

EGU General Assembly 2019

**Short Course SC1.1**

**Data assimilation in the geosciences –**

**Practical data assimilation**

**with the Parallel Data Assimilation Framework**

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*PDAF*

Parallel  
Data  
Assimilation  
Framework

# The Short Course – Overview

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1. Introduction to ensemble data assimilation
2. Implementation concept of PDAF  
(Parallel Data Assimilation Framework)
3. Hands-on Example:  
Build an Assimilation System with PDAF

# 1

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## Introduction to Ensemble Data Assimilation

# Overview

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- What can we expect to achieve with data assimilation?
- What do we need for data assimilation?
- How does ensemble data assimilation work?
- How can we apply ensemble data assimilation?

*Please note:*

We omit equations of assimilation methods because you can apply PDAF without knowing them

(See Short Course SC1.2 on Friday for methodology)

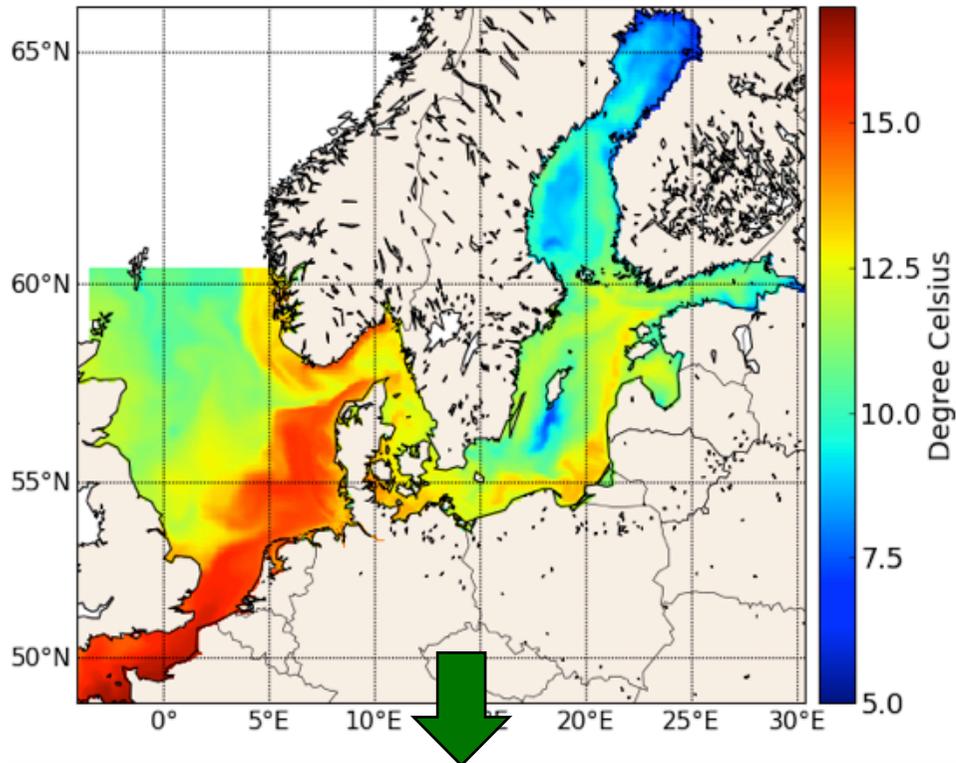
# Application examples

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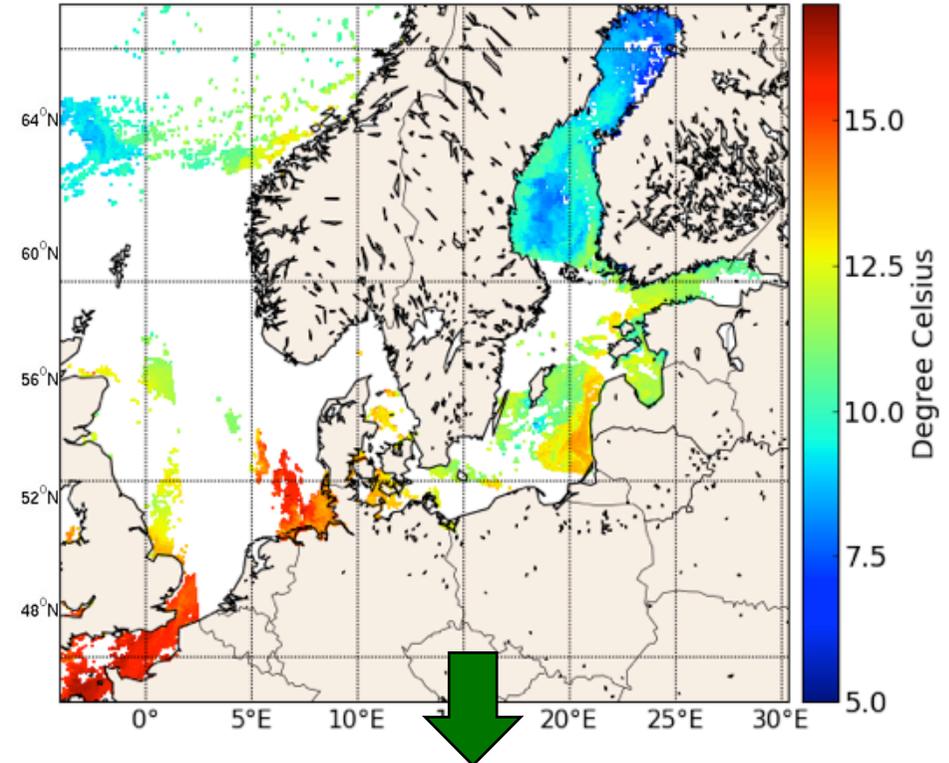
(ocean physics and ocean-biogeochemistry)

# Motivation

*Model* surface temperature



*Satellite* surface temperature

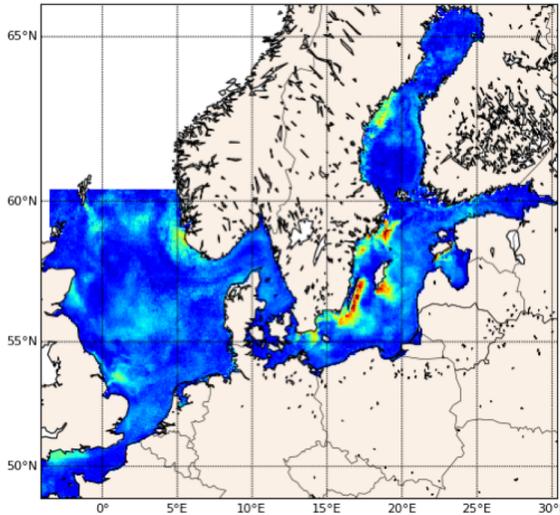


Combine both sources of information  
quantitatively by computer algorithm  
→ Data Assimilation

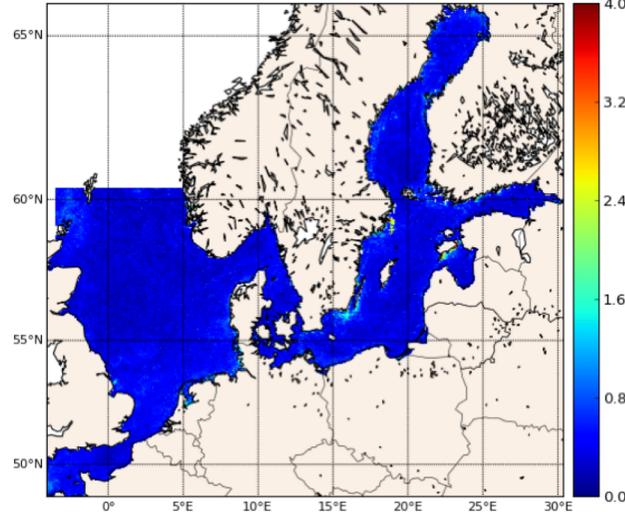
# DA – effect on Temperature (September 2012)

## RMS (root-mean-square) deviation

Free run



Assimilation (analysis)



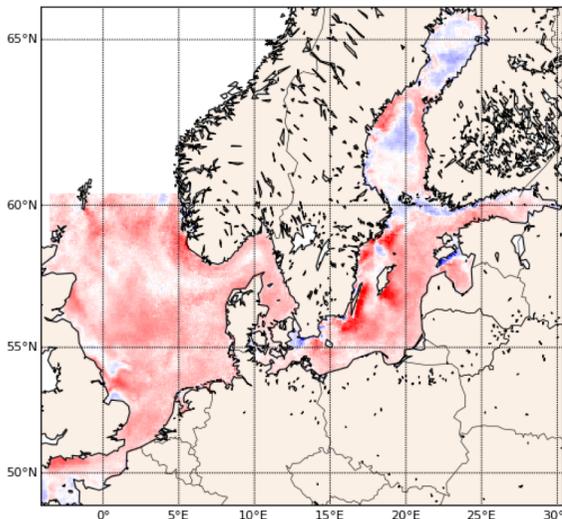
Assimilate surface temperature each 12 h

Compare assimilated estimate with assimilated surface temperature data (monthly average)

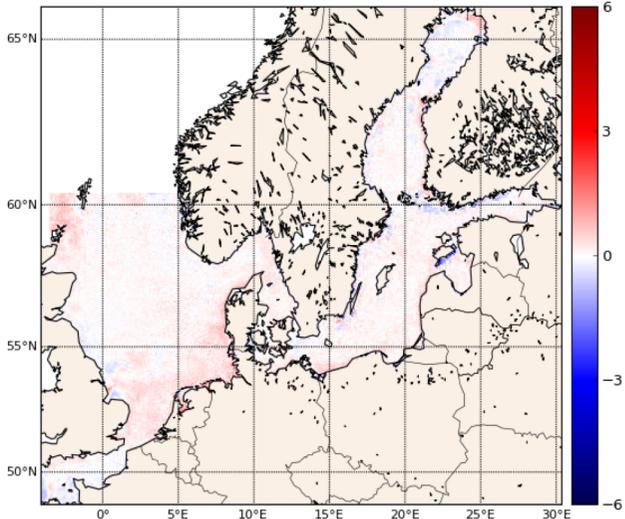
Reduce RMS deviation and mean deviation (bias)

## Mean deviation (observation – model)

Free run



Assimilation (analysis)



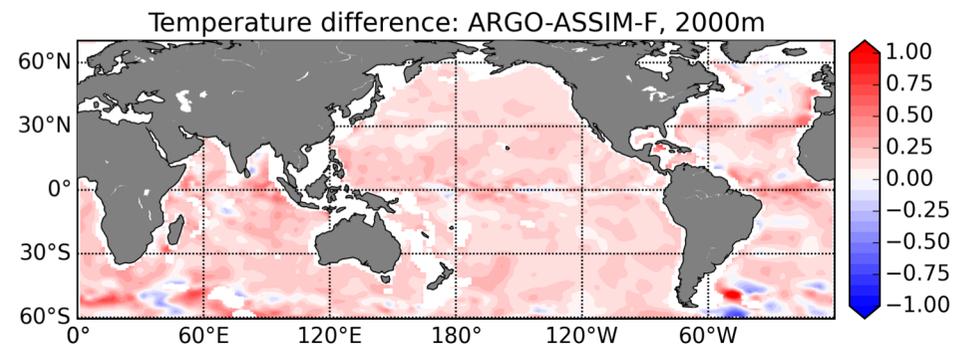
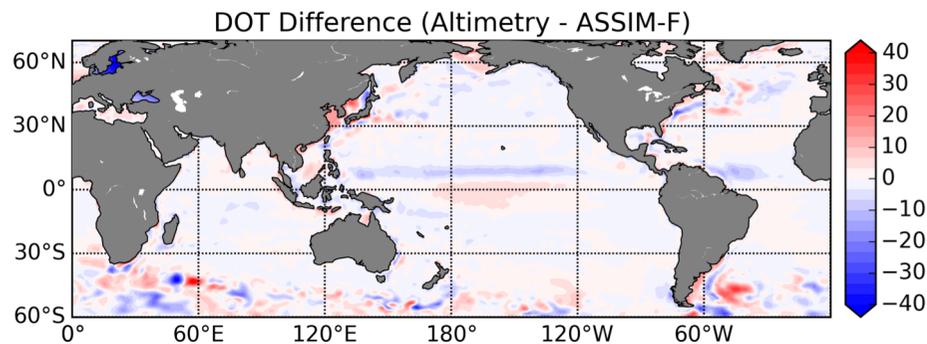
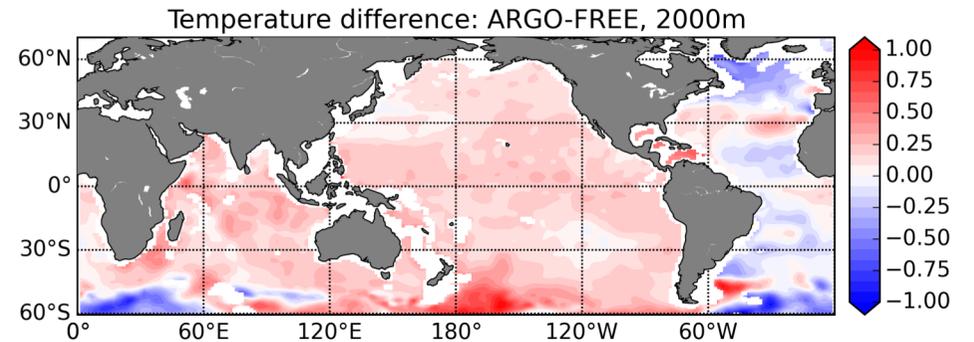
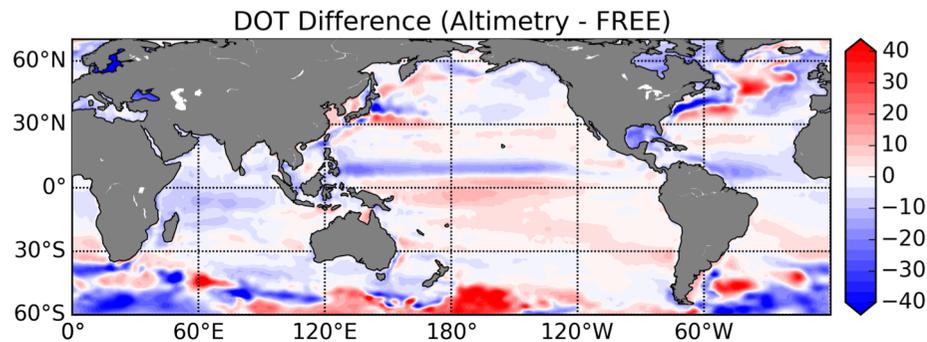
→ necessary effect

