

# SANGOMA: Stochastic Assimilation for the Next Generation Ocean Model Applications

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A. Barth      J.-M. Beckers      Y. Yan      M. Canter  
University of Liège, BELGIUM

P.-J. Van Leeuwen      S. Vetra-Carvalho  
University of Reading, UK

L. Nerger      P. Kirchgessner  
Alfred-Wegener-Institut, GERMANY

A. Heemink      N. van Velzen      M. Verlaan      U. Altaf  
Delft University of Technology, NETHERLANDS

P. Brasseur      J.-M. Brankart      G. Candille      S. Metref  
CNRS-LEGI, FRANCE

P. de Mey  
CNRS-LEGOS, FRANCE

L. Bertino      F. Counillon  
NERSC, NORWAY



### Executive summary

This deliverable is the third code release of the Sangoma project. Several new tools have been added and adapted to the common data model ([deliverable D1.3](#)). This ensures that the tools can be easily used together and integrated in other assimilation systems. In this release the documentation has also been expanded. Examples have been added on how these tools should be used. The software report of [deliverable D2.5](#) includes the documentation of the tools while this deliverable is the released software bundle with installation instructions.

Intended audience: developers installing the software

## Chapter 1

# Technical highlights

The following gives an overview of the tools included in this software release. More information about these tools are given in [deliverable D2.5](#). Tools in the colour [sangoma blue-green](#) are new tools added in this software release. In total 29 new tools have been developed (often from scratch) and included to this software release which includes now 50 tools in total.

### 1.1 Diagnostic Tools

#### *Fortran*

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<b>sangoma_CheckEnsSpread</b>	Compute ensemble spread and deviation of ensemble mean from an input state
<b>sangoma_CheckNormality</b>	Anderson-Darling Test to check normality of a sample
<b>sangoma_CheckWhiteness</b>	Check whiteness of innovations
<b>sangoma_CompareObsDiag</b>	Compare observation-space diagnostics
<b>sangoma_ComputeBRIER</b>	Compute the Brier skill score and its decomposition, and the entropy
<b>sangoma_ComputeCRIGN</b>	Compute CRPF and CRIGN scores
<b>sangoma_ComputeCRPS</b>	Compute the CRPS and its decomposition
<b>sangoma_ComputeEffSample</b>	Compute the effective sample size of a particle filter
<b>sangoma_ComputeEnsStats</b>	Compute ensemble statistics
<b>sangoma_ComputeHistogram</b>	Compute ensemble rank histograms
<b>sangoma_ComputeInvStats</b>	Compute innovation statistics
<b>sangoma_ComputeMutInf</b>	Compute the mutual information
<b>sangoma_ComputeRCRV</b>	Compute the bias & the dispersion of the RCRV
<b>sangoma_ComputeRE</b>	Calculate the relative entropy
<b>sangoma_ComputeSMatrix</b>	Compute scaled ensemble observation anomalies
<b>sangoma_ComputeSensitivity</b>	Calculate the sensitivity matrix with $\mathbf{H}$ as matrix
<b>sangoma_ComputeSensitivity_op</b>	Calculate the sensitivity matrix with $\mathbf{H}$ as operator
<b>sangoma_arm</b>	Calculate array modes
<b>sangoma_armca</b>	Check the consistency of an ensemble using array modes
<b>sangoma_ObsDiag</b>	Compute sampled observation-space diagnostics

***MATLAB/Octave***

<b>computeBRIER</b>	Compute the Brier skill score and its decomposition, and the entropy
<b>computeCRPS</b>	Compute the CRPS and its decomposition
<b>computeRCRV</b>	Compute the bias & the dispersion of the RCRV
<b>computeHistogram</b>	Compute ensemble rank histograms
<b>mutual_information</b>	Compute mutual information in a particle filter
<b>relative_entropy</b>	Compute relative entropy in a particle filter
<b>sensitivity</b>	Compute sensitivity of posterior mean to observations in a particle filter

