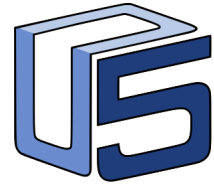


Doc. No. RRT-OSCAR-REP-18006
Revision No 1



OSCAR-5
REACTOR ANALYSIS
PLATFORM

OSCAR-5 Tutorial

Running Core-follow and Related Applications with OSCAR-5

Full-Core Applications Tutorial

OSCAR development team

Last update: 05/04/2019

Radiation and Reactor Theory

Revisions

This document has been revised in accordance with the following schedule:

Rev. No.	Date released	Nature of Revision	Prepared
01	05/04/2019	Original	R H Prinsloo

Abstract

Thus far, you have spent most effort in these tutorials on developing reactor models. We now depart from that somewhat, and focus on some calculational applications for real world reactor problems. As part of your OSCAR-5 distribution you will note that a series of full reactor benchmark problems are included. These serve many purposes, such as support to verification and validation of OSCAR-5, training of users as well as demonstration of the typical usage of more advanced features in the code. These benchmarks also provide you with a good starting point for modelling your own reactor, as many different designs are covered in these test problems. There is a specific user guide covering these problems which you will see later. However, in this tutorial, we will move away from the tutorial space you have been using up to now, and perform the steps described here on the SAFARI-1 validation problem included in the benchmark set. This reactor and experimental description originates from a published IAEA database for research reactor code validation (which is also the case with the other benchmarks distributed with OSCAR-5). In this tutorial you will be briefly introduced to the SAFARI-1 benchmark problem, and then perform a series of typical analysis activities such as core-follow and reload analysis, which can then be compared directly to plant data.

Copyright

Copyright © 2018 [Necsa](#), South Africa

Legal Notice This document is part of the OSCAR-5 calculational system developed by the South African Nuclear Energy Corporation SOC Ltd (Necsa). Neither Necsa, nor any contributor to the code system, nor any person acting on behalf of Necsa makes any warranty, expressed or implied, or assumes any legal liability or responsibility for the accuracy, completeness, usefulness or functioning of any data and related material.

Distribution Notice This document is the property of Necsa. This document is transmitted as part of a license agreement. Any further distribution by any holder is prohibited without the written approval of Necsa.

Contact information

Postal address

Radiation and Reactor Theory
Necsa, Building 1900
P O Box 582
Pretoria
0001
South Africa

Telephone:

+27 (0) 12 305 5042

E-mail:

oscar5@necsa.co.za

Contents

1	Introduction	9
1.1	Background	9
1.2	The Scenario	11
1.3	The OSCAR Approach	11
2	Step-by-Step	13
2.1	Getting Started: The OSCAR-5 Setup for this Scenario	13
2.2	Step 1: Studying the Existing Reactor Model	14
2.2.1	Assembly models	14
2.2.2	The core configuration and facility loading	15
2.2.3	The cOMPoSe model	16
2.3	Step 2: Obtaining a Starting Inventory	16
2.3.1	Residual isotope treatment	16
2.3.2	Importing historic assembly data	18
2.4	Step 3: Facility Loading	20
2.5	Step 4: Performing a Reload Calculation	21
2.5.1	Perform the predictive line calculations	22
2.5.2	Extracting cycle trend parameters	25
2.5.3	Define specific results to calculate	25
2.5.4	Define the reportable quantities	26
2.5.5	Building the report	27
2.6	Step 5: Performing a Core-Follow Calculation	28
2.6.1	Processing plant data	28
2.6.2	Running a core-follow calculation	31
2.7	Step 6: Creating an Independent Core-Follow Line	32
3	Conclusion	34

Symbols and Abbreviations

BOC Beginning-of-Cycle

cOMPoSe OSCAR Model Preparation System

EOC End-of-Cycle

GUI Graphical User Interface

IAEA International Atomic Energy Agency

MCNP Monte Carlo N-Particle Transport Code

OSCAR-5 Overall System for the CAlculation of Reactors, generation 5

SAFARI-1 South African Fundamental Atomic Research Installation, unit 1

1 Introduction

Contents

1.1	Background	9
1.2	The Scenario	11
1.3	The OSCAR Approach	11

Congratulations on completing the first two tutorials that gave an overview of the [OSCAR-5](#) system (*Your First OSCAR-5 Run*) and covered the model building (*Building a Reactor Model with OSCAR-5*) side of the system. This current tutorial will focus on some key applications of the [OSCAR-5](#) system, i.e. core-follow and reload calculations. Additionally, related activities such as importing an inventory, loading a facility, and sharing number densities between codes will also be covered.

1.1 Background

In this tutorial we focus on some calculational applications for real world reactor problems. As part of your [OSCAR-5](#) distribution you will note that a series of full reactor benchmark problems are included. These serve many purposes, such as support to verification and validation of [OSCAR-5](#), training of users as well as demonstration of the typical usage of more advanced features in the code. These benchmarks also provide you with a good starting point for modelling your own reactor, as many different designs are covered in these test problems. There is a specific user guide covering these problems which you will see later. However, in this tutorial, we will move away from the tutorial space you have been using up to now, and perform the steps described here on the SAFARI-1 validation problem included in the benchmark set. This reactor and experimental description originates from a published International Atomic Energy Agency (IAEA) database for research reactor code validation (which is also the case with the other benchmarks distributed with [OSCAR-5](#)).

The objective of any reactor analyst is to simulate the reactor, but this is a process rather than a single step due to the varied nature of reactor operations and the data required at different points of the operating process. The interrelated and cyclic nature of the feedback between operation and calculation is illustrated in [Figure 1.1](#).

After the completion of a reactor cycle, plant data is available for use in a simulation to update the burn-up and number densities of the assemblies used in that cycle, with the process data obtained from the plant computer. Thus, before doing a reload for one cycle, one must perform a core follow calculation of the previous cycle.

It is of the utmost importance to be able to track the number densities of several active nuclides, especially fissile material and decay products. This is the primary purpose of performing

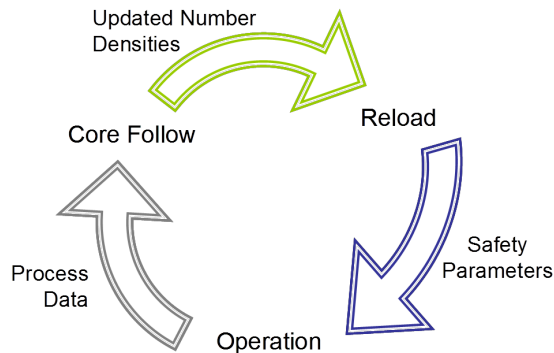


Figure 1.1: A typical reactor calculational cycle.

core follow calculations, and it is important to use the most accurate and up to date process data from the plant for the purpose of inventory tracking.

In this tutorial we assume that you want to start following this process with your new **OSCAR-5** model. However, to get into the process, we first have to find a starting point.

You will learn how to get a starting inventory for your newly built model, load a facility with a set of assemblies from your inventory and then do some reload design calculations for the given core. Assuming that the cycle actually occurred in reality, we move on to perform the core-follow calculation for the cycle after its completion. There will be no detailed discussion on the model building process for the SAFARI-1 benchmark (in general this topic was covered in the previous tutorial on *Building a Reactor Model with OSCAR-5* and *Building a Reactor Model with OSCAR-5 - the nodal model*), but you will be given the opportunity to browse the model as part of this tutorial.

After this tutorial, you will be able to...

Import an inventory from an historical **OSCAR-4** model to your new **OSCAR-5** model, and load a facility for the coming cycle.