# Longe-range effect

**Example:** Assimilate satellite sea surface height data (DOT)

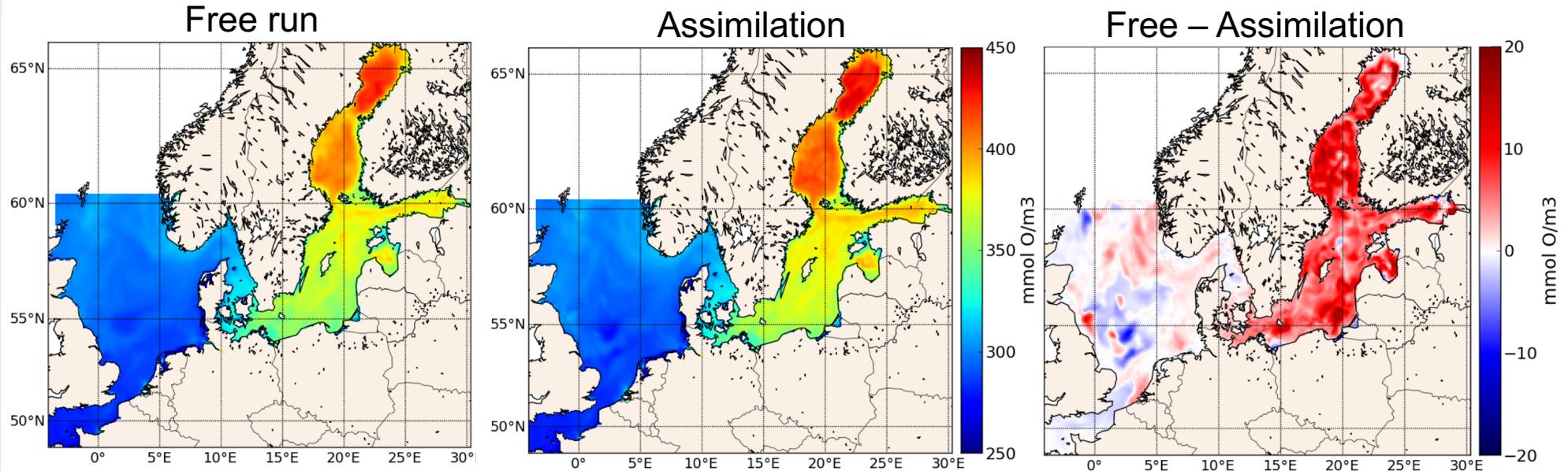
Reduce difference to assimilated data (necessary)

Improve also temperature at 2000m depth



# Biogeochemistry: Coupled data assimilation effect

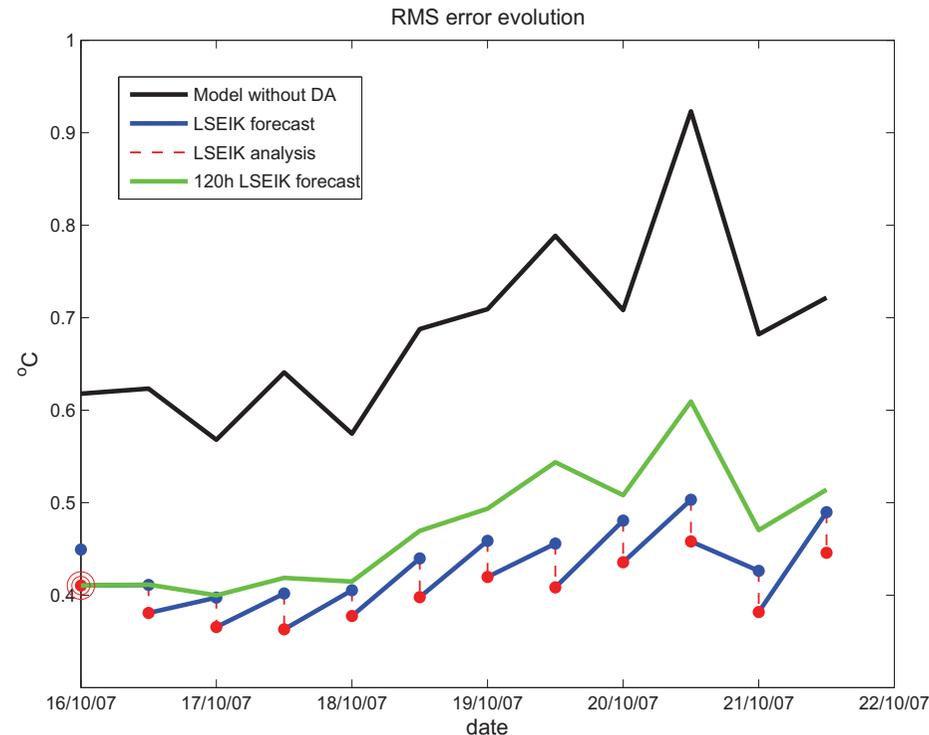
Surface oxygen mean for May 2012 (as mmol O / m<sup>3</sup>)



Coupled data assimilation case: physics and biogeochemistry

- Assimilate satellite sea surface temperature observations
- Assimilation directly changes Oxygen and other biogeochemical variables (strongly-coupled assimilation)

## Impact of Assimilation for temperature forecasts (North & Baltic Seas)

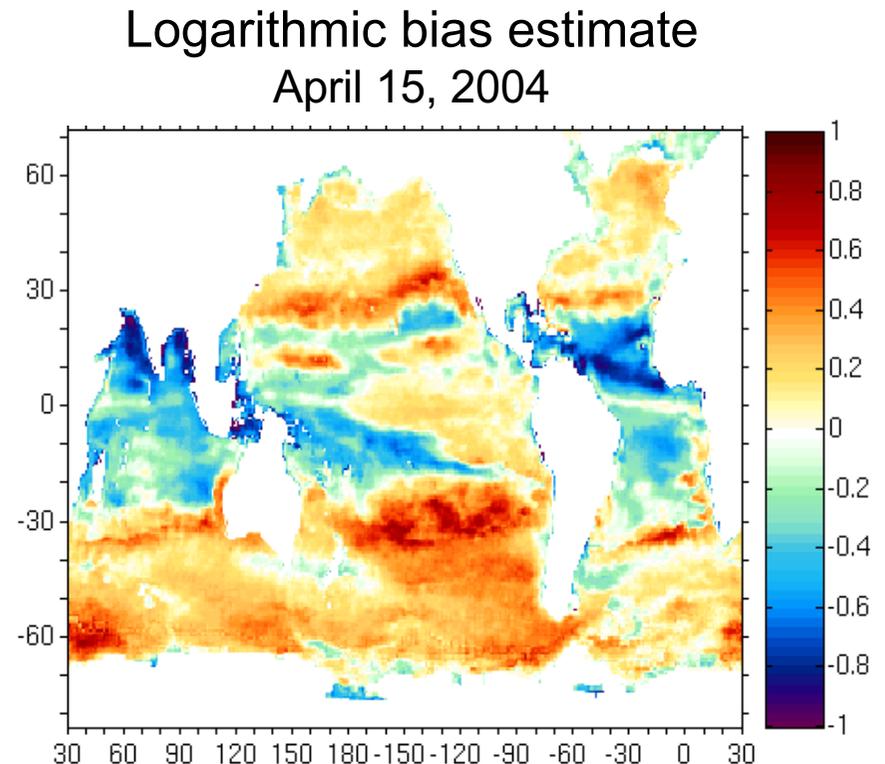


- Very stable 5-days forecasts
- At some point the improvement might break down due to dynamics

# Bias Estimation

**Example:** Chlorophyll bias of a biogeochemical model

- *un-biased system:*  
random fluctuation around true state
- *biased system:*  
systematic over- and underestimation  
(common situation with real data)
- *Bias estimation:*  
Separate random from systematic deviations

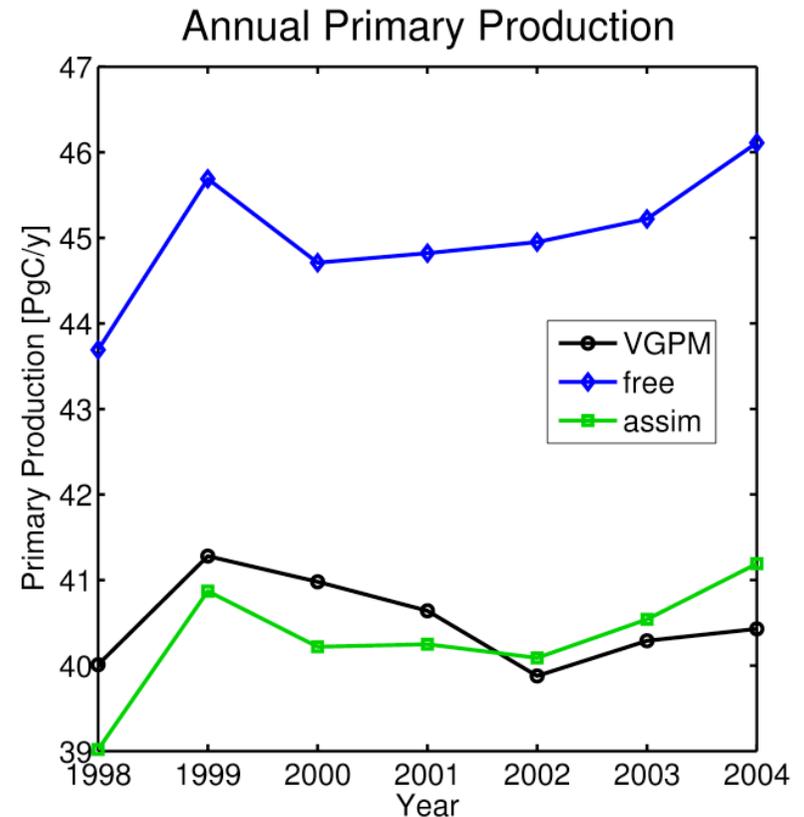


## Estimate a flux (Primary Production)

- *Primary production is a flux*: Uptake of carbon by phytoplankton
- Model: computed as depth-integrated product of growth-rate times Carbon-to-Chlorophyll ratio
- VGPM: Vertical Generalized Production model - satellite data only
- Primary production from assimilation consistent with VGPM-estimate
- *Important*: Concentration change by assimilation is not primary production

(VGPM: Behrenfeld, M.J., P.G. Falkowski., Limnol. Oce. 42 (1997) 1-20)

L. Nerger & W.W. Gregg, *J. Marine Syst.* 68 (2007) 237-254



Mean relative difference to VGPM:  
Free: 11.2%  
Assimilation: -0.5%

# Data Assimilation

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## Combine Models and Observations

# Data Assimilation

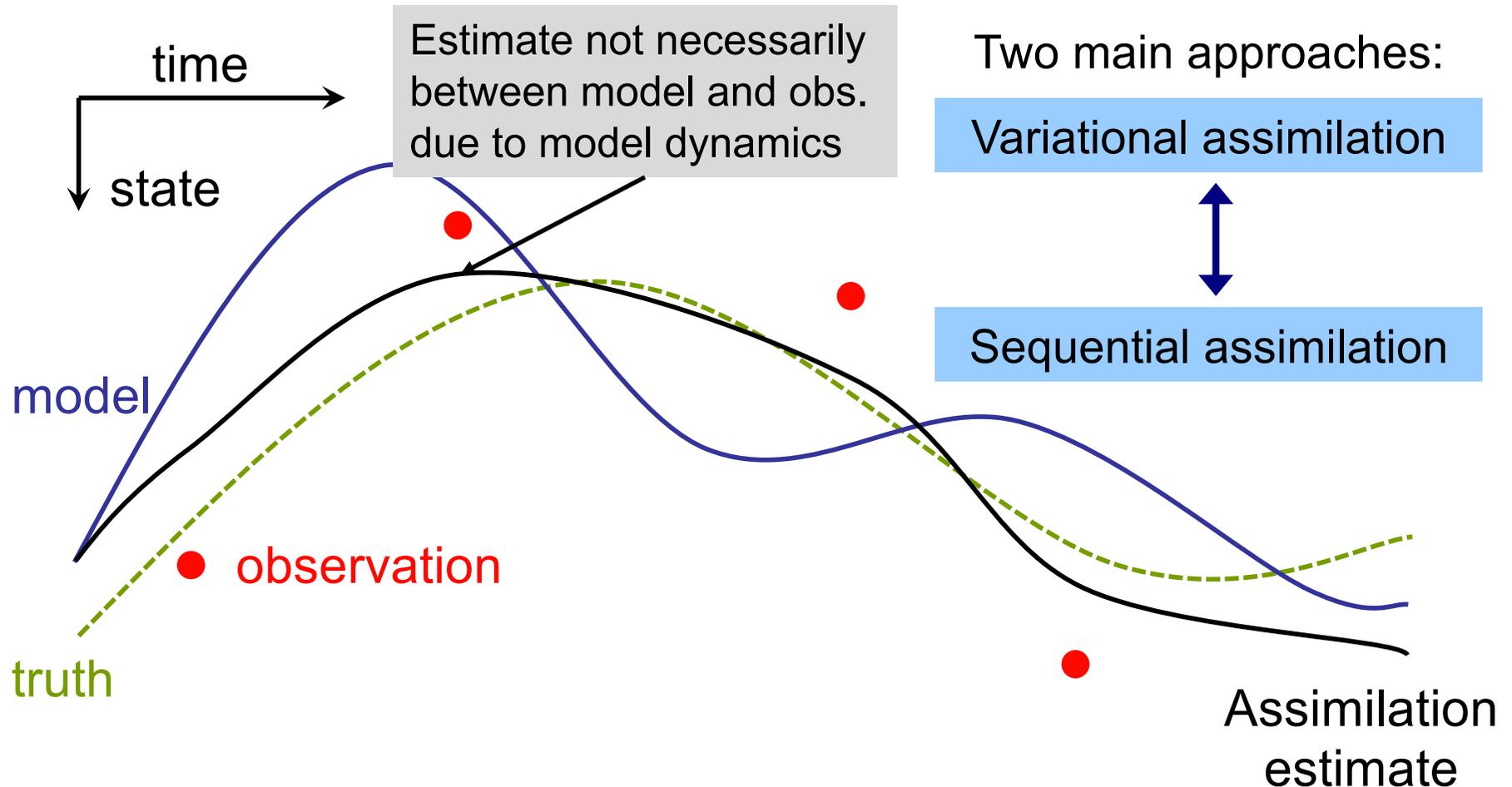
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Combine model with real data

- Optimal estimation of system state:
  - initial conditions (for weather/ocean forecasts, ...)
  - state trajectory (temperature, concentrations, ...)
  - parameters (growth of phytoplankton, ...)
  - fluxes (heat, primary production, ...)
  - boundary conditions and ‘forcing’ (wind stress, ...)
- More advanced: Improvement of model formulation
  - Detect systematic errors (bias)
  - Revise parameterizations based on parameter estimates

# Data Assimilation – a general view

Consider some physical system (ocean, atmosphere, land, ...)



