

Work Package 1: Harmonization

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SANGOMA Annual Meeting: Liège, 01 – 02 April, 2014

Objectives

- ❑ Analyze the existing tools as a series of modules, some of which are common to several assimilation tools.
- ❑ Propose new modules for WP2 and WP4
- ❑ These modules together with the newly developed modules of WP4 and WP2 then serve as the toolbox for designing new data assimilation systems
- ❑ Adoption of common standards and naming convention

Deliverables

- ❑ **Task 1.1** Identification of common tools.
 - List of commonly available tools (**Month 3**)
- ❑ **Task 1.2** Identification of new tools to be shared
 - List of tools to be adapted (**Month 6**)
- ❑ **Task 1.3** Specification of tool interface data model
 - Specification of data model (**Month 6**)
- ❑ **Task 1.4** Documentation of physical interfaces
 - Augmented list of common and new tools (**Month 12**)
 - Documentation of specifications (Living Document)

Task 1.1 – Task 1.4

□ Summary:

- Analyzed the features of available data assimilation toolboxes and essential modules for SANGOMA.
- A list of common and new modules for SANGOMA was prepared.
- Designed a data model that is consistent with the data models of:
 - Existing systems:
(PDAF, OpenDA, OAK, Beluga / Sequoia, SESAM, TOPAZ)
 - MYOcean:
(HYCOM, NEMO).

Task 1.3 Specification of tool interface data model

- ❑ Overview of the presently available data models:
 - Basic dimensions and Arrays
 - Observation operator.
 - Programming concepts.

- ❑ Design a data model that is consistent with
 - Data models of the existing systems
 - MyOcean data models

SANGOMA Data Model

- ❑ Initially specified as logical data model
- ❑ Programming languages (Fortran, Matlab/Octave)
- ❑ Implementation should also be possible in C, Java
- ❑ Data model was discussed in virtual meeting on May 8, 2012.
- ❑ Two main interfaces were finalized:
 - File Interface
 - In-memory Interface

SANGOMA Data Model

□ File Interface

- I. The NetCDF files will be used to connect models to DA tools
- II. SANGOMA tools should work for CF-complaint input files
- III. Model output standards
- IV. Tools should be compiled with version 3 and 4
- V. Separate NetCDF files for each ensemble member

SANGOMA Data Model

□ In-memory Interface

- I. Basic data structures (no complex data types)
- II. Compatibility of Fortran based codes with C, Matlab, Java
- III. No derived datatypes
- IV. Basic dimensions as in PDAF (statevector, ensemble, observations)
- V. Call-back functions to make use of complex data structures

Example (compute_histogram)

- ❑ Description of functionality/purpose : Compute rank histogram of an ensemble about some state (e.g. ensemble mean or true state). Computation is done for a single location. It increments the information stored in a histogram array.
- ❑ Inputs: Ensemble values for a single grid point. Single state entry. Size of ensemble. Histogram array (size ensemble size+1. It has to be initialized to zero before the first call).
- ❑ Outputs: Histogram array
- ❑ Required work: The module needs to be adapted to be generally usable. The input/output is currently different.
- ❑ Language: F95
- ❑ Needs: no libraries
- ❑ Host: AWI

Task 1.4 Documentation of physical interfaces

- A specification of inputs and outputs of SANGOMA modules were prepared

An updated list of modules for SANGOMA

A detailed specification of these modules (Living document)

Example of Specification: POD

1. Function Name: **Compute_POD**

- **Operation:** This function reads an initial ensemble (*nens* ensemble members) from Netcdf files of model states and store them to an array of vectors (*nstate*, *nens*) of size *nstate*. Then the eigenvalue problem is solved using this array of ensemble. An example of such a Netcdf file is already given in data model description. Before solving the eigenvalue problem the individual vectors may be normalized. The outputs of the function are truncated eigenvalues and eigenvectors.
- **Inputs:** Netcdf files containing model states (see section 2.2)
- **Output:** A data file containing truncated eigenvectors and eigenvalues. The elements of output file are:

P(nstate,nvals)

truncated eigenvectors

Evals(nvals)

vector containing eigenvalues

Example of Specification: POD

```
program compute_POD
  read ensemble(nens) // Assumes ensemble already available
  do i = 1 to nens
    read ensemble(i)
    set X(ntate, i) = ensemble(i)
  end do
  Normalize X
  Compute X' X
  Compute EVD(X' X)
  Return P(nstate, nvals)
  Return Evals(nvals)
end compute_POD
```

Summary

- Listed the modules to be shared for SANGOMA
- Logical data model design
- A detailed specification of shared modules is formulated
- A living document (Deliverable 1.5) is available at svn for inclusion of new methods as we agreed last time.