**1.2 Perturbation Tools**

***Fortran***

**sangoma\_pseudornd2D**

Generate random fields with given correlation length

**sangoma\_MVNormalize**  
**sangoma\_EOFcovar**

Perform multivariate normalization  
Initialize covariance matrix from EOF decomposition

### ***MATLAB/Octave***

**Weakly constrained ensemble perturbations**

Create ensemble perturbations that have to satisfy an a priori linear constraint

## **1.3 Transformation Tools**

### ***Fortran***

**sangoma\_Anamorphosis**  
**sangoma\_ComputeQuantiles**

Computes local Gaussian anamorphosis  
Computes ensemble quantiles as input for anamorphosis

### ***MATLAB/Octave***

**Empirical Gaussian Anamorphosis**

Determine the empirical transformation function such that a transformed variable follows a Gaussian distribution

## **1.4 Utilities**

### ***Fortran***

**sangoma\_computepod**

Computes dominant POD modes from an ensemble of snapshots

**sangoma\_costgrad**

Computes the values of Objective function and Gradient using reduced state dimensions

**mod\_sangoma\_utils**

Module of utilities for easy porting from MATLAB

### ***MATLAB/Octave***

**hfradar\_extractf**

Observation operator for HF radar surface currents

## **1.5 Analysis**

### ***Fortran***

<b>sangoma_ens_analysis</b>	Computes the analysis ensemble using the ETKF scheme
<b>sangoma_local_ensemble_analysis</b>	Computes the local analysis ensemble using the ETKF scheme
<b>sangoma_enkf_analysis</b>	Compute analysis ensemble using the EnKF with perturbed observations (globally or with covariance localization)
<b>sangoma_ensrf_analysis</b>	Compute analysis ensemble using the EnSRF with serial observation processing (globally or with covariance localization)
<b>sangoma_estkf_analysis</b>	Compute analysis ensemble using the global ESTKF method
<b>sangoma_etkf_analysis</b>	Compute analysis ensemble using the global ETKF method
<b>sangoma_letkf_analysis</b>	Compute analysis ensemble using the ESTKF method with observation localization
<b>sangoma_letkf_analysis</b>	Compute analysis ensemble using the ETKF method with observation localization
<b>sangoma_netf_analysis</b>	Compute analysis ensemble using the NETF method

***MATLAB/Octave***

<b>sangoma_ensemble_analysis</b>	Computes the analysis ensemble using the EnSRF, EAKF, ETKF, ETKF2, SEIK, ESTKF or EnKF scheme
<b>sangoma_local_ensemble_analysis</b>	Computes the local analysis ensemble using the EnSRF, EAKF, ETKF, ETKF2, SEIK, ESTKF or EnKF scheme (domain localization)
<b>sangoma_local_EnKF</b>	Computes the local analysis ensemble using the EnKF (covariance localization)

**1.6 Requirements**

**1.6.1 For Fortran tools**

- **GNU make**
- A **Fortran compiler** (such as gfortran, ifort,...). Under Linux, make and gfortran can be installed by the package manager. For Windows and Mac OS, pre-compiled binaries of gfortran are available at <http://gcc.gnu.org/wiki/GFortran>.
- **LAPACK** and **BLAS**. Reference versions of those libraries are available at <http://www.netlib.org/lapack/> and <http://www.netlib.org/blas/>. An optimized version of BLAS such as OpenBLAS (<http://www.openblas>).

`net/` or Intel's Math Kernel Library (<https://software.intel.com/en-us/intel-mkl>) should be preferred for better performance.

- **FFTW** for computing the Fast Fourier Transform available at <http://www.fftw.org>.
- **pkg-config** is used to detect the installation path of FFTW. This package is generally installed by default on a Linux system.