Optimal estimate basically by least-squares fitting  
(but constrained by model dynamics)

# Needed for Data assimilation

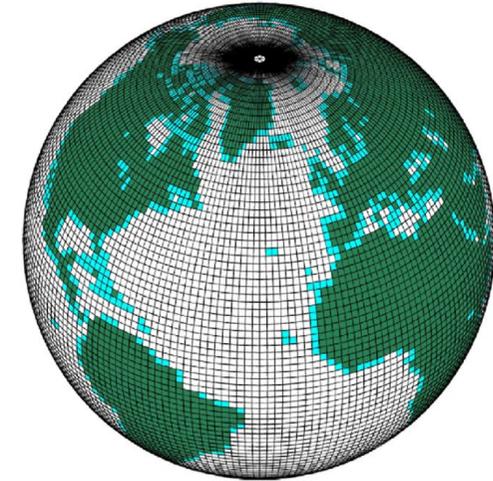
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1. Model
  - with some skill
2. Observations
  - with finite errors
  - related to model fields
3. Data assimilation method

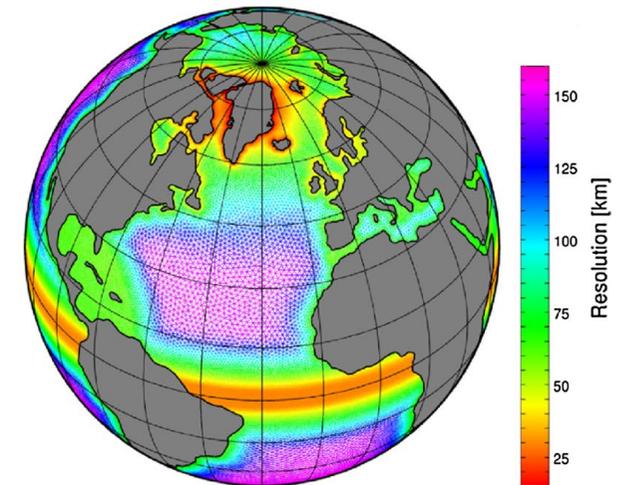
# Models

## Simulate dynamics of ocean

- Numerical formulation of relevant terms
- Discretization with finite resolution in time and space
- “forced” by external sources (atmosphere, river inflows)
- Uncertainties
  - initial model fields
  - external forcing
  - in predictions due to model formulation



*Uniform-resolution mesh*



*Variable-resolution mesh  
(ocean model FESOM)*

# Observations

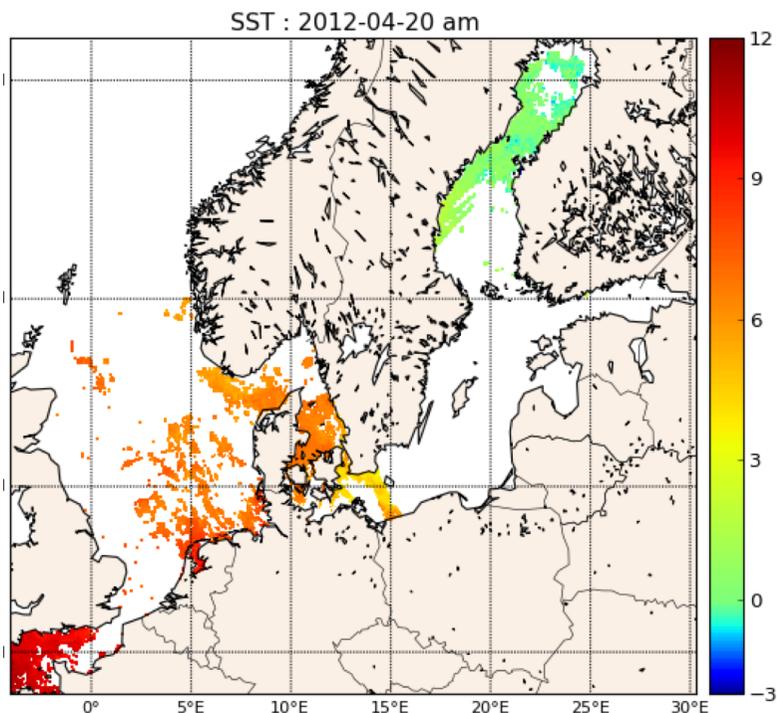
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Measure different fields ... for example in the Ocean

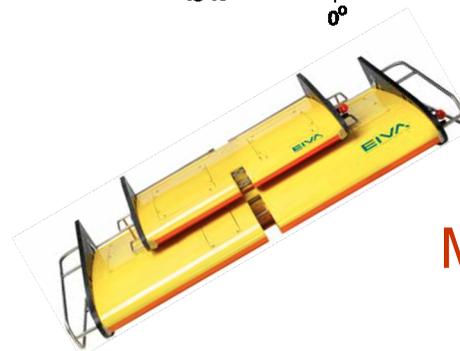
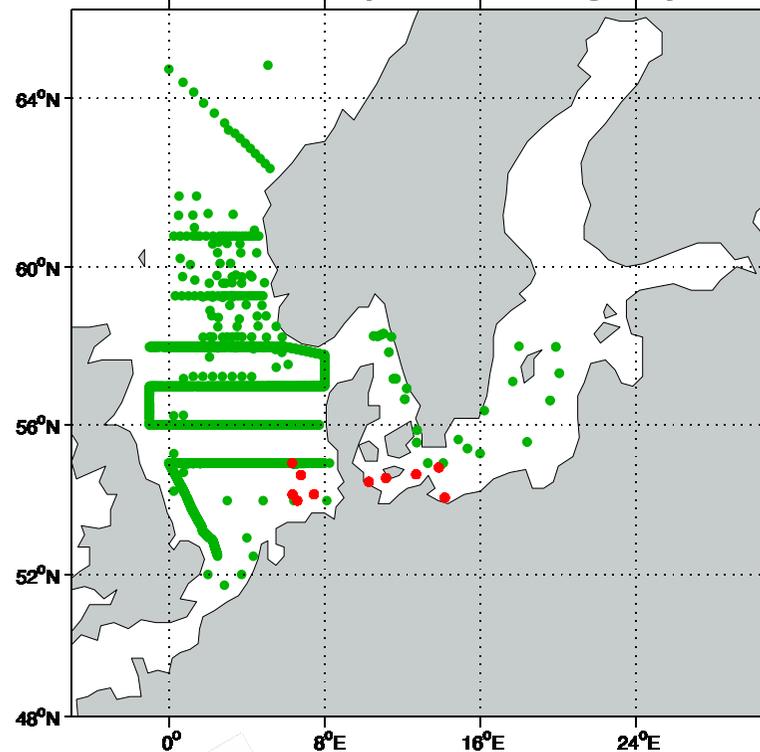
- Remote sensing
  - E.g. surface temperature, salinity, sea surface height, ocean color, sea ice concentrations & thickness
- In situ (ships, autonomous vehicles, ...)
  - Argo, CTD, Gliders, ...
- Data is sparse: some fields, data gaps
- Uncertainties
  - Measurement errors
  - Representation errors:  
Model and data do not represent exactly the same  
(e.g. cause by finite model resolution)

# Example: Physical Data in North & Baltic Seas

Satellite surface temperature  
(12-hour composite)



Available T and S profiles during July 2008



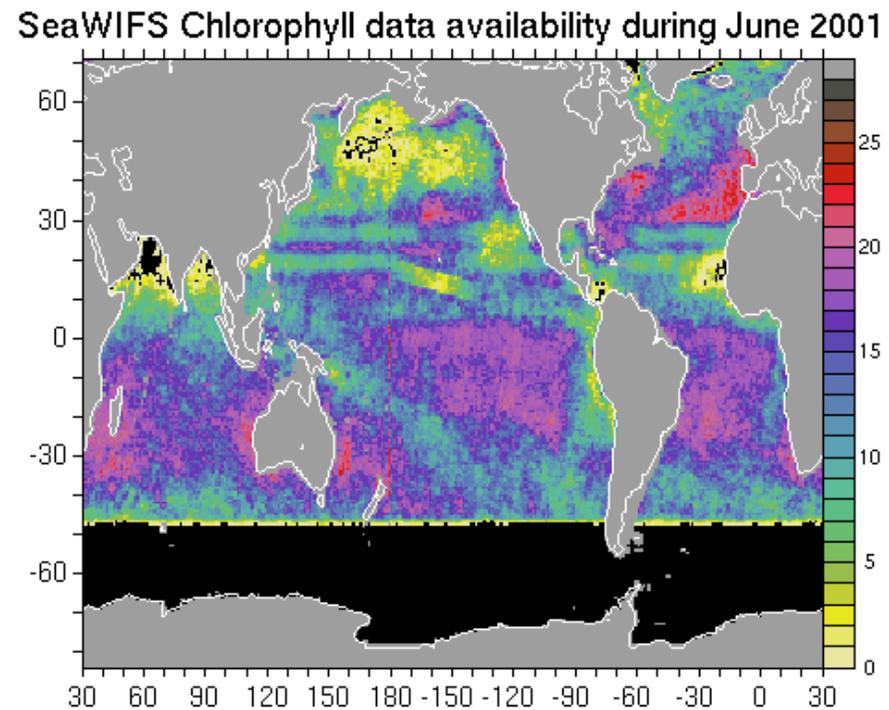
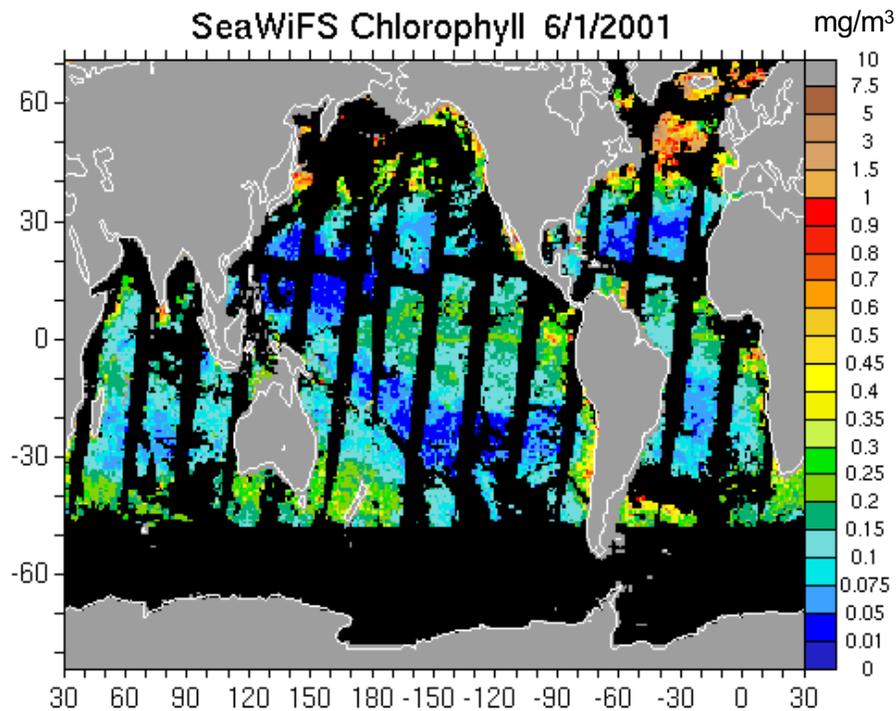
Scanfish and  
CTD profiles

MARNET  
stations



*PDAF*  
Parallel  
Data  
Assimilation  
Framework

## Example: Chlorophyll-a (SeaWiFS)



### Daily gridded SeaWiFS chlorophyll data

- gaps: satellite track, clouds, polar nights
- On model grid: ~13,000-18,000 data points daily (of 41,000 wet grid points)
- irregular data availability

# Observation Error Estimates

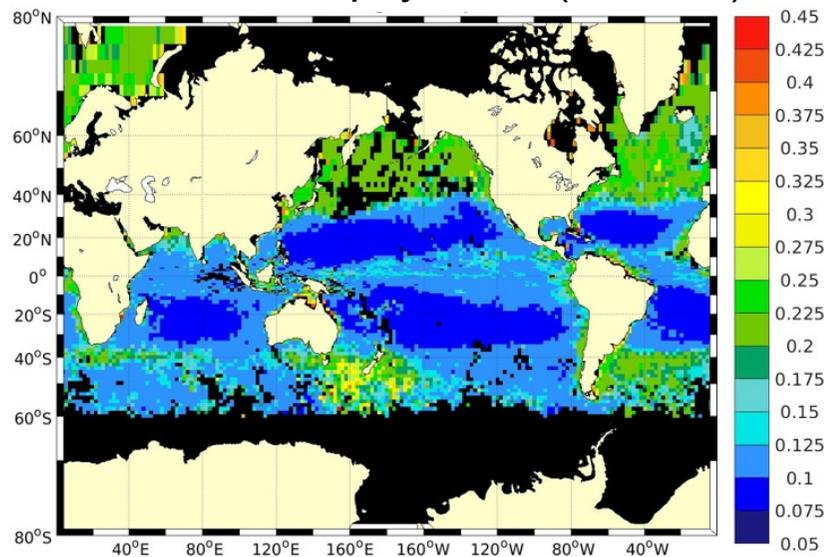
If observation errors available:

- they are typically usable
- usually do not account for representation errors (might be too low)

If no observation errors available:

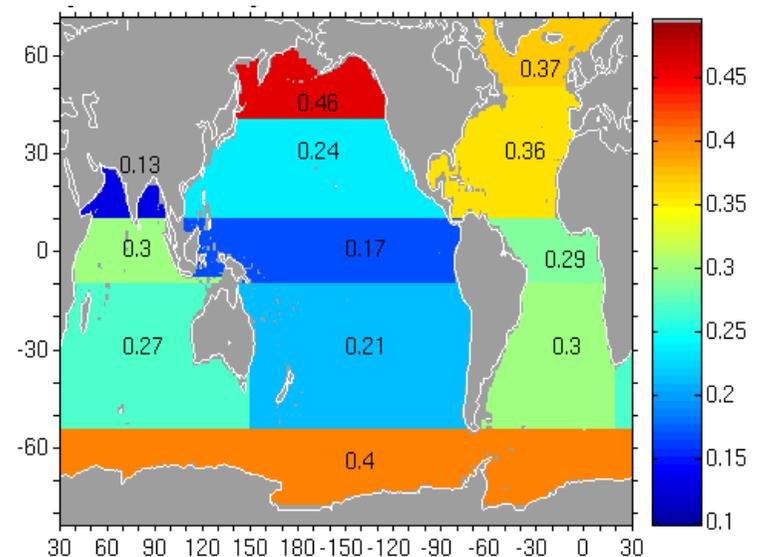
- need to estimate them

logarithmic data errors provided with satellite chlorophyll data (OC-CCI)



Pradhan et al, JGR 2019

data errors from comparison with 2186 collocation points of in situ data (SeaWiFS)



Nerger & Gregg, JMS 2007

# Data Assimilation Methods

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## Combine observations and model state estimate

- Account for uncertainty in observations
- Account for uncertainty in model state estimate
- Account for relations (correlations) between observed part of the model state and unobserved parts