Perform a series of reload calculations to confirm that safety and utilization requirements for the proposed core are met

Process plant data, and then set up and run a core-follow calculation with your nodal model for the cycle.

Deploy the core-follow calculation to Serpent to obtain an independent high fidelity core-follow solution

1.2 The Scenario

You are in the process of migrating your reactor simulation tools from OSCAR-4 to OSCAR-5. You have a functioning OSCAR-4 core-follow set and you have just created a full-core nodal model with OSCAR-5. Now you want to create a starting condition for your new OSCAR-5 models, and model your first reactor cycle in OSCAR-5.

You will need a starting fuel inventory for your model (to be taken from your OSCAR-4 model as example in this case) as well as a planned core loading for the coming cycle. Finally, you would need plant data from the reactor once the new cycle has finished its operation to model fuel depletion in the core. After performing the reload analysis and subsequent core-follow calculation for the cycle, confirm the accuracy of the nodal model by running an independent core-follow calculation in Serpent for the same cycle.

1.3 The OSCAR Approach

To perform this work, we will go through four steps, each with its own input file and application in [OSCAR-5](#). We will:

1. Create a script dedicated to import an existing inventory into the [OSCAR-5](#) inventory manager. There are many ways in which this could be done, and the scripting environment makes it easy to customize this for whatever format you have the data defined in (text files, spreadsheets or other code packages). Typically this data contains, for each fuel assembly in your storage pool or core, a description of the axial distribution of burn-up (and potentially radial) and isotopic number-densities. In this case, we will assume that you are migrating from an existing nodal code, namely [OSCAR-4](#). In [OSCAR-4](#), inventory information for each fuel element was stored in a so-called history file, which contains the required information in time-stamped format. After defining the data source, and choosing the time-point for import, you will specify which fuel elements to import. Finally, the script is executed and the [OSCAR-5](#) code-independent inventory is populated. If you have defined a nodal model for your core, specific inventory files for the nodal code will also be created.
2. Next the facility loading has to be performed, which places a subset of assemblies into a chosen configuration – normally for your core, but it could of course serve to load elements into other facilities such as the pool, the vault or other storage racks. For this task, you will define a cycle specific script, complete it and execute the script. You can also create different types of loadings, such as actual (as it happened in the core), or design (as you are planning it) to name but two.
3. Now that the core is loaded, we can perform some analysis on the proposed loading, in order to confirm that the safety and utilization parameters meet their respective requirements. [OSCAR-5](#) houses a flexible interface for defining your own custom set of reload parameters which you would like to check against their respective limits. These parameters can then be calculated with all codes connected to [OSCAR-5](#), and an automated report is produced summarizing what was defined. The customizable aspect is quite important,

given that different reactors have slightly different definitions of the same parameters. Prime examples hereof would be parameters such as shutdown margin, excess reactivity, peaking factors and predicted cycle length. To perform the reload calculation, we again develop a script, and execute it to perform the reload analysis.

4. It might be of interest to be aware of a second approach to producing a reload report. As an alternative, we could consider using some of the code specific features included in some of the [OSCAR-5](#) sub-codes. For example, the nodal solver in [OSCAR-5](#) contains an automation assistant called [OASYS](#), which has the capability to produce a reload analysis and report specific to the nodal solver. [OSCAR-5](#) also supports this [OASYS](#) sub-code, and can auto-generate the necessary input to run any of the various options available in [OASYS](#). We will not spend much time on this approach further in this tutorial, but this feature is valuable to know for users who used [OASYS](#) as part of the [OSCAR-4](#) code system, and would like to continue doing so. To perform reload analysis in this way, we would deploy the [OSCAR-5](#) model to [OASYS](#) and then continue to work in that environment.
5. Next we march ahead in time and reach the end-of-cycle. At this point plant data for the past cycle is available from the reactor, and we can proceed to perform a core-follow calculation. Here we have to think of two distinct steps. The first is to analyse the plant data (power levels, control rod positions, temperatures and energy delivered as a function of time). It is typical to have a look at the granularity of the data (in this case it will be quite fine – order of minutes) and then to process the data into a reasonable quantization, to allow for an accurate simulation of the cycle, while still retaining acceptable calculational efficiency. This typically means to divide the data into two kinds of steps – [flux](#) steps to calculate the reactivity and neutron flux distribution in the reactor at a given time point, and [burn-up](#) steps (spanning typically hours to days of real time), which performs the depletion of reactor materials with the last available fluxes. Your task will be to use the tools available in [OSCAR-5](#) to process the plant data, and the second task will be to create and run a script to simulate the cycle. As before, you can deploy this core-follow application to either the nodal solver [MGRAC](#) or the Monte Carlo solver [Serpent](#).

2 Step-by-Step

Contents

2.1	Getting Started: The OSCAR-5 Setup for this Scenario	13
2.2	Step 1: Studying the Existing Reactor Model	14
2.2.1	Assembly models	14
2.2.2	The core configuration and facility loading	15
2.2.3	The cOMPoSe model	16
2.3	Step 2: Obtaining a Starting Inventory	16
2.3.1	Residual isotope treatment	16
2.3.2	Importing historic assembly data	18
2.4	Step 3: Facility Loading	20
2.5	Step 4: Performing a Reload Calculation	21
2.5.1	Perform the predictive line calculations	22
2.5.2	Extracting cycle trend parameters	25
2.5.3	Define specific results to calculate	25
2.5.4	Define the reportable quantities	26
2.5.5	Building the report	27
2.6	Step 5: Performing a Core-Follow Calculation	28
2.6.1	Processing plant data	28
2.6.2	Running a core-follow calculation	31
2.7	Step 6: Creating an Independent Core-Follow Line	32

In this chapter you will create a working reload and core-follow set for a newly built SAFARI-1 reactor full core model. There are many ways in which you can approach such a task but we will demonstrate our suggested way using the **OSCAR-5** system. You will be referred to appropriate sections in the **OSCAR-5** user guide for information on other methods and tools.

2.1 Getting Started: The OSCAR-5 Setup for this Scenario

The project space for this tutorial can be found at:

`<OSCAR5Path>/validation/`.

We will therefore refer to this path as the `<TutorialPath>` for this tutorial. For this tutorial, we will use the SAFARI-1 benchmark to perform all the necessary calculations. The model

description for the SAFARI-1 model that you will use can be found here:

```
<OSCAR5Path>/validation/SAFARI_1/document_browser.html
```

As usual, **OSCAR-5** should be installed on your system. The actual calculational codes within the **OSCAR-5** system should of course be pre-installed and their paths defined in the codes within the **OSCAR-5** configuration files.

The **OSCAR-5** user guides will also be of great help during this tutorial. These `.html` guides can be found at

```
<OSCAR5Path>/manual.html,
```

where `<OSCAR5Path>` typically refers to `~/OSCAR5/`.

Task: The first task for you in this tutorial, is to browse the auto generated model documentation produced for the SAFARI-1 model (using the html link given above). Have a look at the fuel design, control rod design, core layout and also the nodalisation used to produced the homogenized model.

2.2 Step 1: Studying the Existing Reactor Model

Given that you are familiar with the building of reactor models after completing the tutorial Building a Reactor Model with **OSCAR-5 Part 1** and **Part 2**, we will only have a quick look at the existing model of the SAFARI-1 reactor, which will form the basis of this tutorial. The model is based on the **SAFARI-1 benchmark specification**, and also contains its own full set of **OSCAR-5** generated **model description documentation**.

