

SANGOMA: Stochastic Assimilation for the Next Generation Ocean Model Applications

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Scale separation

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1 Problem definition

2 Solution

3 Sangoma relevance

Additive processes

Different independent processes, each one with a different covariance matrix \mathbf{B}_s

\Rightarrow the covariance of the total field is $\mathbf{B} = \sum_s \mathbf{B}_s$

Kalman gain:

$$\bar{\mathbf{K}} = \mathbf{B}\mathbf{H}^T(\mathbf{H}\mathbf{B}\mathbf{H}^T + \mathbf{R})^{-1} = \sum_s \mathbf{B}_s \mathbf{H}^T \left(\sum_i \mathbf{H}\mathbf{B}_i\mathbf{H}^T + \mathbf{R} \right)^{-1} = \sum_s \bar{\mathbf{K}}_s \quad (1)$$

and

$$\bar{\mathbf{K}}_s = \mathbf{B}_s \mathbf{H}^T \left(\sum_i \mathbf{H}\mathbf{B}_i\mathbf{H}^T + \mathbf{R} \right)^{-1}. \quad (2)$$

Additive processes

$$\bar{\mathbf{K}} = \sum_s \bar{\mathbf{K}}_s; \quad \bar{\mathbf{K}}_s = \mathbf{B}_s \mathbf{H}^T \left(\sum_i \mathbf{H} \mathbf{B}_i \mathbf{H}^T + \mathbf{R} \right)^{-1}. \quad (3)$$

If $\left(\sum_i \mathbf{H} \mathbf{B}_i \mathbf{H}^T + \mathbf{R} \right)^{-1}$ can be "inverted": No need to go further

Otherwise ...

Special individual covariance matrices

- Spline methods (covariances not explicitly calculated)
- 3DVar-NMC implementations
- Reduced rank approaches (EnKF, SEEK,...)
- OI with localization
- ...

using a covariance specification and Kalman gain

$$\mathbf{K}_s = \mathbf{B}_s \mathbf{H}^T (\mathbf{H} \mathbf{B}_s \mathbf{H}^T + \mathbf{R})^{-1}. \quad (4)$$

Special individual covariance matrices

Common property: Using a single covariance matrix specification does lead to efficient algorithms

⇒ K_s can easily be applied to a (big) data array

So what ?

$$\mathbf{K}_s = \mathbf{B}_s \mathbf{H}^T (\mathbf{H} \mathbf{B}_s \mathbf{H}^T + \mathbf{R})^{-1}. \quad (5)$$

can easily be applied to a (big) data array but

$$\bar{\mathbf{K}} = \sum_s \bar{\mathbf{K}}_s; \quad \bar{\mathbf{K}}_s = \mathbf{B}_s \mathbf{H}^T \left(\sum_i \mathbf{H} \mathbf{B}_i \mathbf{H}^T + \mathbf{R} \right)^{-1}. \quad (6)$$

can not.

Two processes (scales)

Matrix identity

$$\bar{\mathbf{K}}_1 = \mathbf{K}_1 - \mathbf{K}_1 \mathbf{H} (\mathbf{I} - \mathbf{K}_2 \mathbf{H} \mathbf{K}_1 \mathbf{H})^{-1} \mathbf{K}_2 (\mathbf{I} - \mathbf{H} \mathbf{K}_1) \quad (7)$$

$$= \mathbf{K}_1 - \mathbf{K}_1 \mathbf{H} \mathbf{K}_2 (\mathbf{I} - \mathbf{H} \mathbf{K}_1 \mathbf{H} \mathbf{K}_2)^{-1} (\mathbf{I} - \mathbf{H} \mathbf{K}_1) \quad \text{Method P1a} \quad (8)$$

$$= (\mathbf{I} - \mathbf{K}_1 \mathbf{H} \mathbf{K}_2 \mathbf{H})^{-1} \mathbf{K}_1 (\mathbf{I} - \mathbf{H} \mathbf{K}_2) \quad (9)$$

$$= \mathbf{K}_1 (\mathbf{I} - \mathbf{H} \mathbf{K}_2 \mathbf{H} \mathbf{K}_1)^{-1} (\mathbf{I} - \mathbf{H} \mathbf{K}_2) \quad \text{Method P1b} \quad (10)$$

but conservation of difficulty because of $(\mathbf{I} - \mathbf{H} \mathbf{K}_2 \mathbf{H} \mathbf{K}_1)^{-1}$?