# Ensemble Data Assimilation

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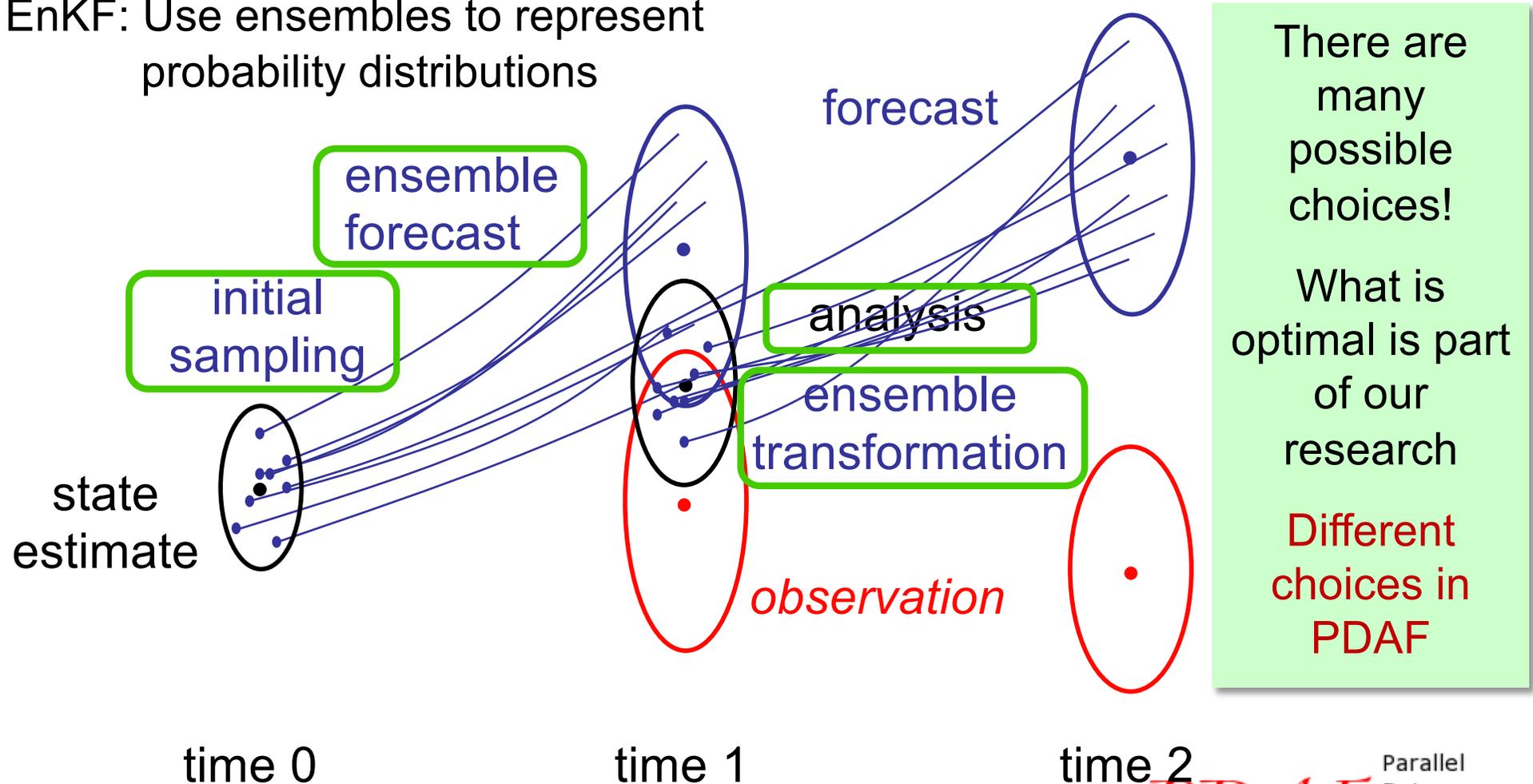
**Estimate uncertainty**

# Ensemble Kalman Filters

First formulated by G. Evensen (EnKF, J. Geophys. Res. 1994)

Kalman filter: express probability distributions by mean and covariance matrix

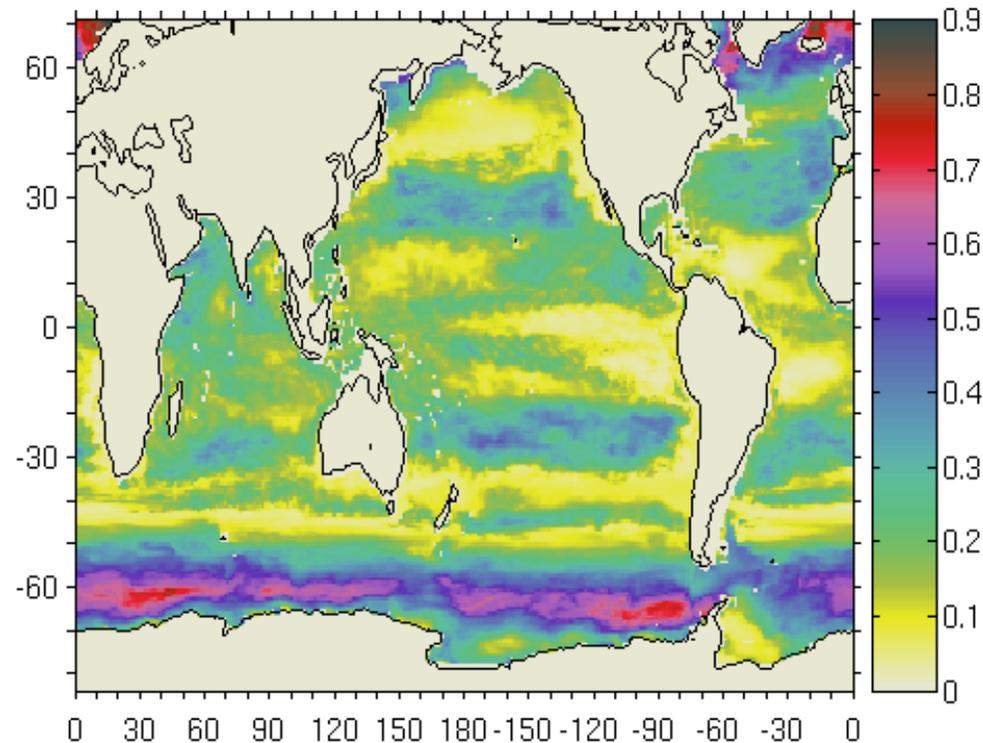
EnKF: Use ensembles to represent probability distributions



# Ensemble Covariance Matrix

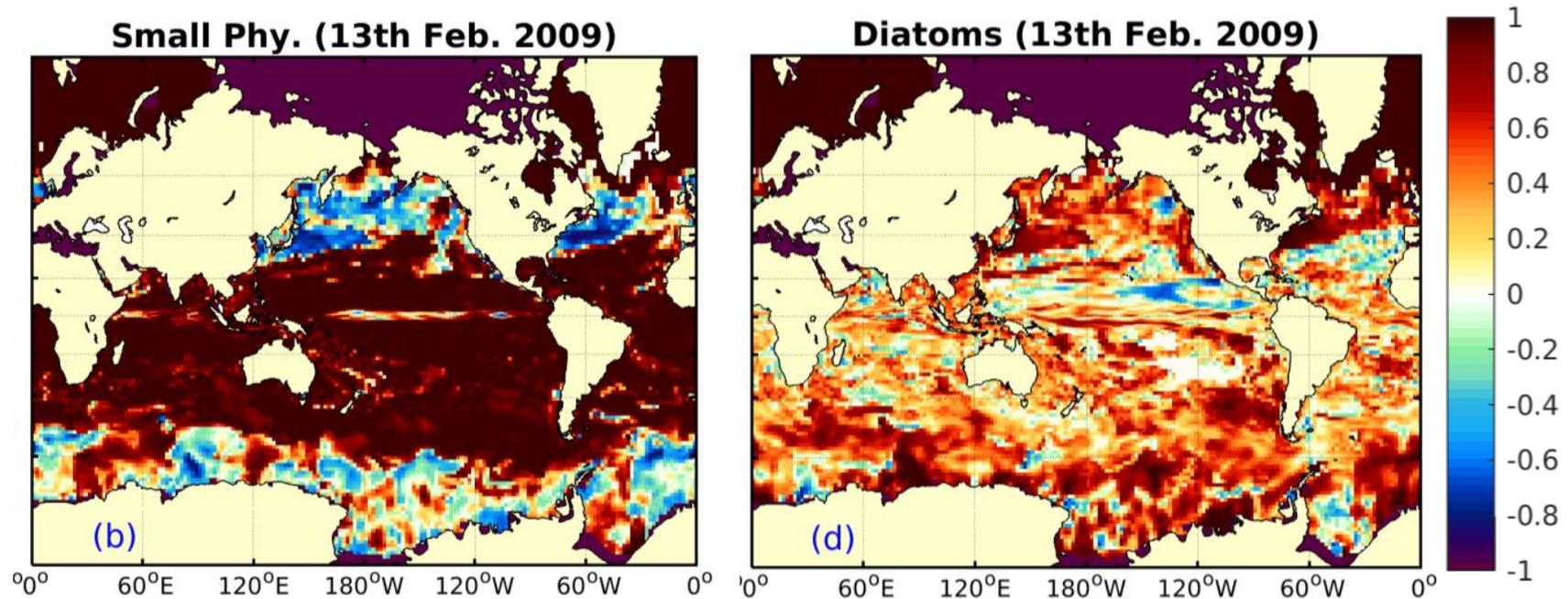
- Provide uncertainty information (variances + covariances)
- Generated dynamically by propagating ensemble of model states

Uncertainty: Std. deviation of log Chlorophyll



# Ensemble-estimated Cross-correlations

Cross correlations between total chlorophyll and chlorophyll in phytoplankton groups



Cross-correlations are used to correct non-observed quantities from observed ones

# Validation of assimilation results

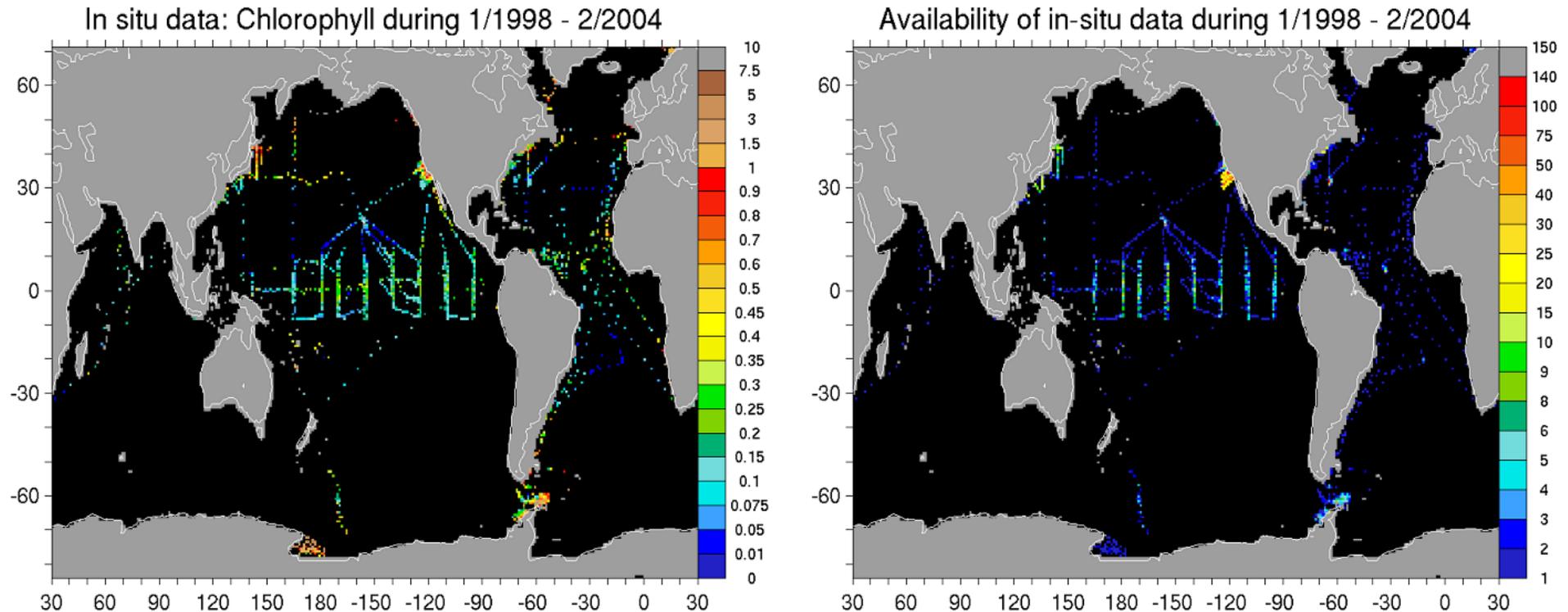
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# Validating a data assimilation system

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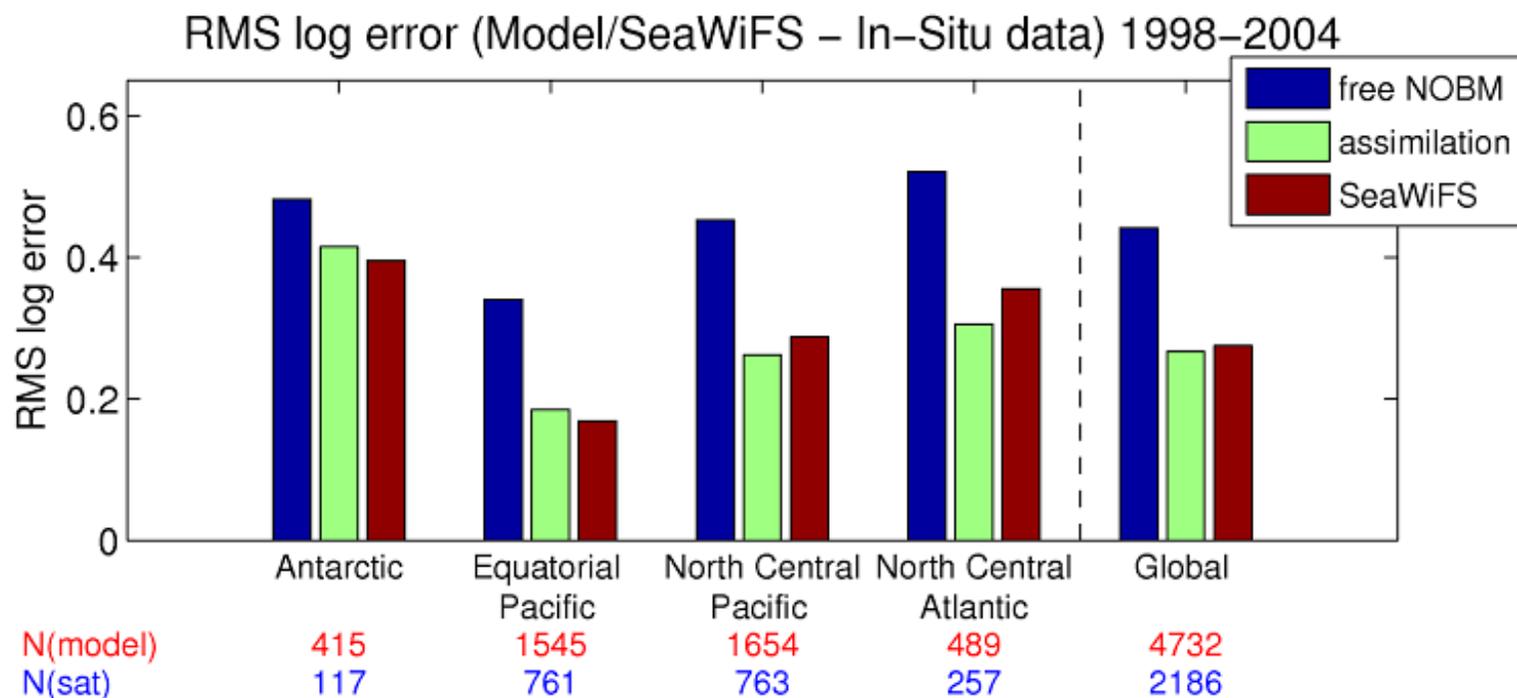
- Need independent data for validation
  - Necessary, but not sufficient:  
Reduction of deviation from assimilated data
  - Required:
    - Reduction of deviation from independent data
    - Reduction of errors for unobserved variables
  - Ideally:
    - Reduce error below that of model and data alone
- Want to assimilate all available data (in the ocean)
  - Data-withholding experiments
  - Twin experiments
  - Validate with data of small influence

# Validation: In-Situ chlorophyll data



- In situ data from SeaBASS/NODC over 1/1998-2/2004
- Independent from SeaWiFS data (only used for verification of algorithms)
- North Central Pacific dominated by CalCOFI data
- North Central Atlantic dominated by BATS data

## Comparison with independent data



- Shown basins include about 87% of data
- Compare daily co-located data points
- ⇒ Assimilation reduces errors significantly
- ⇒ Error from assimilation lower than SeaWiFS error in many basins and globally