2.2.1 Assembly models

One notable difference between the assembly models that you have encountered in previous tutorials and those in this model, is that many of these have parts, or even the full assembly model, imported from an existing **MCNP** input. One example of only parts being imported is the fuel assembly, found in

```
<OSCAR5Path>/validation/SAFARI_1/model/leu_fuel_assembly.py .
```

In this case the fuel plate bundle is constructed with the `PlateAssembly` macro, while all the other regions, such as the end adapters, are imported from **MCNP**. We will have a quick look at a simpler case where the whole model is imported from **MCNP**. If you open the file

```
<OSCAR5Path>/validation/SAFARI_1/model/hollow_beryllium.py ,
```

you will see the following import statement:

```
from core import *
from material_library.moderators import LightWater
import materials
from extensions.mcnp_model_importer import *
```

Notice the reference to `extensions.mcnp_model_importer`, which gives access to the built-in system interface with MCNP input. Inside the `build()` method, you will see the familiar material definitions, followed by the following statements:

```
src = create_mcnp_importer(utilities.path_relative_to(__file__, 'sources
/RRT_SAFARI-1_Model_v2.i'))

# pick a universe
u = src.universe(20)
u.materials = {120: lwt, 112: al, 130: be}
# extract all cells from the universe
cells = u.extract_cells()
# add all the cells to the components
hb.add_cells(*cells)
```

You can see that it starts by specifying the path to the MCNP input that should be used, whereafter it indicates the specific universe and materials that correspond to this assembly, and finally adds the cells that were extracted from that universe to this model.

You are welcome to look at the other assembly models and compare their contents to what you now know about modelling of assemblies in OSCAR-5.

Task: Visualize some of the assemblies, especially those that were not used in previous models, such as the shielded ringas or the pool.

2.2.2 The core configuration and facility loading

The basic structure of the core configuration is the same as before, although it now contains more loadable facilities, a number of stationary reflector assemblies, and several control rods that are grouped into different banks. To see what the reactor you will be using to do full core calculations on, visualize the file

`<OSCAR5Path>/validation/SAFARI_1/configurations/base_hot_core.py`.

It is possible to have several different configurations that can form the basis for different types of applications. Examples of this can be found in the set of reactor models included in the OSCAR-5 distribution.

The facility loading is an interesting case, given that it forms part of the bridge between the high level conceptual model of the assemblies and core configuration, and the physical core for which calculations are performed. Take a look at the file

`<OSCAR5Path>/validation/SAFARI_1/facility_loading/tutorial_C1109_1.py`.

It should look quite familiar, except that the core map now contains assemblies that were not

imported with the fresh importer as before. Section 2.3.2 will discuss where the data for them comes from and how to obtain it.

2.2.3 The cOMPoSe model

This is the first model you encounter that was not built with `auto_compose` interface. It is quite an extensive model, and creating it is beyond the scope of this tutorial series. To see the axial cuts, visualize the model with the command:

```
> oscar5 SAFARI_1.compose.operational visualization
```

In this cOMPoSe model, both the fuel and control follower sections are taken from infinite lattice environments.

2.3 Step 2: Obtaining a Starting Inventory

We will be working with SAFARI-1 cycle C1109-1 as our starting point. In order to initiate an inventory at that point, it is assumed that you have an OSCAR-4 inventory. This means that you already have a database of partially burnt assemblies, and thus you are of course not starting with a fresh core.

OSCAR-5 offers many options for obtaining a starting inventory (see the OSCAR-5 user guide by opening `<OSCAR5Path>/manual.html`).

Although we plan to use an OSCAR-4 model as source in this tutorial, the same process is used if you have data defined in some other format. OSCAR-5 uses the concept of an `importer`, which provides the set of rules for reading the historic data files. Although a series of typical importers are predefined for you, you can easily add your own. In this case though, we will use an importer called the `OSCAR4HistoryImporter`.

OSCAR-4 keeps track of material compositions of all loadable components. For each loadable component, e.g. fuel assemblies and follower type control rods, there are associated component history files. The history file contains the irradiation history of the particular component. It lists date and time stamped material data including fuel exposures and nuclide number densities at specific time points. In this specific historic OSCAR-4 model, data was written to the history at two time points for each cycle i.e. at the Beginning-of-cycle (BOC) and at the End-of-Cycle (EOC). These history files then make up the starting inventory for the new model that you intend to use. In this step you will import elements from this already existing inventory so that it becomes the starting material composition data for partially burnt components. Note that we intend to import all assemblies that were already burned at the beginning of cycle C1109-1, as fresh assemblies will be added in facility loading actions (next section). Of course we have no guarantee that the historical model has the same meshing or list of isotopes as our new model, but this should not be a problem as the importer will take care of these differences.

2.3.1 Residual isotope treatment

Before we get into the step-by-step process, we have to introduce additional information on the process of creating an OSCAR-5 element inventory. We want it to be code independent.

However, it is typical for deterministic reactor modelling tools to keep track only of a subset of the most important actinides and fission products. This is the case for the historic [OSCAR-4](#) inventory which we are importing from. This list typically contains between 20 and 50 isotopes, and allows the core solver to perform core depletion with this subset of neutronicallly important isotopes explicitly during the core diffusion calculation. The remainder of the fission products and actinides (called residual isotopes), not selected for explicit core tracking (typically a few hundred of them) are often modelled as a single macroscopic lump material. This material is not depleted explicitly at the core diffusion stage, but simply tabulated during the lattice code calculation as a function of burn-up. To capture their effect during the core calculation, a lookup from the lattice code output for the lump macroscopic cross-section is performed when needed.

We face a challenge then, since our aim of defining a code independent inventory, would require that our database carry explicit number densities for all fission products, actinides and structural material. How do we reconstruct these when importing?

The [OSCAR-5](#) approach to this is the following:

1. Perform, for each loadable, burnable assembly type, a so-called exposure calculation, which depletes the assembly in an infinite-lattice (or your own choice of colourset) environment at nominal conditions. At each burn-up step, all microscopic isotopes present in the assembly are tabulated as a function of burn-up. This database is stored for later use. Just a note, that [OSCAR-5](#) contains two ways to do this – either via a dedicated exposure calculation (which we will show you below) or via a special input option added to the lattice calculation input when performing lattice cut calculations with [Serpent](#).
2. Import the historic assembly data, and instruct [OSCAR-5](#) to fetch information on the residual isotopes from the exposure database by matching the exposure of the imported assembly to that of the exposure database. In that way we replace the macroscopic lump with an explicit list of isotopes. You could even choose a single representative isotope if you prefer, as [OSCAR-5](#) allows you to choose an isotope and calculate its number density in such a way as to retain the reaction rate of the lumped material. B-10 (boron 10) is a typical choice for such a representation.

We are not going to cover the exposure calculation in detail here, as the process is very similar to that which you did in the first tutorial to perform a lattice cut calculation for a fuel element. However, to see what the input looks like, you can open the file:

```
<OSCAR5Path>/validation/SAFARI_1/exposure/leu_fuel_assembly.py
```

To run the exposure calculation, we would use the following command:

```
> oscar5 SAFARI_1.exposure.leu_fuel_assembly --config-file buttercup.cfg \  
execute --threads 20
```

after which you can post process the result to produce the database with

```
> oscar5 SAFARI_1.exposure.leu_fuel_assembly \  
--config-file buttercup.cfg post
```

You do not need to perform this calculation, as the exposure database for both the fuel and control follower components has already been created, and you will just use it.

