *shortName* key as found in the GRIB message. This is an independent version key and it's only related to the variable itself and not to the *generating centre* of the file, since this is already coded in GRIB API tables. That is, in pyg2p the concepts of generating centre and table parameter versions don't need to exist.

Mapping variables

To "translate" the parameter code **x.y** used in grib2pcraster to the *shortName* form of pyg2c, just follow these simple steps:

- 1. Identify the parameter in grib2pcraster's parameters.xml file using **y** to find the *gribTablesVersionNo* and **x** to find the single item *Param*.
- 2. Take note of abbreviation and field description strings of the Param above.
- 3. Use some basic GRIB tools (and meteorological) skills to find the *shortName* value of the grib:

grib get -p shortName,parameterName /path/to/grib/file.grib2

Known parameters mapping with conversions

Variable	grib2pcraster selectors (indicatorOfParameter.gribT ablesVersionNo)	pyg2p.py shortName	Configured conversions in grib2pcraster
2 meters Dew- point	• 17.2 • 168.128	• 2D	• x-273.15
2 meters Temperature	11.2500011.0167.128	t_2mtd_2m2t	• x-273.15
Total precipitation	• 61.2 • 228.128	• tp	Only cut offx*1000 + cut off
Latent heat flux	• 500086.0 • 121.2	• alhfl_s	 x*(-0.0353) + cut off [default] x*(-0.00147) + cut off
Surface Latent heat flux	• 147.128	• slhf	No conversion
Surface Runoff	• 8.230 • 8.128	• sro	• x*1000 + cut off
Sub-Surface Runoff	• 9.230 • 9.128	• ssro	• x*1000 + cut off
Large scale precipitation	• 142.128	• Isp	• x*1000 + cut off
Convective precipitation	• 143.128	Not tested. It needs to be added to parameters.xml if used.	• x*1000 + cut off
Evaporation	• 182.128	• e	• x*(-1000)
Runoff	• 205.128	Not tested. It needs to be added to parameters.xml if used.	 x*1000 + cut off (not as default, it needs to be set by command line withunit parameter)
Surface precipitation amount, rain, grid scale	• 102.201	• rain_gsp	Only cut off
Surface precipitation amount, rain,	• 113.201	• rain_con	Only cut off

pyg2p 2.0

convective			
Large scale rain rate	• 500134.0	Not tested. It needs to be added to parameters.xml if used.	Only cut off
Snow fall	• 144.128	Not tested. It needs to be added to parameters.xml if used.	• x*1000
Snow depth	• 141.128	Not tested. It needs to be added to parameters.xml if used.	X*1000 (not as default, it needs to be set by command line withunit parameter)

Execution parameters mapping

Argument	grib2pcraster	pyg2p.py
Input file	-i	-i
Variable	-p x.y	Xml ⁷ (shortName)
Output directory	-0	-0
EPS perturbationNumber	-m	-m
Variable level	level	Xml
Aggregation type	Default in parameters.xml. Overwriting withma	Xml (type)
Aggregation step	-t	Xml (step)
Start step	-S	-s or Xml (tstart)
End step	-е	-e or Xml (tend)
Output map unittime	unittime	Xml (unitTime)
Interpolation	-r,-l,-a or –intertable	Xml (see relative section in this manual).
Conversion	parameters.xml	parameters.xml and Xml (applyConversion)
Correction	Two formulas hardcoded and configured by default in parameters.xml or with	Xml (attributes correctionFormula, gem, dem)

In this table, "Xml" means that the corresponding parameter configuration is in the execution xml template where you configure your extraction and pass to the pyg2p with the -c input argument. If not indicated otherwise, the xml tag/attribute name is the same as the grib2pcraster input argument (e.g.: --level=10 becomes the xml attribute level="10").

	temp andevap.	
Map extension naming	fmap,ext	Xml
Map prefix name	-n	Xml
Logging	report. Loggers directory is under program installation folder	Configurable in logger- configuration.xml. Console logger level overwritable with -I (default is INFO). Default loggers dir ./logs overwritable with -d.