# Quantifying the quality of the assimilation result

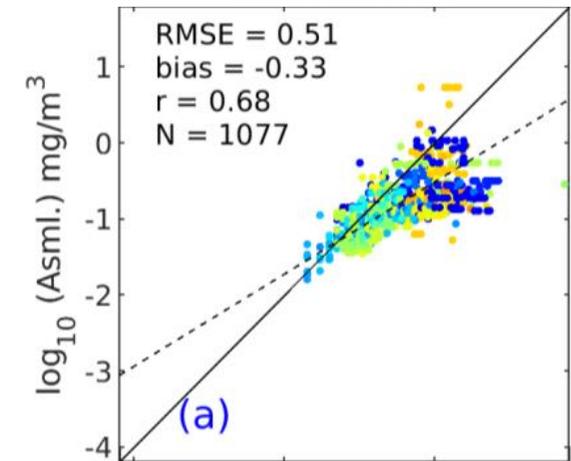
## Assess ensemble mean state:

### Common choices

- RMS (root mean square) errors
- Bias (mean error)
- Correlation

compared to observations

## Scatter plot for validation

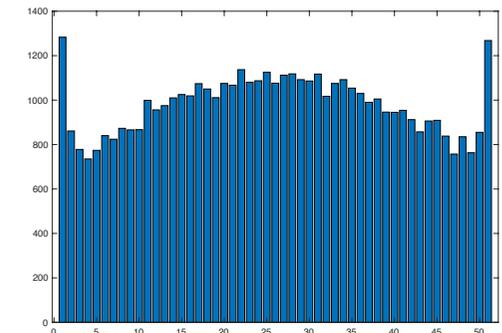


## Assess ensemble quality:

- Rank histogram
- CRPS (continuous ranked probability score)
- Relative entropy

Particularly relevant when using nonlinear assimilation methods (e.g. particle filters)

## Rank histogram, N=50



# Essential “Fixes” for Ensemble Filters

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**Covariance Inflation**

**Localization**

# Covariance inflation

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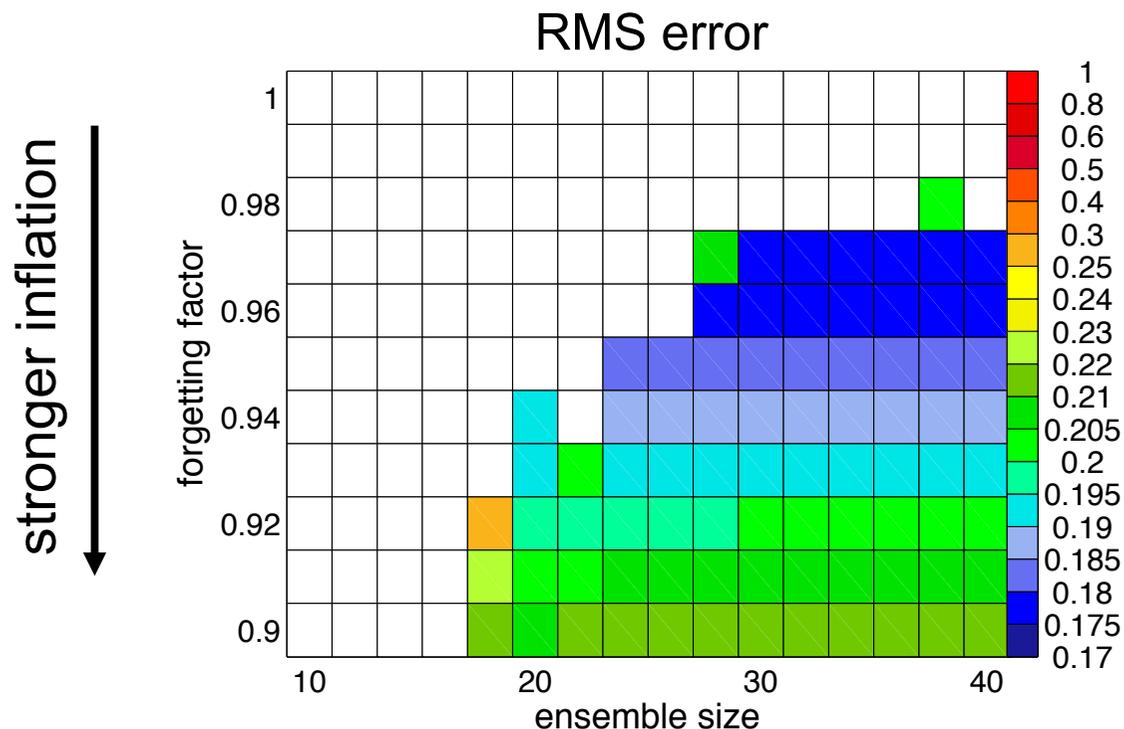
- True variance is always underestimated
  - small ensemble size
  - sampling errors (unknown structure of P)
  - model errors

→ can lead to filter divergence
- Simple remedy
  - Increase error estimate before analysis
- Inflation
  - Increase ensemble spread by constant factor
  - Some filters allow multiplication of a small matrix (“forgetting factor”  $\leq 1$ ; computationally very efficient)
  - Needs to be experimentally tuned

(Mathematically, this is a regularization)

# Impact of inflation on stability & performance

Experiments with Lorenz96 model  
(available with PDAF)



Lorenz96 model

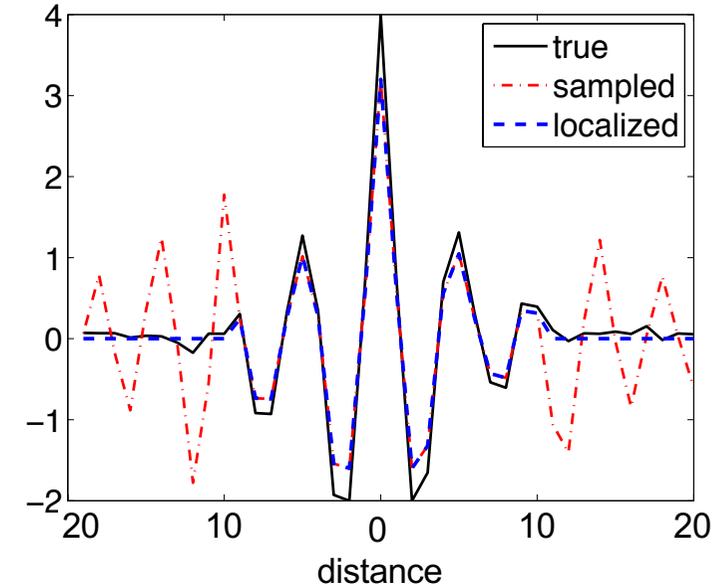
- a widely used toy model
- one-dimensional period wave
- chaotic dynamics
- included in PDAF release

- white: filter fails („diverges“)
- increased stability with stronger inflation (smaller forgetting factor)
- optimal choice for inflation factor

# Localization: Why and how?

- Combination of observations and model state based on estimated error covariance matrices
- Finite ensemble size leads to significant sampling errors
  - particularly for small covariances!
- Remove estimated long-range correlations
  - Increases degrees of freedom for analysis (globally not locally!)
  - Increases size of analysis correction

Example: Sampling error and localization

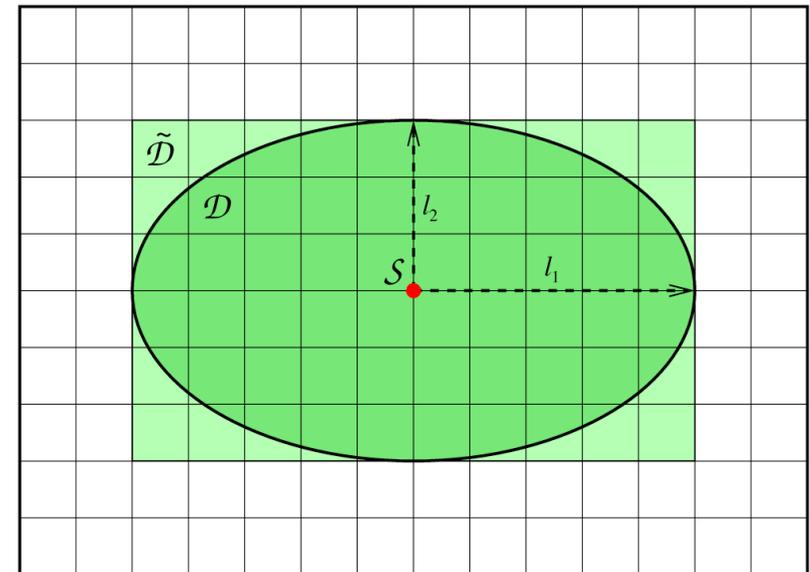


(introduced for EnKFs by Houtekamer & Mitchell 1998)

# Observation Localization

## Local Analysis:

- Update small regions (like single vertical columns) allows to define distance
- Use only observations within some distance around this region
- State update and ensemble transformation fully local



S: Analysis region

D: Corresponding data region

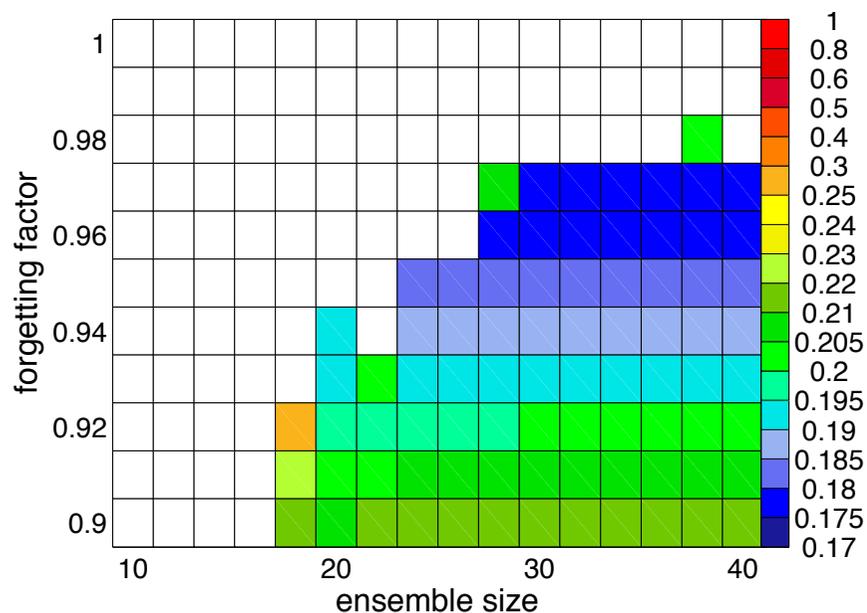
## Observation localization:

- Down-weight observations with increasing distance

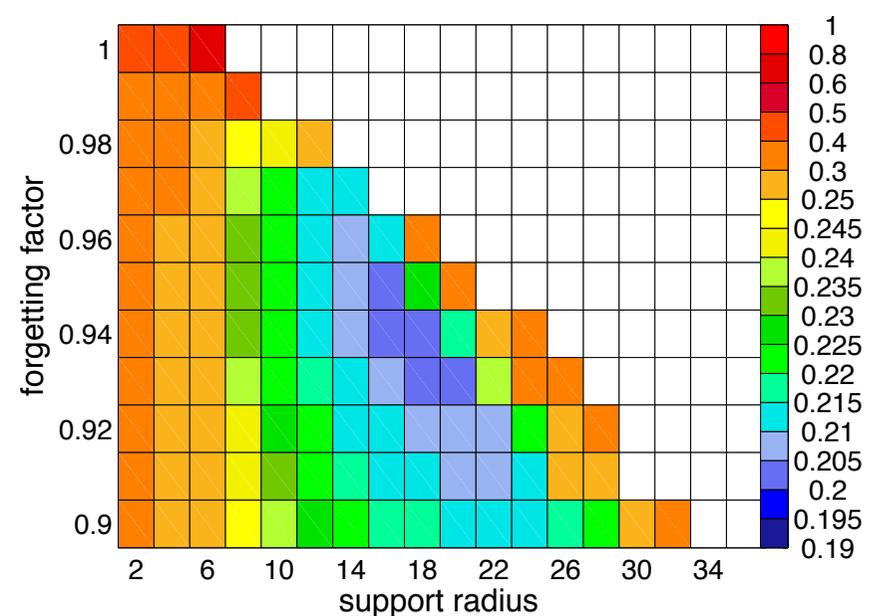
# Impact of inflation and localization

## Experiments with Lorenz96 model

Global filter



Localized, ensemble size 10



- smaller ensemble usable with localization
- optimal combination of forgetting factor and support radius

# Overview

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(See Short Course SC1.2 on Friday for methodology)

# 2

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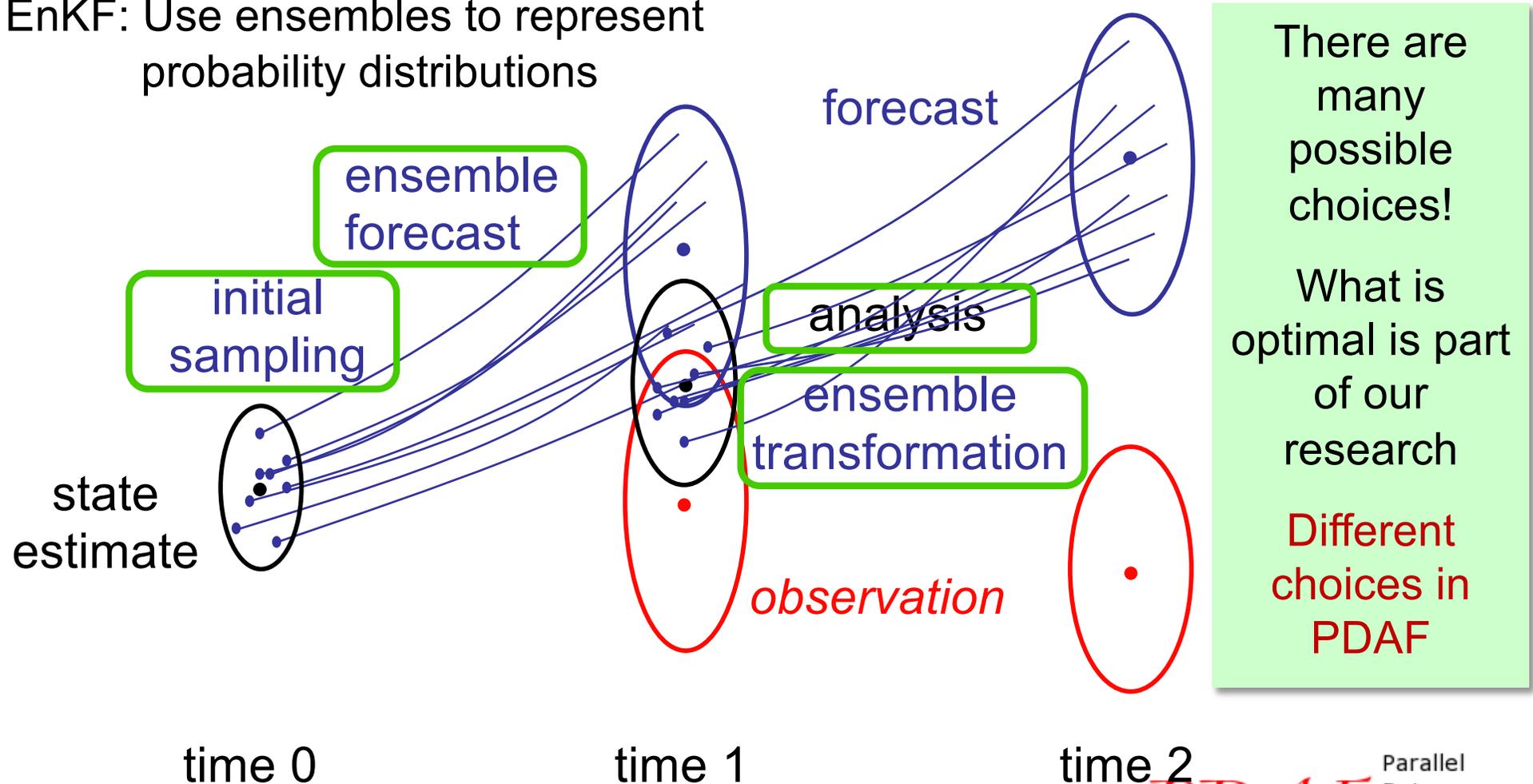
## Implementation Concept of PDAF (Parallel Data Assimilation Framework)