2.3.2 Importing historic assembly data

As we would like to perform calculations starting from cycle C1109-1, our starting inventory should have data up to the end-of-cycle C1108-1. We start of by inspecting the inventory importing script. This is a complete script and you will not have to change it, but simply understand its contents. Open the file:

```
<OSCAR5Path>
/validation/SAFARI_1/facility_loading/tutorial_initiate_inventory.py
```

to inspect the script. The script has been divided into two sections as shown below. As has been the case in previous tutorials, the top part of the script imports some relevant modules from [OSCAR-5](#) and sets some preliminaries. We note that the facility management modules are imported, as well as the specific history importer we intend to use, along with the core configuration and assembly library which you have seen before in most of the previous scripts. You will note that the path to the inventory is set, as well as the path to the facility loading database. The former defines the space where the inventory files will be stored, and the latter refers to the file which will keep a list of facility loading states.

```
import os
from applications.facility_management import *
from core import *
from extensions.oscar_history_importers import OSCAR4HistoryImporter,
    residual_isotope_options
from ..configurations.base_hot_core import model
from ..model import assemblies

model.inventory_manager.inventory = utilities.path_relative_to(__file__, '
    ../SAFARI_1_inventory_tutorial')
model.facility_manager = utilities.path_relative_to(__file__, '../
    SAFARI_1_facility_tutorial')
parameters.model = model
```

Next we set important run parameters for performing the initialization of the inventory and the facility database.

```
# Time at which inventory should be populated
parameters.time = '2011-10-10 00:37:20'
# =====
# Import fuel elements
source_path = '/home/oscar/projects/oscar5_tutorials/FULL_CORE_APPLICATIONS/
    historical_model'

importer = OSCAR4HistoryImporter()
# specify base file paths
importer.base_path = {
    'CNTRL_': os.path.join(source_path, 'assdef', 'CNTRL_ROD.BASE'),
    'F_LEU-': os.path.join(source_path, 'assdef', 'FUEL_LEU.BASE'),
}
# library file
```

```

importer.library = \
    os.path.join(source_path, 'library/SAFARI_LIBRARY.LNX')
# history
importer.inventory_path = os.path.join(source_path, 'hist')

importer.residual_isotope_treatment = residual_isotope_options.
    replace_with_exposure

importer.load_time = '2011-10-10 00:00:00'      # Slightly earlier to allow
    decay from previous cycle

# -----
# Fuel assemblies
fuel = parameters.add_batch(assemblies.SAFARI_1_LEU_fuel, importer=importer)
fuel.add_all(importer.inventory_path, 'EF*L.HIST')

# -----
# Control assemblies
followers = parameters.add_batch(assemblies.SAFARI_1_LEU_fuel_follower,
    importer=importer)
followers.add_all(importer.inventory_path, 'C*L.HIST')

if __name__ == '__main__':
    main()

```

We begin by setting `parameter.time`, which is the date at which a planned facility change should take place. Typically this would be the date and time when the core was physically loaded. However, in this case it will also be interpreted as the time at which the inventory is to be populated.

Next we define the setup for the importer which will search the source (set here via the `source_path` variable) and fetch burn-up data. In this case we define the importer as an `OSCAR4HistoryImporter`. In the `OSCAR-4` world (as is the case for the `OSCAR-5` nodal solver), each assembly (defined by its `HIST` file) has a type, and the basic properties of each assembly type is defined in a so-called `BASE` file. We therefore define the `base_path` and the path to the old nodal cross-section library to tell the importer about the types of assemblies in the historical model (we supply the type name and the file name here). From these historical `BASE` files (and the old cross-section library), the importer knows information such as the axial material mesh of the old assemblies and the list of microscopic isotopes that were historically tracked. You can learn more about the file types associated with the nodal solver in the `OSCAR-4` user-guides and tutorials if you wish, but it is not necessary for this process at this time.

Three more pieces of information is passed to the importer:

- ▷ The `inventory_path` is defined relative to the top-level source path. This is the directory which will be searched for assembly data.
- ▷ We set `residual_isotope_treatment` to `replace_with_exposure`. This makes sure that the inventory is not only populated with the limited list of microscopic isotopes tracked in the historical model (typically between 20 – 50 in nodal codes). The more complete list of fission products and actinides (maybe 300 – 400) may now be populated

in the code independent inventory based on the burn-up value of the given layer in the assembly. The additional isotopes are interpolated (via burn-up) based on a predefined database of isotopes tabulated from a single assembly infinite lattice burn-up calculation for each assembly type. This database is created via a so-called exposure calculation which **OSCAR-5** performs on an assembly type. As discussed above, we accept that this database already exists and you will use it.

- ▷ We set the `importer.load_time`. This is the time at which the historical data source is queried for assembly isotopic and burn-up data. This time is typically set to some time before the `parameter.time` (facility loading time), but after the EOC of the previous cycle when data was saved to the historical model. In practice then, the old inventory is searched for a time stamp that is earlier than or equal to the `load_time`. If such a step is found, the assembly is added to the inventory at its historically saved time-step (and not at the `load_time`). Note that imported data is always added to the inventory at a time-stamp found in the historical database, as this is the time stamp for which the data is valid.
- ▷ Finally, we create assemblies via defining batches of certain types. These new types now have to match some of the heterogeneous assembly types we defined in the **OSCAR-5 model** directory and which thereafter reside in the assembly library. We go further and assign a filter to each batch to assist the importer in selecting which elements have to be imported in that batch. In this tutorial we import a fuel and a control batch.

To execute the script, import inventory, and update the facility loading database, we run:

```
> oscar5 SAFARI_1.facility_loading.tutorial_initiate_inventory \  
execute --force
```

Task: Perform the initialization of your inventory and facility loading database. You can thereafter browse the inventory folder and see what files have been added to it.

We might be done at this point, but we anticipate that we would like to later have a second independent core-follow calculational line with Serpent. To that end, we would need a second inventory for future use, as each core tracking code should store its own depletion information.

Task: Repeat the above task, but create a second inventory called `SAFARI_1_inventory_tutorial_serpent` in anticipation of later having a second independent core-follow line with Serpent. You will have to change the name of the target inventory in the script.

2.4 Step 3: Facility Loading

Now that we have a starting point in time, we can perform the actual facility loading for cycle C1109-1. This script is relatively simply, and serves to store the loading of the core facility at the load time of the cycle. It performs two functions:

- ▷ Add assemblies to the inventory which are fresh in this cycle, by the same batch mechanism as before.
- ▷ Create a loading of type `actual` and define the core load map for this cycle. The `actual` keyword implies that this is an action in reality, as opposed to generating a loading for, let's say, some conceptual core. The loading will be stored at time `parameter.time`, which will be the time stamp used to retrieve this loading in the future.

The facility loading input is primarily just an administrative support tool to make sure that the states of the facilities are updated. Here again we provide you with a completed input. Open the file:

```
<OSCAR5Path>/validation/SAFARI_1/facility_loading/tutorial_C1109.py.
```

The content of the file should now be rather self-explanatory. Browse the file and make sure the content is clear to you. To execute the script, import inventory, and update the facility loading database, we run

```
> oscar5 SAFARI_1.facility_loading.tutorial_C1109_1 execute --force
```

Task: Create a core loading and save the loading in the facility loading database. Note that the file `SAFARI_1_facility_tutorial.py` is created and contains the loading for this core.