Lucky fact

$\mathbf{H}\mathbf{K}_1$ and $\mathbf{H}\mathbf{K}_2$ are so called "hat" matrices, "smaller than one".

For Ξ "smaller than one"

$$(\mathbf{I} - \Xi)^{-1} = (\mathbf{I} + \Xi(\mathbf{I} + \Xi(\mathbf{I} + \Xi(\dots)\dots))). \quad (11)$$

When applied to a vector \mathbf{x} , this immediately provides the algorithm to calculate $\mathbf{y} = (\mathbf{I} - \Xi)^{-1}\mathbf{x}$ as follows

$$\mathbf{y} \leftarrow \mathbf{x}$$

Loop

$$\mathbf{y} \leftarrow \mathbf{x} + \Xi\mathbf{y} \quad (12)$$

End loop

Algorithm 1 : Approximate matrix inversion



Successive applications of existing tools

Even if iterations are stopped, some of the formulas yield identical results and there remain

$$\bar{\mathbf{K}}_1 = \mathbf{K}_1 - \mathbf{K}_1 \mathbf{H} \mathbf{K}_2 (\mathbf{I} - \mathbf{H} \mathbf{K}_1 \mathbf{H} \mathbf{K}_2)^{-1} (\mathbf{I} - \mathbf{H} \mathbf{K}_1) \quad \text{Method P1a} \quad (13)$$

$$= \mathbf{K}_1 (\mathbf{I} - \mathbf{H} \mathbf{K}_2 \mathbf{H} \mathbf{K}_1)^{-1} (\mathbf{I} - \mathbf{H} \mathbf{K}_2) \quad \text{Method P1b} \quad (14)$$

and also

$$\bar{\mathbf{K}}_2 = \mathbf{K}_2 - \mathbf{K}_2 \mathbf{H} \mathbf{K}_1 (\mathbf{I} - \mathbf{H} \mathbf{K}_2 \mathbf{H} \mathbf{K}_1)^{-1} (\mathbf{I} - \mathbf{H} \mathbf{K}_2) \quad \text{Method P2a} \quad (15)$$

$$= \mathbf{K}_2 (\mathbf{I} - \mathbf{H} \mathbf{K}_1 \mathbf{H} \mathbf{K}_2)^{-1} (\mathbf{I} - \mathbf{H} \mathbf{K}_1) \quad \text{Method P2b} \quad (16)$$

But which to chose ?

Synthetic test case

Controlled experiment leads to very clear guidelines:

- Label the process with the highest signal-to-noise ratio as 1 and the other process as 2 and apply $P1a + P2b$ if we want to achieve the best analysis.
- If both processes have a similar signal-to-noise ratio, we should label the larger scale process as process 1 so that the same formulas are still the best.
- For individual processes, $P1a$ should be used for the iterated versions, whereas K_1 is the best choice for a simple analysis for process 1; for process 2 version $P2b$ and $K_2(I - HK_1)$ are indicated respectively for the iterated approach or a simple approach.
- For iterative methods, if scales are well separated or at least one of the processes has a small signal-to-noise ratio, only very few iterations are needed.



Implementation

$$\phi \leftarrow \mathbf{K}_1 \mathbf{d}$$

$\mathbf{w}_1 \leftarrow \mathbf{H}\phi$ (Use only analysis at data points without clouds)

$\mathbf{w}_1 \leftarrow (\mathbf{d} - \mathbf{w}_1)$ (Residuals)

Calibrate or check OI parameters at this point using \mathbf{w}_1

$\mathbf{w}_2 \leftarrow \mathbf{w}_1$ (Start of iterative matrix inversion)

Loop n times

$\mathbf{w}_2 \leftarrow \mathbf{H}\mathbf{K}_2\mathbf{w}_2$ (apply tool 2 and retrieve solution at data points)

$\mathbf{w}_2 \leftarrow \mathbf{H}\mathbf{K}_1\mathbf{w}_2$ (apply tool 1 and retrieve solution at data points)