Difference in dealing with multiresolution

Multiresolution extraction can be made in one step with pyg2p, using both -i and -I for input files or even a single multiresolution file. The grib2pcraster application needs to perform two executions, instead.

Differences in Correction

There is a sort of bug in grib2pcraster when come to correction, since the dem values it uses are stored in its interlookup tables and, if the DEM has changed for some reasons, the corrected values will be wrong unless the interlookup table is not regenerated using the new DEM map.

However, here the main difference is that the correction formulas are editable directly by the user in pyg2p.

Differences in Aggregation

The type of aggregation and its step window must be specified in pyg2p, using the *Aggregation* XML tag.

In grib2pcraster, instead, the type of aggregation for the specific variable is being extract (i.e. manipulation) is read from *ManipulationFlag* attribute in *parameters.xml* file: "1" for average and "2" for accumulation. The step window is 24 hours by default and it can be overwritten with -t parameter.

Differences in Interpolation

Resampling of grids in grib2pcraster is made using GRIB API. Available interpolation methods are:

- nearest neighbours
- inverse distance

The interpolation table file is declared with the input argument –intertable, or otherwise by configuration, using one of the tables in *interpolation.xml*.

In **pyg2p** there are two additional resampling modes (see the relative Interpolation modes paragraph in this document). Results are very similar to GRIB API based interpolation methods but interlookup table

pyg2p 2.0

is being created in seconds rather than hours (in our tests, interlookup table creation for high resolution or global scale grids take almost two days with GRIB_API methods while it takes one minute with scipy based methods).

They are *nearest* and *invdist* modes, performed using scipy routines to find neighbours and distances.

Interlookup tables are read/written by default using \$PYG2PHOME/configuration/intertables as folder where to load existing intertables or save new ones. An alternative folder can be specified using the intertableDir attribute in the Interpolation XML tag of the xml template configuration.

Appendix C - Glossary

Brief description of terms you find in this manual, with their synonims.

Term	Synonims	Description
Interpolation lookup table	Intertable, interlookup table	It's a numpy binary file containing indexes correspondences and coefficients/distances. Each combination of GRIB grid/PCR aster target coordinates maps/Interpolation mode has its own intertable. Since there are four interpolation modes using intertables, there can be four different intertables for each combination of GRIB grid/PCR aster lat/lon maps.
Aggregation		Average or accumulation
Interpolation	Resampling	Resampling between the original GRIB grid and the target raster lat/lon coordinates.
Execution XML template file	Execution template, commands file, execution file, xml template, xml configuration	The XML file where most of execution parameters are configured. It's being passed by the mandatory CLI argument -c .
Parameter	Variable	The meteorological variable stored in the input grib file.
Command line Input argument	Input argument, CLI argument, option	Input arguments to the application passed via command line. They are -i,-I,-o,-m,-I,-d,-c
Execution parameter	Parameter, input argument	All configured options (via CLI or XML files) that define the execution context.
GRIB grid	Grid	It's the set of geodetic attributes identifying the coordinate system and the specific projected area. The set includes gridType , first and last values of latitude and longitudes, number of points along the meridian.
GRIB message's gridType	Grid type, grid	It's the specific grid type of the GRIB message and can be regular_rr, regular_II, rotated_rr, rotated_II, reduced_II and so on.
GRIB message	Grib	It's a single message extracted from a GRIB file .
GRIB API		C library (and its wrappers) from ECMWF to deal with grib files .
GRIB file	Grib	An input file encoded using GRIB format.

pyg2p 2.0

Term	Synonims	Description
Geopotential GRIB values	Geopotential	Geopotential values encoded in grib files, used for correction.
Gem value	Gem	In correction, the gem value is being obtained using the formula configured with gem XML attribute, where z is the geopotential value. E.g. gem = " $(z/9.81)*0.0065$ "