2.5 Step 4: Performing a Reload Calculation

It is now time to perform an analysis of the proposed core. The required parameters to define a core as acceptable is highly variable. It differs between reactor types, but even between similar reactor designs simply because of the way that the safety analysis for that reactor was defined. It is therefore important for a modelling support system to allow a high level of customization in defining the set of reload parameters and how to calculate them. In *OSCAR-5*, we go about this in four stages outlined below. Remember that in *OSCAR-5* a result may be calculated with any code connected to it – hence any of the calculated reload parameters discussed here may be equally calculated by the nodal solver or a Monte Carlo code – however, some choices would be more practical than others. For the sake of this part of the tutorial, we are going to perform the calculations with the nodal solver. To define a reload calculation, and generate a report on the results:

- ▷ Define the set of `predictive_lines` you require. A predictive line has a given core loading determined by placement of fuel, control, irradiation rigs and samples, and a cycle progression spanning BOC to EOC. You may have a number of these. For example, you may have a line from which you want to extract power peaking related quantities, and as such want to define an allowable core state which maximizes peaking. In the case of SAFARI-1, this could imply a hot core state, where all power producing rigs are unloaded,

as to maximize the power density in fuel elements. From such a calculation you could aim to extract quantities such as maximum relative peaking factor, maximum local heat-flux or maximum power density. Similarly, you could define a reactivity limiting core from which you could aim to extract quantities such as shutdown margin, control rod worth or excess reactivity. You could also add a line from where you would like to extract utilization information such as typical fuel burn-up or isotope production yields, and load the most typical isotopic load into the target positions.

- ▷ Define a set of results or trending parameters to `extract` from the predictive lines. These give an overview of how the cycle would progress and typically would include time-dependent plots of bank position or mass change over the cycle. This could also include maps of powers, fluxes or masses at various points throughout the cycle.
- ▷ Next we define specific results to `calculate`. Here we think of power peaking, bank worth or other derived quantities. `OSCAR-5` has a series of these known to it, and you can select from an available list, or even add your own. `OSCAR-5` will assess the list of results requested, and set up a set of calculations, called *cases*, to generate the results.
- ▷ Finally, you choose which of the above you would like to include in the final `report`, and provide some additional information such as titles, limits or specific formatting. The requested parameters are then collated into both a `pdf` and `html` report.

In the coming sub-sections, a series of code-snippets are extracted from the relevant input files. Note that in some cases, the order of code extracts do not follow exactly the order of the input files, and might be grouped differently to facilitate the explanation.

2.5.1 Perform the predictive line calculations

To get you started, we have prepared an input file for you that already contains sufficient definition to produce a simplified reload report. Actually, we have prepared two input files. The reason for this is that a large part of the input required to generate a reload report, is common to all cycles for which you would want to do this. Hence it makes sense to separate the cycle-independent part of the input into a reusable definition of what you required in a given reload report, and have a second input file, specific to each particular cycle. In this case you will find the following two relevant files:

1. `reload/generic_report.py` - this is the common input to all cycles for a given definition of what should be included in a reload report.
2. `reload/tutorial_C1109_1.py` - this is the cycle specific input used to perform the calculations and generate the report for cycle C1109-1.

Lets start by having a look at the `reload/generic_report.py` file. As before, we are going to highlight some important parts of the input file, and select relevant parts to reproduce in the blocks below as needed. We look at the part of the file defining the predictive lines:

```

from ..configurations.base_hot_core import model, assemblies

parameters.model = model

peaking =
parameters.add_predictive_calculation('peaking',
                                     'Predictive calculation for the
                                     peaking limiting core',
                                     steps=[0.1 * units.days,
                                             0.25 * units.days,
                                             0.75 * units.days,
                                             1.5 * units.days,
                                             3.0 * units.days])

peaking.bank_search = ['control', 'regulating']
peaking.bank_search.target_keff = 1.00

```

Here we fetch our standard model for the core as built in the *configuration* stage, and create a prediction line – we call it **peaking**. The standard model, as defined in:

```
../configurations/base_hot_core.py
```

does not have fuelled target plates loaded in irradiation positions, and is thus already the correct core layout for use in a peaking calculation. We add the predictive line called **peaking**, and define its cycle progression. The specification for the day increments is followed as specified, with the last step automatically applied multiple times until the planned cycle length is reached. The fine scale steps at the start of the cycle is added for accurate modelling of Xenon build-up. Planned cycle length however is only specified in the cycle specific input, as cycles may vary in planned length. Finally we choose which banks to include in the bank searches, as the cycle progression will perform critical searches and thus automatically find the critical bank at each step in the cycle. We also set for this purpose the target k_{eff} to define the model specific value for criticality (models generally have an off-set in their critical prediction).

We can also have a look at the definition of a second predictive line called **reactivity**. The relevant parts of the input for the second prediction line are given below:

```

# Rig loading (Dummy rigs) c3 = assemblies.SAFARI_1_Homogenized_Mo99(name='
Mo99C3')
e3 = assemblies.SAFARI_1_Homogenized_Mo99(name='Mo99E3')
g3 = assemblies.SAFARI_1_Homogenized_Mo99(name='Mo99G3')
b6 = assemblies.SAFARI_1_Homogenized_Mo99(name='Mo99B6')
b8 = assemblies.SAFARI_1_Homogenized_Mo99(name='Mo99B8')
d8 = assemblies.SAFARI_1_Homogenized_Mo99(name='Mo99D8')
f8 = assemblies.SAFARI_1_Homogenized_Mo99(name='Mo99F8')
f6 = assemblies.SAFARI_1_Homogenized_IPR(name='IPRRF6')
d6 = assemblies.SAFARI_1_Homogenized_IPR(name='IPRRD6')

rig_map = \
[[0, 1, 2, 3, 4, 5, 6, 7, 8, 9],
['A', '-', '-', '-', '-', '-', '-', '-', '-', '-'],
['B', '-', '-', '-', '-', '-', b6, '-', b8, '-'],
['C', '-', '-', c3, '-', '-', '-', '-', '-', '-']]

```

```
[ 'D', '-', '-', '-', '-', '-', d6, '-', d8, _ ],
[ 'E', '-', '-', e3, '-', '-', '-', '-', '-', _ ],
[ 'F', '-', '-', '-', '-', '-', f6, '-', f8, _ ],
[ 'G', '-', '-', g3, '-', '-', '-', '-', '-', _ ],
[ 'H', '-', '-', '-', '-', '-', '-', '-', '-', _ ] ]

reactivity limiting reactivity =
parameters.add_predictive_calculation('reactivity',
                                     'Predictive calculation for the
                                     reactivity limiting core',
                                     steps=[0.1 * units.days,
                                             0.25 * units.days,
                                             0.75 * units.days,
                                             1.5 * units.days,
                                             3.0 * units.days])

reactivity.model.add_assembly_facility_load_map('Irradiation-position',
                                                rig_map)
```

In this case all fuelled rigs are defined, placed in a rig map, and loaded into rig positions. You have already learned how to do this in a previous tutorial, so please look back at tutorial 2 if this rig loading seems unfamiliar. These fuelled rigs are loaded as we would like to have the model in its most reactive state for the sake of conservatism. For now ignore the rest of this input file, and let's move on to the cycle specific input so that we can run these predictive lines. The cycle specific input can be found in `reload/tutorial_C1109_1.py` and contains only a few simple definitions:

```
from generic_report import parameters, reactivity, peaking

parameters.start_time = '10/10/2011 00:37:20'
parameters.cycle_name = 'C1109-1'
parameters.working_directory = utilities.path_relative_to(__file__, '
    tutorial_C1109_1')
parameters.planned_cycle_length = 30 * units.days
```

This file should be largely self-explanatory by now. The `parameters.start_time` is important, as this is the time at which a facility loading will be fetched from the facility loading database. We also note the import of the `generic_report` module, and the setting of some basic parameters and the cycle length for this cycle.