$\mathbf{w}_2 \leftarrow \mathbf{w}_1 + \mathbf{w}_2$

End loop

$$\omega_1 \leftarrow \mathbf{K}_2 \mathbf{w}_2$$

ω_1 now contains scale 2 field $\bar{\mathbf{K}}_2 \mathbf{d}$; can be saved for other uses

$$\phi \leftarrow \phi + \omega_1$$
 (Add it to $\mathbf{K}_1 \mathbf{d}$)

$$\mathbf{w}_1 \leftarrow \mathbf{H}\omega_1$$

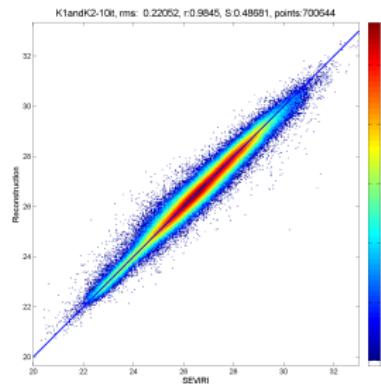
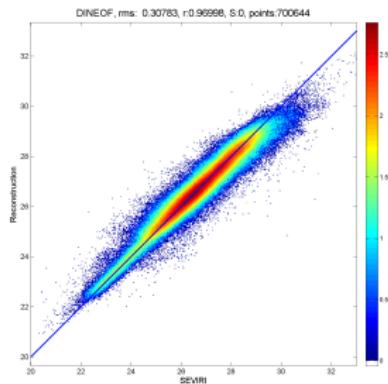
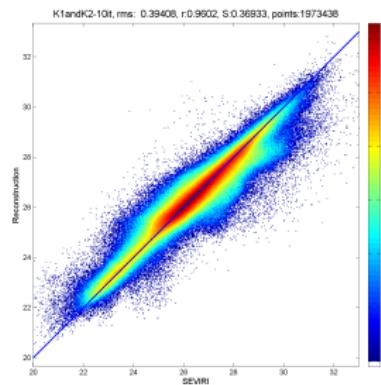
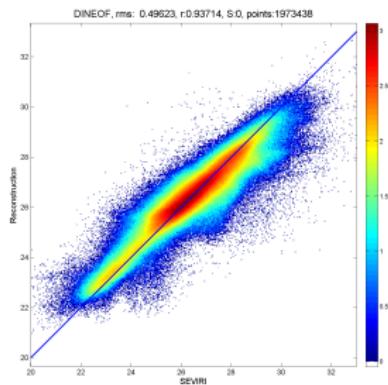
$$\omega_1 \leftarrow \mathbf{K}_1 \mathbf{w}_1$$
 (Calculates $\mathbf{K}_1 \mathbf{H} \bar{\mathbf{K}}_2 \mathbf{d}$)

$$\phi \leftarrow \phi - \omega_1$$
 (Final analysis)

If scale 1 solution is requested, subtract saved scale 2 solution from ϕ and save it

DINEOF+OI, also outline for 3DVar-NMC

Example



Relevance for SANGOMA

- Possibly combine different covariance specifications (Reduced rank+classical OI)
- Possibility to include correlated observational errors into tools not allowing for them (treat R as a covariance matrix of a process (noise)) and isolate it
- Bias analysis

Bias analysis

- Process 1: model field x
- Process 2: bias b

and we want the unbiased field $u = x - b$.
with property

$$\bar{K}_1 = K_1 (I - H \bar{K}_2) \quad (17)$$

and similarly

$$\bar{K}_2 = K_2 (I - H \bar{K}_1) \quad (18)$$

Analysis

$$\mathbf{x}^a = \mathbf{x}^f + \bar{\mathbf{K}}_1 (\mathbf{y} - \mathbf{H}\mathbf{u}^f) \quad (19)$$

$$\mathbf{b}^a = \mathbf{b}^f - \bar{\mathbf{K}}_2 (\mathbf{y} - \mathbf{H}\mathbf{u}^f) \quad (20)$$

and also

$$\mathbf{u}^a = \mathbf{u}^f + (\bar{\mathbf{K}}_1 + \bar{\mathbf{K}}_2) (\mathbf{y} - \mathbf{H}\mathbf{u}^f) \quad (21)$$