# Ensemble-based Kalman Filter

First formulated by G. Evensen (EnKF, J. Geophys. Res. 1994)

Kalman filter: express probability distributions by mean and covariance matrix

EnKF: Use ensembles to represent probability distributions



# Computational and Practical Issues

- Running a whole model ensemble is costly
- Ensemble propagation is naturally parallel (all independent)
- Ensemble data assimilation methods need tuning
- No need to go into model numerics (just model forecasts)
- Filter step of assimilation only needs to know:
  - Values of model fields and their location
  - Observed values, their location and uncertainty

→ Ensemble data assimilation can be implemented in form of a generic code + case-specific routines

→ Can be used without knowing the exact details of the filter algorithm

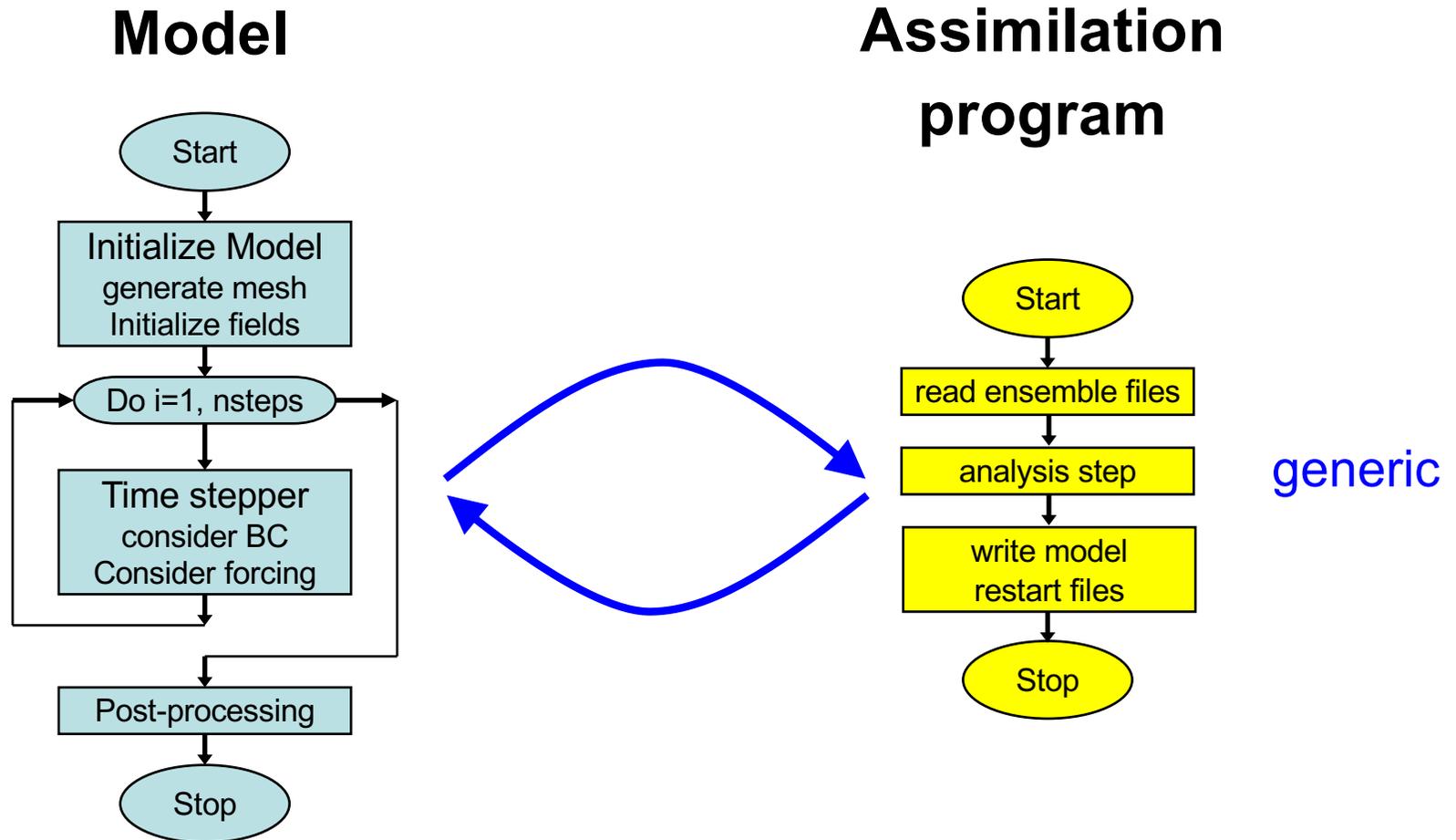
## PDAF - Parallel Data Assimilation Framework

- a program library for ensemble data assimilation
- provide support for parallel ensemble forecasts
- provide fully-implemented & parallelized filters and smoothers (EnKF, LETKF, NETF, EWPF ... easy to add more)
- easily useable with (probably) any numerical model (applied with MITgcm, NEMO, FESOM, HBM, TerrSysMP, ...)
- run from laptops to supercomputers (Fortran, MPI & OpenMP)
- first public release in 2004; continued development
- ~350 registered users; community contributions

Open source:  
Code and documentation available at

<http://pdaf.awi.de>

# Offline coupling – separate programs



- For each ensemble state
- Initialize from restart files
  - Integrate
  - Write restart files

- Read restart files (ensemble)
- Compute analysis step
- Write new restart files

# Offline coupling - Efficiency

Offline-coupling is simple to implement but can be very inefficient

## Example:

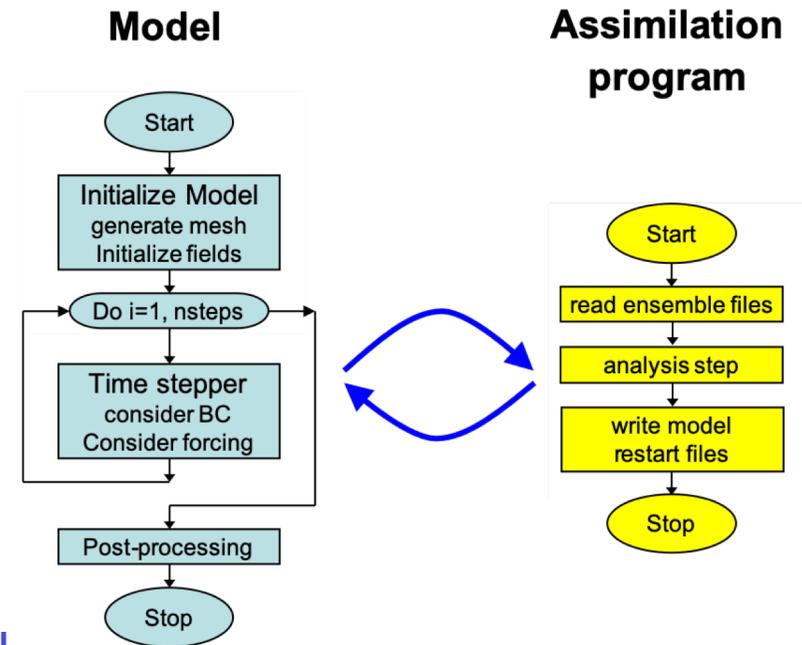
Timing from atmosphere-ocean coupled model (AWI-CM) with daily analysis step:

Model startup:	95 s	} overhead
Integrate 1 day:	28 s	
Model postprocessing:	14 s	

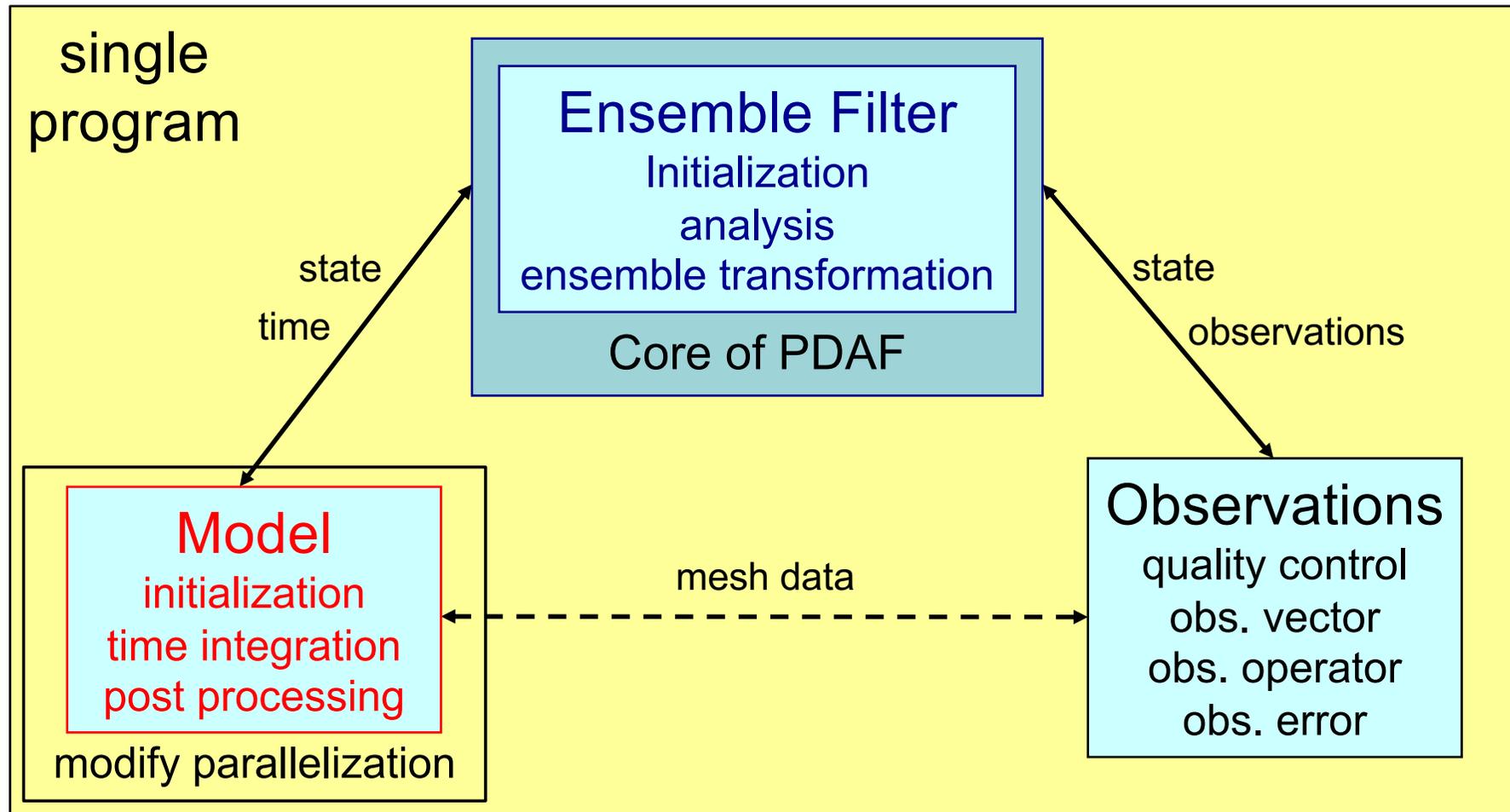
Analysis step: 1 s

Restarting this model is ~3.5 times more expensive than integrating 1 day

→ avoid this for data assimilation



# Components of an Assimilation System

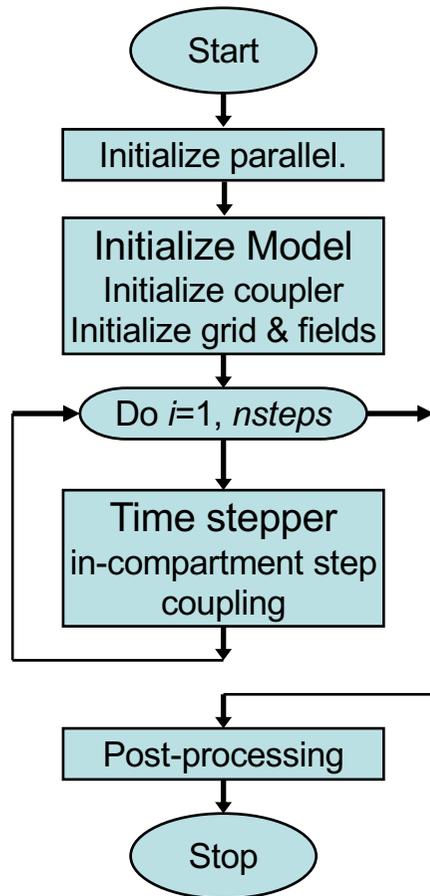


↔ Explicit interface

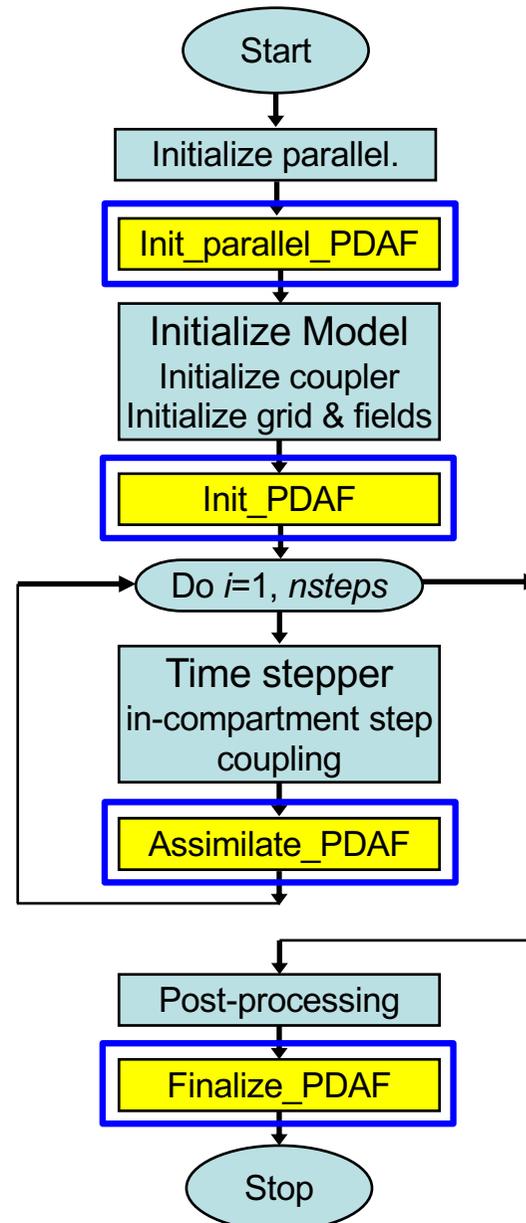
⋯ Indirect exchange (module/common)