To now perform the two predictive line calculations, we can either launch them both at the same time via:

```
> oscar5 SAFARI_1.reload.tutorial_C1109-1 prediction --target-mode MGRAC
execute --force
```

or call, for example, only the peaking line via:

```
> oscar5 SAFARI_1.reload.tutorial_C1109-1 prediction peaking \
--target-mode MGRAC execute --force
```

Task: Run both peaking and reactivity predictive lines which simulate the cycle progression for the core with and without targets loaded in irradiation positions.

2.5.2 Extracting cycle trend parameters

As discussed in the introduction to the core reload analysis, we next define which parameters we would like to extract directly from the prediction lines to characterise the cycle progression. In this case we choose (from the generic report file):

```
days = parameters.extract(cycle_progression, reactivity)
banks = parameters.extract(bank_positions(bank='control'), reactivity)
masses = parameters.extract(u235_mass, reactivity)
boc_map = parameters.extract(u235_mass_map(at=BOC), reactivity)
eoc_map = parameters.extract(u235_mass_map(at=EOC), reactivity)
```

You will see here that we identify the result of interest with a keyword, followed by the predictive line from which we would like to extract it. In this case we ask for `cycle_progression`, `bank_positions`, `u235_mass` and `u235_mass_map` at BOC and EOC. For a full list of possible keywords, consult the user manual. In this case, we choose them explicitly from the reactivity calculation, as we want these parameters extracted from the core with targets loaded. To extract these parameters, we have to post-process the prediction calculations. We do this, simultaneously for all prediction lines, via:

```
> oscar5 SAFARI_1.reload.tutorial_C1109-1 prediction \
--target-mode MGRAC post --force
```

The post processing step here is optional, as the overall post-processing can be called at the very end, and subsequent steps which need post-processed results will call the post-processing automatically. You thus do not need to perform any task at this step.

2.5.3 Define specific results to calculate

The next step is to define the set of particular results of interest to calculate. Here we have freedom to use the power of the scripting environment in order to define particular results and manipulate them to report correctly defined safety and utilization parameters. The input file you have contains quite a number of examples - and we show a few in the input file extract below. These parameters require additional calculation, as they often require tallies or additional solution options to be activated at the time-points of interest. From the input file we have:

```
avg = parameters.calculate(average_power_density, peaking, 0)
mx = parameters.calculate(maximum_power_density, peaking, 0)
hottest_channel = parameters.calculate(maximum_power_density_channel,
    peaking, 0)

total = configure_banks(all=['control', 'regulating'])
all_in = parameters.calculate(total.banks_inserted(), reactivity, 0) # keff
    with rods in
all_out = parameters.calculate(total.banks_extracted(), reactivity, 0) #
    keff with rods out
```

This block shows how some specific results, such as k_{eff} , average power density or maximum power density with given bank position can be calculated. Once again, for each requested result, you specify from which line to take it, and at which time-point during the cycle. These results to be calculated are grouped together by the code into a set of calculational scenarios, or cases,

which you can launch in a batch, or of course as usual one at a time. To spawn the calculation for these requested parameters, use the command:

```
> oscar5 SAFARI_1.reload.tutorial_C1109-1 cases --target-mode MGRAC \
execute --force
```

after which you should again post-process the results via:

```
> oscar5 SAFARI_1.reload.tutorial_C1109-1 cases --target-mode MGRAC \
post --force
```

Task: Calculate all requested parameters and results for all cases. You once again do not need to perform the post-processing step here

2.5.4 Define the reportable quantities

Not everything you calculate has to be reported, and the reported value is often a derived quantity as compared to what was calculated. To facilitate this, the next step is to collate all the above into a set of reportable results. The following input file extract shows a series of such definitions of quantities to report.

```
parameters.report(days, 'Days')
parameters.report(banks, 'Bank Positions')
parameters.report(masses, 'Core U235 Mass')
parameters.report(boc_map, 'BOC U235 Mass Distribution')
parameters.report(eoc_map, 'EOC U235 Mass Distribution')

parameters.report(mx / avg, 'Peaking Factor', ranges=(None, 3.5), fmt='{:.2f}
}'.format)
parameters.report(mx, 'Peak Power Density', fmt=format_unit(units.W / units.
cc, '{:.2f} W/cc'.format))
parameters.report(hottest_channel, 'Peak Power Channel')
```

The above extract only shows a few of the results identified for reporting, but you should be able to get a sense of how these are handled. Each reportable result is given in terms of the previous extracted and calculated parameters, is flagged with a name, and some further customization such as setting limits and defining formatting is optional. The system will know when a given quantity is a 1D-plot, a map or simply a single-valued result, and report it appropriately.

The reporting process is highly customizable. In essence the process up to now has defined a series of available quantities for reporting. Once these are defined, you process and calculate the final reportable quantities with a post-processing step. This is done via:

```
> oscar5 SAFARI_1.reload.tutorial_C1109-1 --target-mode MGRAC post --force
```

Task: Post-process the report results in preparation of the report compilation.

2.5.5 Building the report

We need to provide the report builder with a template of how the results should be put together in the report itself. This is done by building a report template, which uses a lightly marked up text based description called *restructured text* (or *rst* files). For the set of reportable results you have defined, we have already built a basic report template for you, which you can browse under `reload/report_template/`. Here you will see a simple top level index file called `index.rst` which refers to two chapters – the first called `summary` and the second called `cycle_description` (these reside in the associated *rst* files also). You can have a look at both the `summary.rst` and `cycle_description.rst` files in order to see how the reportable result names you defined in the previous step are placed in the report. Of course the use of restructured text format will take a little getting used to, but remember that you generally only set this template up once to match your reporting requirements and use it for all cycles in future. You will note that you input script refers to this report template toward the end of the script with the line:

```
parameters.report_template = utilities.path_relative_to(__file__, '
report_template')
```

You can therefore have multiple templates for multiple types of reports and use them as required. We are now ready to produce our report. This requires two steps

Prepare the report

Generate the final version of your *rst* based report with the actual data as opposed to the named result place-holders currently in the template via:

```
> oscar5 SAFARI_1.reload.tutorial_C1109-1 --target-mode MGRAC prepare-report
```

After this step, you will see that a populated version of the *rst* files has now appeared within your working directory (should be `reload/tutorial_C1109_1/report/source`)

Build the report

To compile and build the actual final report in both pdf and html formats, use the command:

```
> oscar5 SAFARI_1.reload.tutorial_C1109-1 --target-mode MGRAC build-report
```

This step will produce a browsable html report under

`reload/tutorial_C1109_1/report/build/html/index.html`

and a pdf version under

`reload/tutorial_C1109_1/report/build/latex/C1109-1.pdf`.

Task: Prepare and build the final report in both *html* and *pdf* formats.
Browse the report and consider the behaviour of the proposed core.

2.6 Step 5: Performing a Core-Follow Calculation

We assume that time has now passed and the cycle has operated (in reality) and reached EOC. It is now our job to perform the core-follow calculation, which will update the isotopics of the assemblies in the core to their EOC state. In order to perform the depletion calculation, we first process the plant data, and then perform the simulation.