All requirements can be installed on Ubuntu (14.04) by the following command:

```
sudo apt-get install make gfortran libfftw3-dev \  
libblas-dev liblapack-dev pkg-config
```

For Red Hat/Fedora/Centos, these packages can be installed by:

```
yum install make gcc-gfortran fftw-devel blas-devel lapack-devel pkgconfig
```

## 1.6.2 For GNU Octave/MATLAB tools

You need to have GNU Octave or MATLAB installed on your machine. GNU Octave is an open-source clone of MATLAB available at <http://www.gnu.org/software/octave/>.

Under Ubuntu (14.04) you can install octave by the following command:

```
sudo apt-get install octave
```

For Red Hat/Fedora/Centos, octave can be installed by (as root):

```
yum install octave
```

## 1.7 Installation instructions

Download the file <http://sourceforge.net/projects/sangoma/files/sangoma-2.0.tar.gz> from the Sourceforge site and decompress it. The decompressed files will be in a folder named `sangoma-2.0`

```
wget http://sourceforge.net/projects/sangoma/files/sangoma-2.0.tar.gz  
tar -xvzf sangoma-2.0.tar.gz
```

### 1.7.1 Fortran tools

The Fortran tools can be compiled by issuing `make` in sub-directory `Fortran/` of the `sangoma-2.0` directory:



```
$ make
gfortran -O3 -fdefault-real-8 -Wall -I/usr/include -c sangoma_base.F90 -o sangoma_base
...
ar: creating libsangoma_tools.a
ranlib libsangoma_tools.a
```

This creates a library called `libsangoma_tools.a` and some module files. The sangoma tools can be used from a Fortran or C program. A Fortran program must be compiled by using the option `-I` (indicating the path to the module files) and the option `-L` (the path of the library) and the name of the library `-lsangoma_tools`. For example:

```
gfortran -I/path myprogram.f90 -L/path -lsangoma_tools
```

For a C program, only the options `-L/path` and `-lsangoma_tools` must be specified.

If a different Fortran compiler is used instead of `gfortran`, then the variables `FC` and `FFLAGS` in the Makefile need to be adapted. Note that the default floating point type `REAL` is assumed to be in double precision. A compiler option like `-fdefault-real-8` (`gfortran`) or `-r8` (`ifort`) is thus required.

### 1.7.2 MATLAB/GNU Octave tools

The MATLAB/GNU Octave tools in the sub-directories of the directory `MATLAB/` can be used by including the corresponding directory in your MATLAB/GNU Octave search path by using the script `sangoma_addpath` for including all tools. Individual tools can be added by using:

```
addpath('/path/to/subfolder');
```

Examples for the tools are provided in the sub-directory `examples/` that is include in each category-directory. In case of the Fortran codes, there is a Makefile in the directory that compiles all examples.

## 1.8 Documentation

Documentation of the tools is available at <http://www.data-assimilation.net/Documents/sangomaDL2.5.pdf>.

## 1.9 Access to the development version

The code is available in a [subversion](#) repository. If you are unfamiliar with subversion you can consult the book [Version Control with Subversion](#), the [sourceforge subversion documentation](#) or one of the many tutorials found online (for example [Version Tracking With Subversion For Beginners](#)).

### 1.9.1 Read-only access

You can get the latest version of the sangoma tools by using the following command:

```
svn checkout svn://svn.code.sf.net/p/sangoma/code/tools/trunk sangoma-tools
```

There is no need to be registered for read-only access of the repository. There is also a [web-interface](#) available.

### 1.9.2 Read and write access

Please contact Alexander Barth (a.barth at ulg.ac.be) or Jean-Marie Beckers (jm.beckers at ulg.ac.be) with your sourceforge username (you can [register here](#)) and with a small description of the changes that you plan to make. After you have been added as a developer, you can get the development's version by using the following command:

```
svn checkout --username=your_sourceforge_username \  
  svn+ssh://your_sourceforge_username@svn.code.sf.net/p/sangoma/code/tools/trunk\  
  sangoma-tools
```

You need to replace `your_sourceforge_username` in the previous command.

### 1.9.3 License

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## Deviation from of the Description of Work

This deliverable conforms to the description of work.