# Extending a Model for Data Assimilation

**Model**  
*single or multiple executables*  
*coupler might be separate program*



revised parallelization enables ensemble forecast



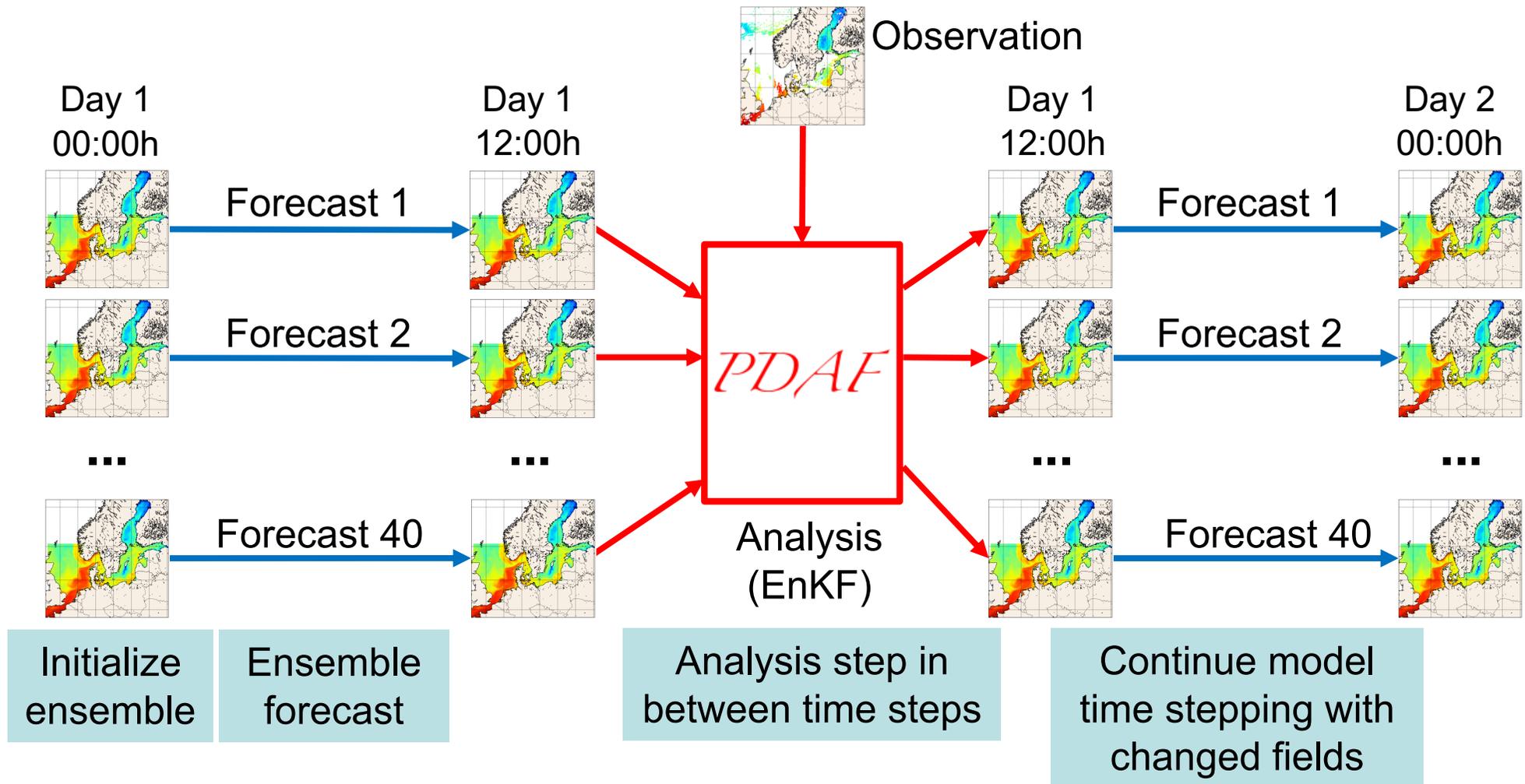
Extension for data assimilation

*plus:*  
 Possible model-specific adaption  
 for MITgcm:  
 adapt name of STDOUT files for ensemble

# Augmenting a Model for Data Assimilation

Couple PDAF (Parallel Data Assimilation Framework) with model

- Modify model to simulate ensemble of model states
- Insert correction step (analysis) to be executed each 12 model hours
- Run model as usual, but with more processors and additional options



# PDAF model binding routines

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## Interface routines

- `init_parallel_pdaf`, `init_pdaf`, `assimilate_pdaf`,  
`finalize_pdaf`

## Call-back routines

- Set number of time steps between analysis steps
- Write model fields into PDAF's state vector and back into model fields
- Observation handling

## PDAF release includes set of model binding routines for MITgcm

- for a simple test case
- just download and adapt for your needs
- (NEMO will be next)

## PDAF interface structure

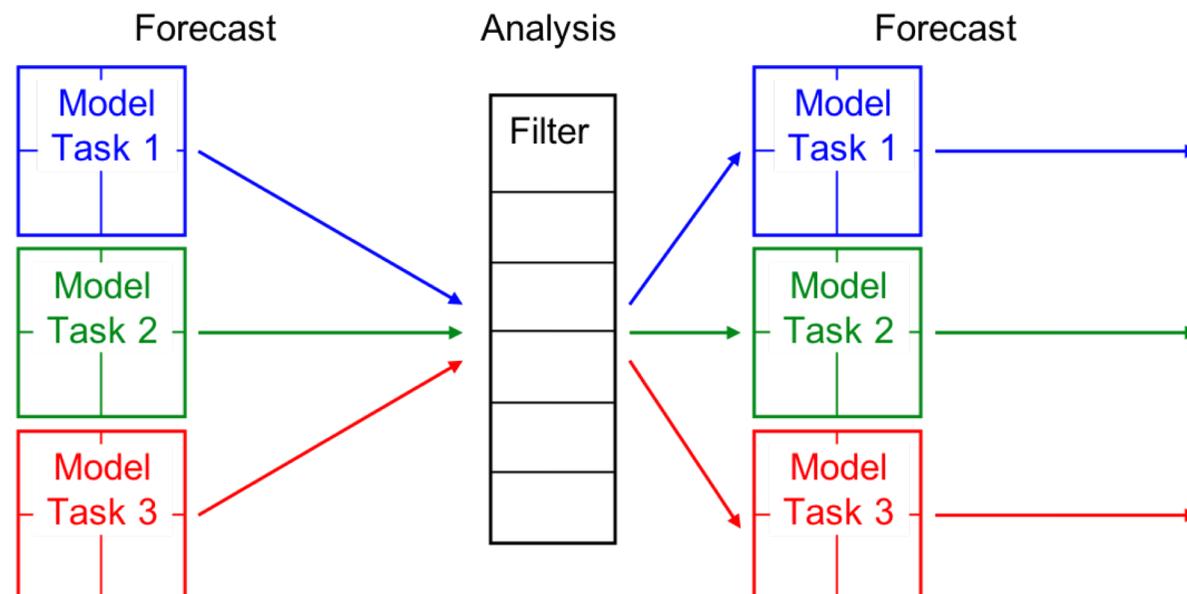
- Interface routines call PDAF-core routines
- PDAF-core routines call case-specific routines provided by user (included in model binding set)
- User-supplied call-back routines for elementary operations:
  - field transformations between model and filter
  - observation-related operations
- User supplied routines can be implemented as routines of the model  
(for MITgcm: Fortran-77 fixed-form source code)



We use MPI (Message Passing Interface)

- standard for highly scaling parallelization
- used by most large-scale models

Only need to do this once for a model (e.g. done for MITgcm)



Set parameters, for example

- select filter
- set ensemble size

Calls `PDAF_init`

- initialization routine of framework
- provide parameters according to interface
- provide MPI communicators
- provide name of routine for ensemble initialization

Ensemble initialization routine – called by `PDAF_init`

- a “call-back routine”
- defined interface: provides ensemble array for initialization
- user-defined initialization

# Simple Subroutine Interfaces

Example: ensemble initialization

```
SUBROUTINE init_ens_pdaf(filtertype, dim, dim_ens, state,  
matrU, ens, flag)
```

```
IMPLICIT NONE
```

```
! ARGUMENTS:
```

```
INTEGER, INTENT(in) :: filtertype ! Type of filter  
INTEGER, INTENT(in) :: dim        ! Size of state vector  
INTEGER, INTENT(in) :: dim_ens   ! Size of ensemble  
REAL, INTENT(out)  :: ens(dim, dim_ens) ! state ensemble  
INTEGER, INTENT(inout) :: flag    ! PDAF status flag
```

Task to be implemented:

➤ Fill `ens` with ensemble of initial model states

*calls* PDAF\_assimilate

- checks whether ensemble integration reached time for analysis step
- **If false:**
  - return to model and continue integration
- **If true:**
  - Write forecast fields into state vectors (call-back routine)
  - Compute analysis step of chosen filter
  - Set length of next forecast phase (call-back routine)
  - Write state vectors into model field arrays (call-back routine)

Clean-up at end of program

- Display timing and memory information for PDAF
- Deallocate arrays inside PDAF

Calls to

`PDAF_print_info` (memory and timing info)

`PDAF_deallocate` (deallocate arrays)

# Filter analysis implementation

---

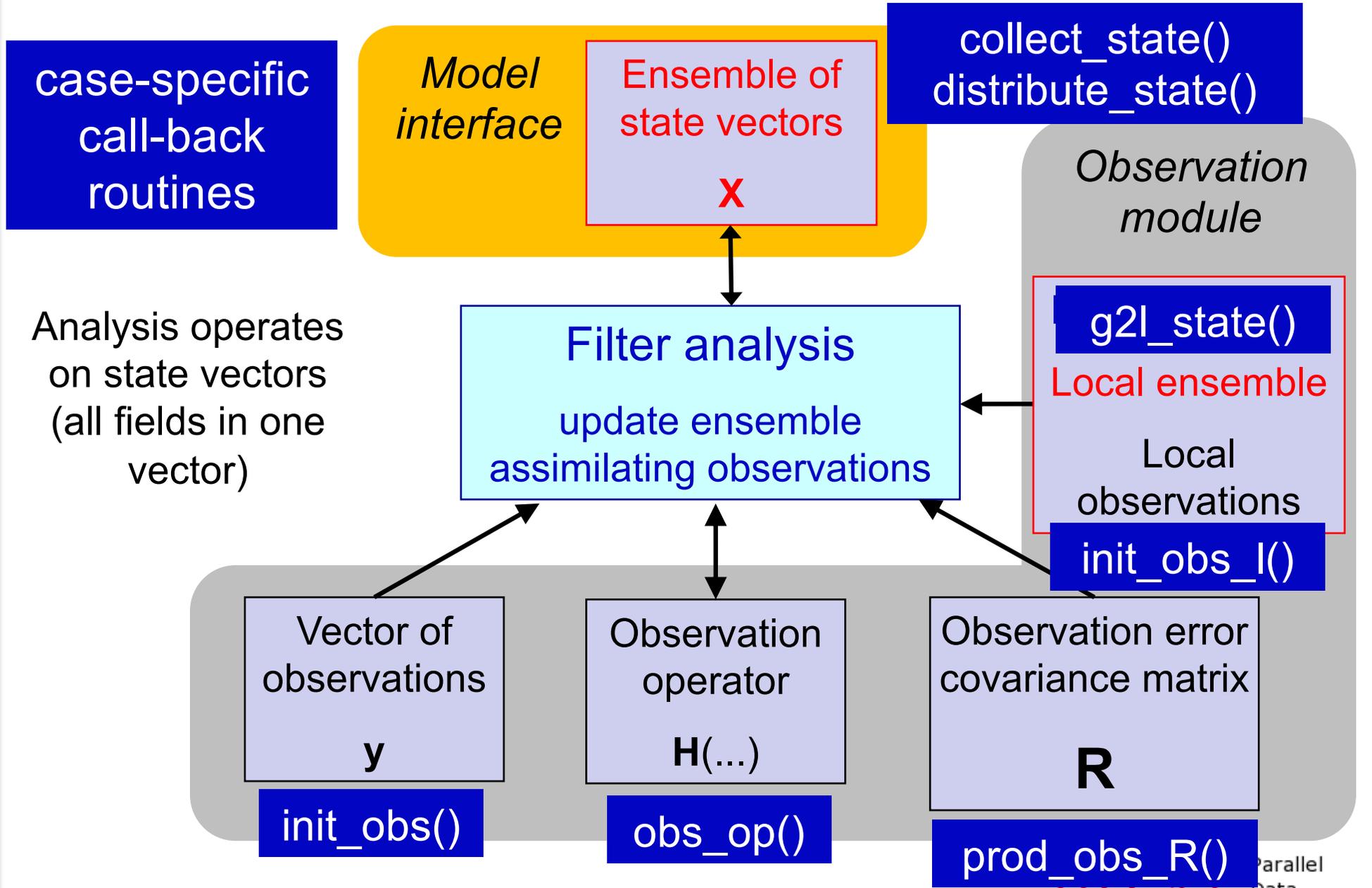
Operate on state vectors

- Write all model fields into a 1-dimensional vector
  - Filter doesn't know about 'fields'
  - Computationally most efficient
  - Call-back routines for
    - Transfer between model fields and state vector
    - Observation-related operations
    - Localization operations

For forecast

- Transfer data from state vector to model fields

# Ensemble Filter Analysis Step



PDAF originated from comparison studies of different filters

## Filters and smoothers

- EnKF (Evensen, 1994 + perturbed obs.)
- ETKF (Bishop et al., 2001)
- SEIK filter (Pham et al., 1998)
- ESTKF (Nerger et al., 2012)
- NETF (Toedter & Ahrens, 2015)

Not yet released:

- serial EnSRF
- particle filter
- EWPF

## All methods include

- global and localized versions
- smoothers

## Model bindings

- MITgcm, Lorenz96

Not yet released:

- NEMO

# Execution times (weakly-coupled, DA only into ocean)

## MPI-tasks

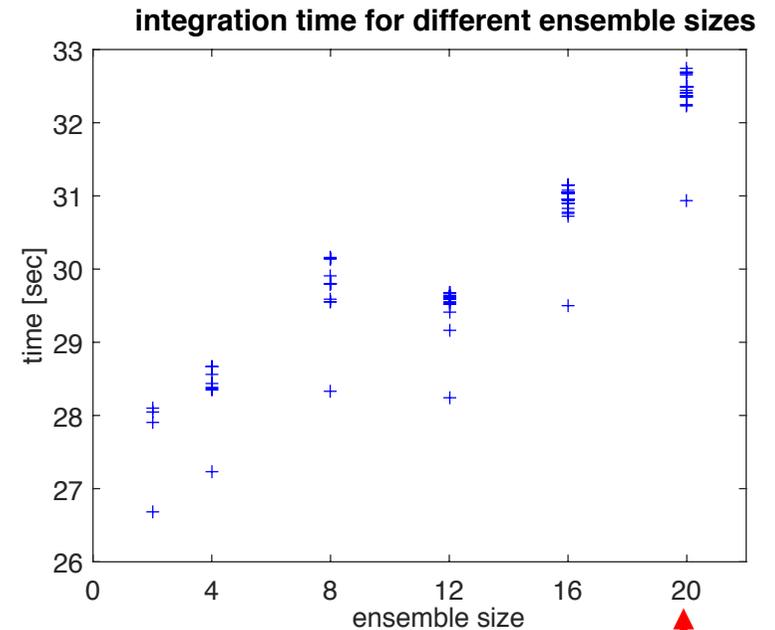
- ECHAM: 144
- FESOM: 384

## Timings (1 day):

- Ens. forecast: 27 – 23 sec
- Analysis step: 0.5 – 0.9 sec

## A remaining issue:

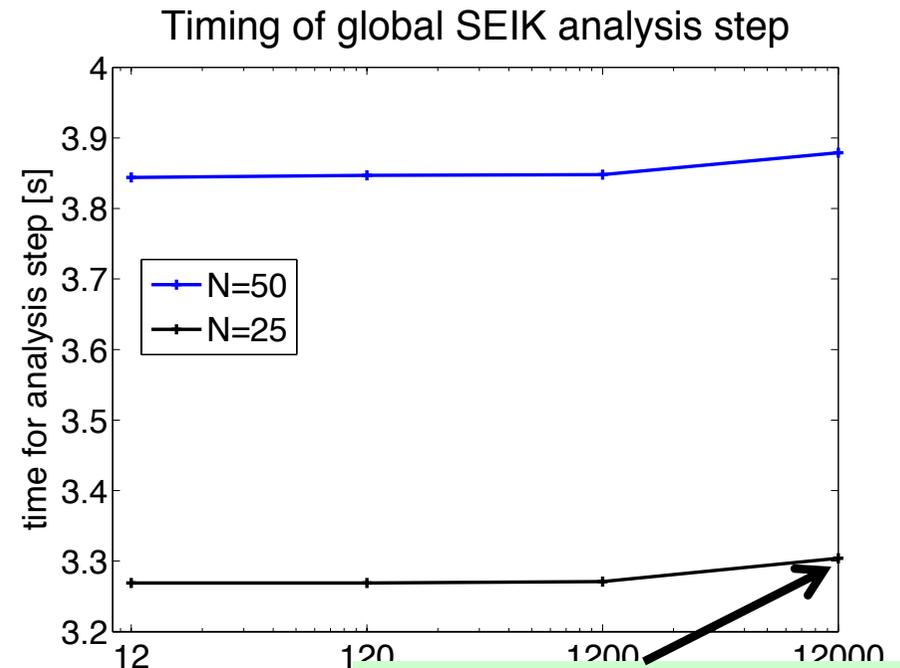
- Increasing integration time with growing ensemble size (only 16% due to more parallel communication)
- some variability in integration time over ensemble tasks
- Need optimal distribution of programs over compute nodes/racks (here set up as ocean/atmosphere pairs)



10,560  
processor  
cores

# Very big test case

- Simulate a “model”
- Choose an ensemble
  - state vector per processor:  $10^7$
  - observations per processor:  $2 \cdot 10^5$
  - Ensemble size: 25
  - 2GB memory per processor
- Apply analysis step for different processor numbers
  - 12 – 120 – 1200 – 12000
- Very small increase in analysis time ( $\sim 1\%$ )  
(Ideal would be constant time)
- Didn't try to run a real ensemble of largest state size (no model yet)



State dimension:  
 $1.2e11$   
Observation  
dimension:  $2.4e9$

# Implementation concept of PDAF

For ensemble data assimilation with PDAF

- Augment program for ensemble data assimilation
- Assimilation methods provided by PDAF
- Model-binding routines required
  - provided for Lorenz96 and for MITgcm for test case
  - easy to code yourself

Next look into an example



pdaf@awi.de

Slides are available online:

<http://pdaf.awi.de>

*PDAF*

Parallel  
Data  
Assimilation  
Framework

# 3

---

## **Hands-on Example: Build an Assimilation System with PDAF**

# Get the tutorial code

---

## Download the tutorial

Directory layout:

`make.arch`

- build configurations

`src`

- source files

`tutorial`

`online_2D_serial`

`model`

- serial model code

`model_coupled_to_pdaf`

- final assimilation code

`pdaf`

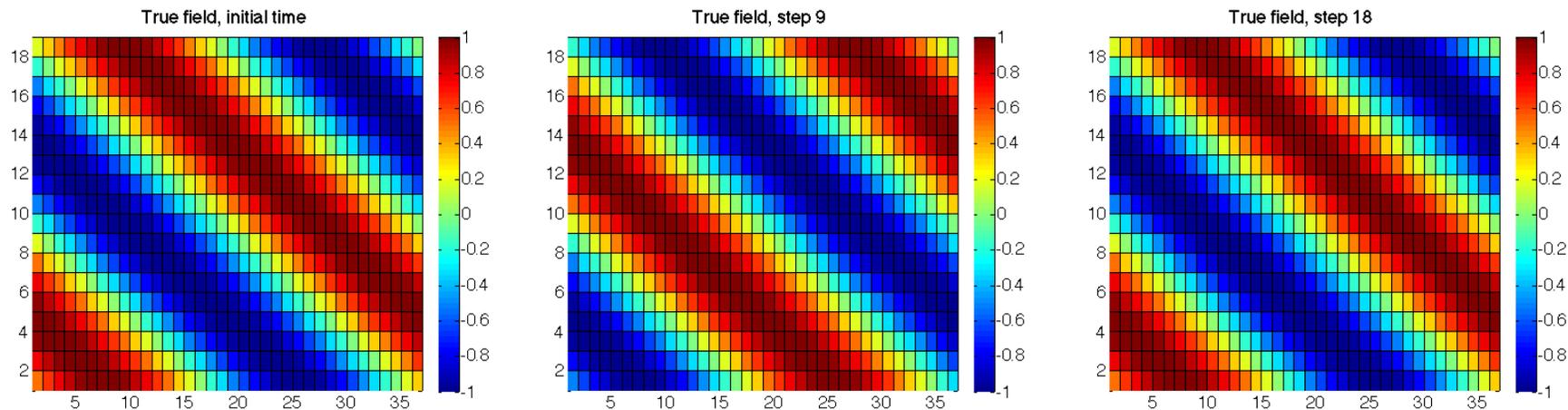
- code to be added to the model

`online_2D_serial.noMPI`

- alternative code without MPI

## 2D „Model“

- Simple 2-dimensional grid domain
- 36 x 18 grid points (longitude x latitude)
- True state: sine wave in diagonal direction (periodic for consistent time stepping)
- Simple time stepping:  
Shift field in vertical direction one grid point per time step
- Output to text files (18 rows) – `true_step*.txt`



## General program structure: model/main.f90

---

```
program main
```

```
  initialize initialize model information:
```

- set dimensions
- allocate model field array
- read initial field

```
  integrate perform time stepping
```

- shift model field
- write new model field

```
end program
```

**No parallelization!**

## Files in the tutorial directories

---

The model source code consists of the following files (`model/`):

- `main.F90`
- `mod_model.F90`
- `initialize.F90`
- `integrate.F90`
- `Makefile`

## Files in the tutorial directories

---

The PDAF coupling code consists of (`pda_f/`)

- interface subroutines (called from the model code)
  - `init_parallel_pda_f.F90`
  - `init_pda_f.F90`
  - `assimilate_pda_f.F90`
  - `finalize_pda_f.F90`
- user subroutines (called from the PDAF library), eg.
  - `collect_state_pda_f.F90`
- “supporting” modules and subroutines (used in the interface and user subroutines), eg.
  - `mod_assimilation.F90`
  - `init_pda_f_parse.F90`

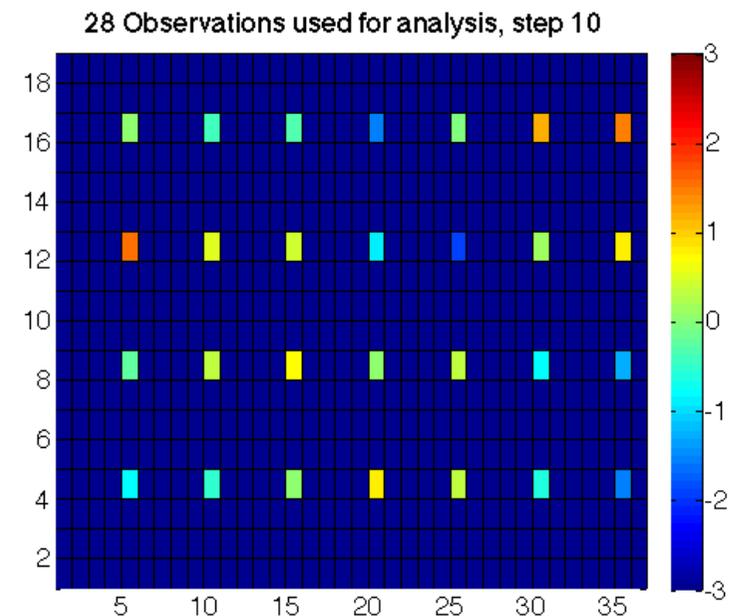
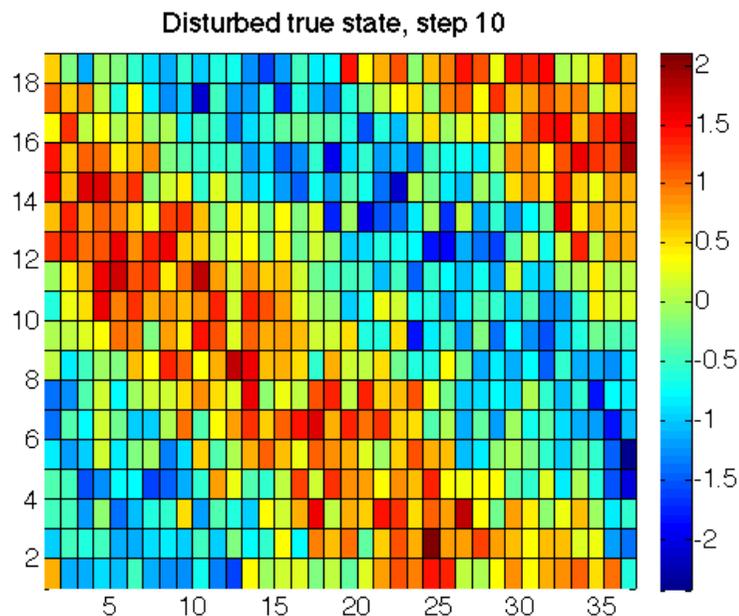
## Running the tutorial model

---

- `cd` to `tutorial/online_2D_serialmodel/model`
- Set environment variable `PDAF_ARCH`  
`export PDAF_ARCH=linux_gfortran_openmpi`
- Run `make`
- Run the model with `./model`
  
- Inputs are read in from `tutorial/inputs_online`
- Outputs are written in  
`tutorial/online_2D_serialmodel/model`  
eg. `true_step10.txt`

# Observations

- Add random error to true state (standard deviation 0.5)
- Select a set of observations at 28 grid points
- File storage (in `inputs_online`):  
text file, full 2D field, -999 marks 'no data' – `obs_step*.txt`  
one file for each time step



## Coupling the model to PDAF: Online mode

---

- Combine model with PDAF into single program
  - modify `Makefile` to build `model_pdaf`
- Add 4 subroutine calls:
  - `init_parallel_pdaf`- add parallelization
  - `init_pdaf` - initialize assimilation
  - `assimilate_pdaf` - perform assimilation
  - `finalize_pdaf` - clean up
- Implement user subroutines, e.g. for
  - observation operator
  - initialization of observation vector
  - transfer between state vector and model fields

<http://pdaf.awi.de/trac/wiki/OverviewOfUserRoutinesWithDefaultNames>

## Online coupling: Parallelization

---

- Online coupling avoids writing to disk to exchange state vectors between the model and PDAF
- Add MPI to the model to run several model instances in parallel
- Run the parallel version with

```
mpirun -np <n> ./model_pdaf ...
```

- *Alternative:* PDAF's "flexible" approach:  
<http://pdaf.awi.de/ModifyModelForEnsembleIntegration>
  - `cd to tutorial/online_2D_serialmodel.noMPI/model`

# Files to copy from pdaf to model

`init_parallel_pdaf.F90`

`mod_parallel_pdaf.F90`

`parser_mpi.F90`

parallelization

`finalize_pdaf.F90`

clean up

PDAF interface subroutine -  
called from the model

helper module/subroutine for  
the interface

PDAF user subroutine - called  
from PDAF library

`init_pdaf.F90`

`mod_assimilation.F90`

`init_pdaf_info.F90`

`init_pdaf_parse.F90`

`init_ens.F90`

initialization

`next_observation_pdaf.F90`

`distribute_state_pdaf.F90`

ensemble forecast

`prepoststep_ens_pdaf.F90`

post step

... (continued on next slide)

# Files to copy from pdaf to model

... (continued from previous slide)

**assimilate\_pdaf.F90**

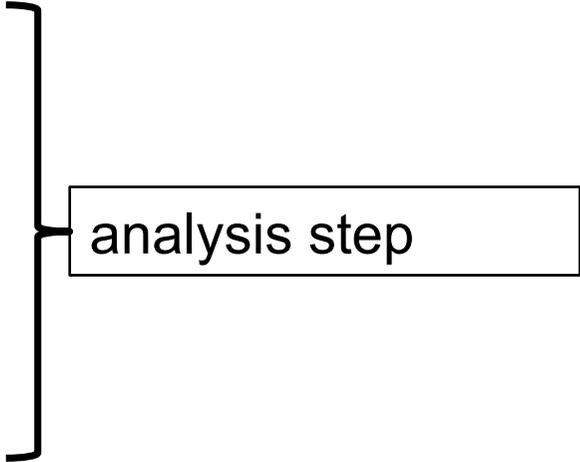
collect\_state\_pdaf.F90

init\_dim\_obs\_pdaf.F90

obs\_op\_pdaf.F90

init\_obs\_pdaf.F90

prodrinva\_pdaf.F90



analysis step

- Each file contains a short summary what the subroutine does

## Files to be adapted in model

- `main.F90` - add calls to PDAF interface
- `integrate.F90` - add calls to PDAF interface
- Makefile** - add linking to PDAF library, PDAF interface and user subroutines

- *Reference solutions for the modified files are in `model_coupled_to_pdaf`*

- When complete, run `make` again

- Then run

```
mpirun -np 9 ./model_pdaf -dim_ens 9
```

- Outputs are written to

```
ens_<i>_step<j>_for.txt
```

```
ens_<i>_step<j>_ana.txt
```

This runs a filter without localization with ensemble size 9

# Plotting

---

- When your coupling is working, lookt at the results
- With Matlab/Octave you can use

```
load ens_01_step02_for.txt  
pcolor(ens_01_step02_for)
```

- Or use the Python scripts

```
./plot_file.py ens_<i>_step<j>_for.txt  
./plot_ens.py <i> <j>
```

## More PDAF experiments

---

- Find PDAF command line parameters in

```
./pdaf/init_pdaf_parse.F90
```

- Try for example

```
mpirun -np 4 ./model_pdaf -dim_ens 4
```

(this runs a filter (ESTKF) without localization with ensemble size 4; it gives a worse result than ensemble size 9)

```
mpirun -np 9 ./model_pdaf -dim_ens 9 -filtertype 7
```

(this runs a filter (LESTKF) with localization and localization radius 0, i.e. correcting only at observed grid points)

```
mpirun -np 9 ./model_pdaf -dim_ens 9 -filtertype 7  
-local_range 5
```

(this runs a filter (LESTKF) with localization and localization radius of 5 grid points)

## Feedback, Questions, more code, ...

---

Full PDAF package contains

- more tutorial code, more filters, and the fully implemented Lorenz-96 model and MITgcm model binding

Web site provides an extensive tutorial for self-study

For further questions

- Contact us at [pdaf@awi.de](mailto:pdaf@awi.de)
- Poster A.14, Friday 14:00–15:45 (L. Nerger)



[pdaf@awi.de](mailto:pdaf@awi.de)

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