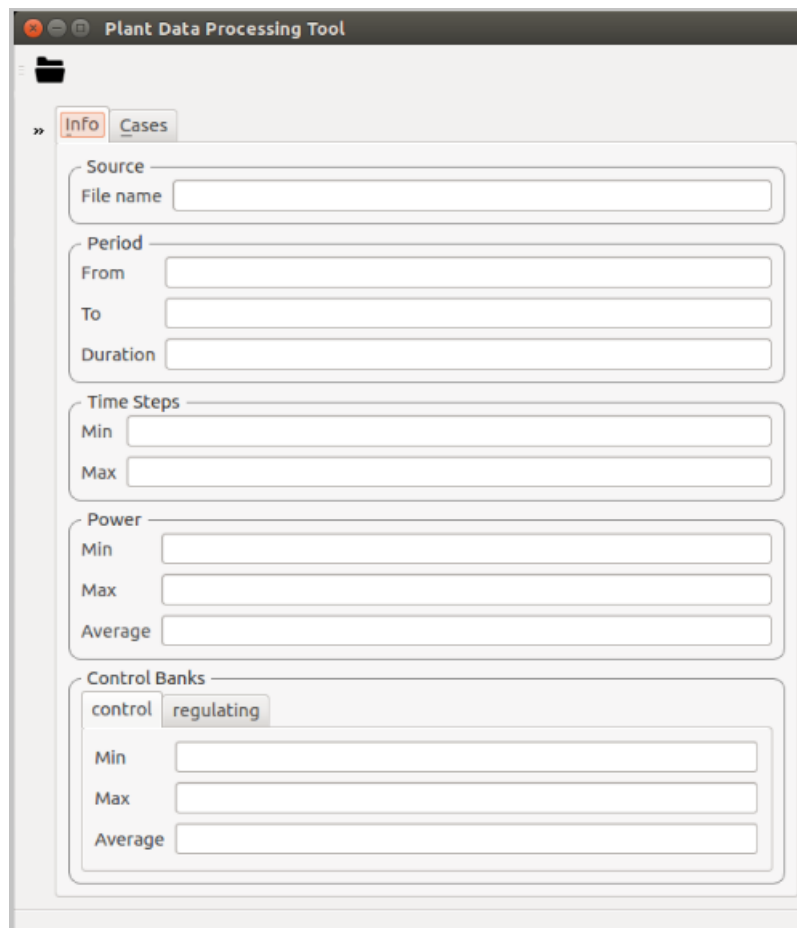
2.6.1 Processing plant data

The first step in a core-follow calculation is to process the reactor operational data or plant data. **OSCAR-5** has a built-in plant data processor. For this part of the tutorial, which is GUI-based, you can follow along with the instructions as we go, and you do not have to wait for a task description to move on. To start the process, execute the following script, which is a generic script placed automatically inside any reactor space you create:

```
> oscar5 SAFARI_1.core_follow.process_plant_data
```

The application window should open as in Figure 2.1.

Click on the file icon in the top left-hand corner, which will enable you to select the plant data for processing. For this tutorial, we have already placed the plant data dump file from the reactor in the benchmark space. To access the aforementioned data, navigate to the `plant_data` folder under your **OSCAR-5 validation/SAFARI-1** folder, and select the `1109-1Adroit.csv` file. Click on the maximize icon to reveal the plant data plot, as seen in Figure 2.2. The plot indicates the reactor power and control rod bank positions as a function of time.



The screenshot shows the 'Plant Data Processing Tool' application window. It features a dark title bar with standard window controls. Below the title bar, there are two tabs: 'Info' (selected) and 'Cases'. The 'Info' tab contains several input sections:

- Source:** A single text input field labeled 'File name'.
- Period:** Three stacked text input fields labeled 'From', 'To', and 'Duration'.
- Time Steps:** Two stacked text input fields labeled 'Min' and 'Max'.
- Power:** Three stacked text input fields labeled 'Min', 'Max', and 'Average'.
- Control Banks:** Two sub-tabs, 'control' (selected) and 'regulating'. Below these are three stacked text input fields labeled 'Min', 'Max', and 'Average'.

Figure 2.1: Plant data processing application window.

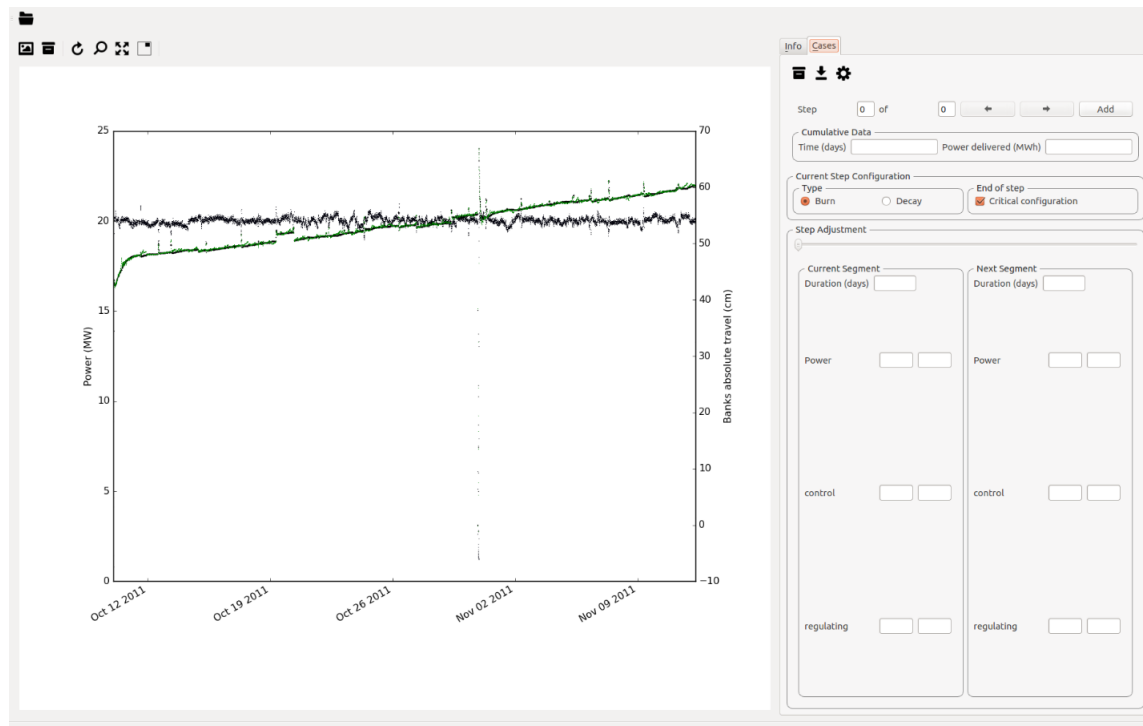


Figure 2.2: Loaded plant data.

The next step is to add critical steps, and burn steps on the graphical representation of the plant data. This is done by selecting the “cases” tab. In the options panel and select add. This will add a burn step on the plant data. The user can now select the characteristics of the step, and define it as a critical burn step, non-critical burn step or a decay step. This choice will depend on the operational characteristics of the reactor, at the specified time step. The duration of the time step can also be varied between very small time steps and a maximum time step (customizable, but here set to 3 days), depending on how rapidly the plant data is changing. It is recommended that at the Beginning of Cycle (BOC), the user specifies a few extra small non-critical time steps to account for rapid changes related to control rod movement and Xenon behaviour. For your convenience we have prepared a plant data file with burn steps for the first couple of days of the cycle. To open the burn step file, select the download icon in the top left corner of the options panel, navigate to the core follow folder under your [OSCAR-5](#) tutorial set, and select the [C1109_1_tutorial.pkl](#) file. If the aforementioned is completed successfully, the burn-up steps will appear on the plant data graph as seen in Figure 2.3. You will see that a few burn steps at the BOC have been added to the plant data, it is now your task to add the rest of the critical burn-up steps.

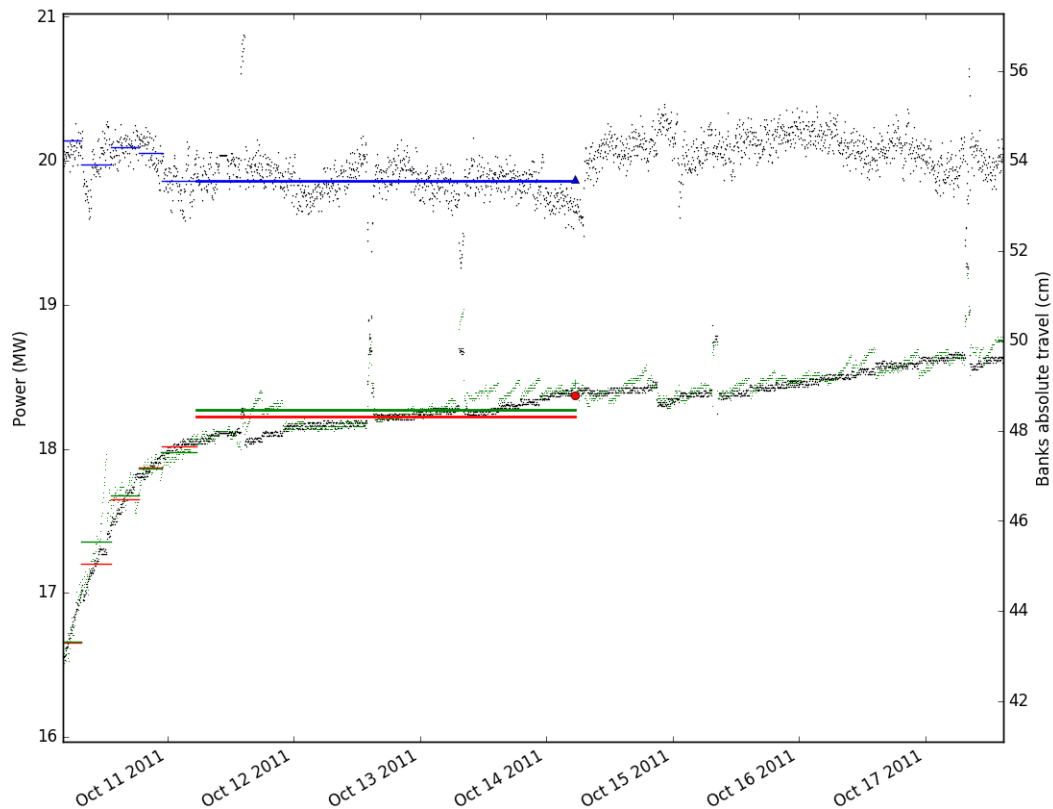


Figure 2.3: Plant data with burn steps at the Beginning of Cycle (BOC) – zoomed view.

A plant data processing file with burn steps for the entire cycle C1109_1 was added to the core follow folder. When proceeding with the core-follow calculations use the `C1109_1_final.pkl` file.

2.6.2 Running a core-follow calculation

As with many of the other scripts in the system, the `manager` utility allows the automatic addition of a core-follow cycle input to the system. You can then populate the script yourself there-after. For example, if you wanted to add a cycle named `TUT_C1109_1` to an existing `OSCAR-5` model, you would:

```
> oscar5 SAFARI_1.manager core-follow TUT_C1109_1 \
--start-time '10/10/2011 00:37:20' \
--configuration base_hot_core
```

Note here that we have pre-specified the configuration and the cycle start time.

Task: Generate a templated core-follow input with the command above. Browse the file to see which data you will have to set to complete the input

You now have to customize this input. Set the following parameters in the input file

1. Set the inventory manager to `'SAFARI_1_inventory_tutorial'`. This points the core-follow calculation to the inventory of fuel assemblies which you created earlier
2. Set the facility manager to `'SAFARI_1_facility_tutorial'`. This database contains the loading of the core at the given start time, as added by you earlier.
3. Set the nodal configuration to `'assemblies.nodal_configuration.Op()'`. This specifies which nodal representation to use when deploying this model to the nodal code, in this case to the standard operational model created for the SAFARI-1 benchmark.
4. Set the plant data to `'../plant_data/C1109_1/1109-1Adroit.csv'`. This refers to the plant data file received from the reactor, which you processed to produce the step wise representation
5. Set the irradiation history to `'C1109_1_final.pkl'`. This file is the result of the plant data processing you performed in the previous step, but in this case does not refer to your produced version, but to a reference file we produced earlier to ensure consistency with the remainder of the tutorial.

Once you have defined these few parameters, it is time to run the core-follow calculation. You can do this via:

```
> oscar5 SAFARI_1.core_follow.TUT_C1109_1 --target-mode MGRAC execute --force
```

Following the calculation, you can inspect the result via

```
> oscar5 SAFARI_1.core_follow.TUT_C1109_1 --target-mode MGRAC post --show
```

Task: Run the core-follow calculation for cycle C1109-1 with the nodal solver, and analyse the behaviour of the critical estimates throughout the cycle. If you wish you can use the pre-completed input named `tutorial_C1109_1.py`. Also look through the working directory structure under `core_follow` and find the output file from the nodal solver `MGRAC`. Open it to see what kind of results are produced by the nodal solver itself.

2.7 Step 6: Creating an Independent Core-Follow Line

Now that you have a working core-follow set for your nodal model, it is good practice to validate this set against a high fidelity code such as [Serpent](#). We will now set up an independent core-follow line in [Serpent](#) and repeat the core-follow calculation. All the work in this regard has

however already been done. You simply have to update the inventory to which the calculation is pointing (as you earlier defined a separate inventory for the high-fidelity line) and point the core-follow script to [Serpent](#). You can do this via:

```
> oscar5 SAFARI_1.core_follow.TUT_C1109_1 --target-mode SERPENT \  
--config-file buttercup.cfg execute --threads 20 --force
```

Once again, since this takes a while, the result of this core-follow calculation has already been post processed and saved for you. In order to compare the results with the nodal solver, you can open the result of the Serpent run via (this now points to the one which we ran for you):

```
> oscar5 SAFARI_1.core_follow.tutorial_C1109_1 --target-mode SERPENT \  
post --show
```

This should open the same graphical display as before, but this time show you the results from Serpent. You should also note a small icon top left of the window, which allows you to compare your result to other cycle data. You can navigate through this option to select the output of the [MGRAC](#) post processing, namely `TUT_C1109_1/MGRAC/C1109_1.res`. You should now be able to compare the runs.

Task: Post process the Serpent core-follow calculation, and compare the results to those you obtained with the nodal solver. Are these differences within expectation? What would be your expectation and how would you check in [OSCAR-5](#) what differences you can expect for this comparison (hint: think of the [cOMPoSe](#) process)

3 Conclusion

Congratulations on surviving your third [OSCAR-5](#) tutorial! You have learned where to start with and how to create a core-follow set for a newly built [OSCAR-5](#) model. You have also been shown how to assess the accuracy of such a core model, by validating it against plant data for control rod calibrations and by doing code-to-code comparisons for both core-follow and rod calibration calculations.

You have now been introduced to the [OSCAR-5](#) system, you have seen how to build reactor models, how to assess the accuracy of said models and how to do key applications with your model such as core-follow and reload calculations. Hopefully you now have the confidence to explore the [OSCAR-5](#) system and apply it to your own reactor designs and calculations.