A little formula shuffling

$$\mathbf{b}^a = \mathbf{b}^f - \bar{\mathbf{K}}_2 (\mathbf{y} - \mathbf{H}\mathbf{u}^f) \quad (22)$$

$$\mathbf{x}^a - \mathbf{b}^a = \mathbf{x}^f - \mathbf{b}^a + \mathbf{K}_1 (\mathbf{y} - \mathbf{H}(\mathbf{x}^f - \mathbf{b}^a)) \quad (23)$$

This is equivalent to the unsimplified Dee's approach:

Dee's method

Their method uses bias variables β in a reduced space from which bias on the model grid b is obtained by a linear relationship $b = L\beta$ (for bias parameters given in model space $L = I$). The method uses an augmented state vector $[\beta^T x^T]^T$ and the solution of the minimization as

$$\beta^a = \beta^f - K_\beta [y - H(x^f - b^f)] \quad (24)$$

$$u^a = (x^f - b^a) + K_x [y - H(x^f - b^a)] \quad (25)$$

with Kalman gain matrices

$$K_x = P_x H^T (H P_x H^T + R)^{-1} \quad (26)$$

$$K_\beta = P_\beta L^T H^T [H (P_x + L P_\beta L^T) H^T + R]^{-1} \quad (27)$$

Is our multiscale version if $P_x = B_1$ and $P_\beta = B_2$.



Dee's method in practice

In Dee's approach then strong simplification are done because of the impossibility to afford inversion involved in the update of β .

Very strong assumption: $L = I$ and

$$P_\beta = \gamma P_x \quad (28)$$

with a tuning parameter γ .

Justification: seems to work AND is practicable see expression of

$$K_\beta = P_\beta L^T H^T \left[H \left(P_x + L P_\beta L^T \right) H^T + R \right]^{-1} \quad (29)$$

Back to our idea

With iterations stopped early $\bar{\mathbf{K}}_2 \sim \mathbf{K}_2 (\mathbf{I} - \mathbf{H}\mathbf{K}_1)$

$$\mathbf{r} = \mathbf{y} - \mathbf{H}\mathbf{u}^f \quad (30)$$

$$\tilde{\mathbf{r}} = \mathbf{r} - \mathbf{H}\mathbf{K}_1\mathbf{r} \quad (31)$$

$$\mathbf{b}^a = \mathbf{b}^f - \mathbf{K}_2\tilde{\mathbf{r}} \quad (32)$$

$$\mathbf{u}^a = \mathbf{x}^f - \mathbf{b}^a + \mathbf{K}_1 \left(\mathbf{y} - \mathbf{H} \left(\mathbf{x}^f - \mathbf{b}^a \right) \right) \quad (33)$$

Yet another ordering of operations

$$\mathbf{u}^f = \mathbf{x}^f - \mathbf{b}^f \quad (34)$$

$$\tilde{\mathbf{y}} = \mathbf{y} - \mathbf{H}\mathbf{K}_2\mathbf{y} \quad (35)$$

$$\tilde{\mathbf{u}}^f = \mathbf{u}^f - \mathbf{K}_2\mathbf{H}\mathbf{u}^f \quad (36)$$

$$\mathbf{x}^a - \mathbf{b}^f = \mathbf{u}^f + \mathbf{K}_1 \left(\tilde{\mathbf{y}} - \mathbf{H}\tilde{\mathbf{u}}^f \right) \quad (37)$$

$$\mathbf{b}^a = \mathbf{b}^f - \mathbf{K}_2 \left(\mathbf{y} - \mathbf{H}(\mathbf{x}^a - \mathbf{b}^f) \right) \quad (38)$$

$$\mathbf{u}^a = \mathbf{x}^a - \mathbf{b}^f - (\mathbf{b}^a - \mathbf{b}^f) \quad (39)$$

Test case

Small benchmark case with bias in equations.

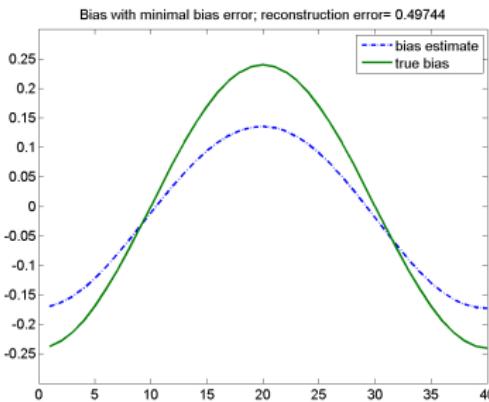
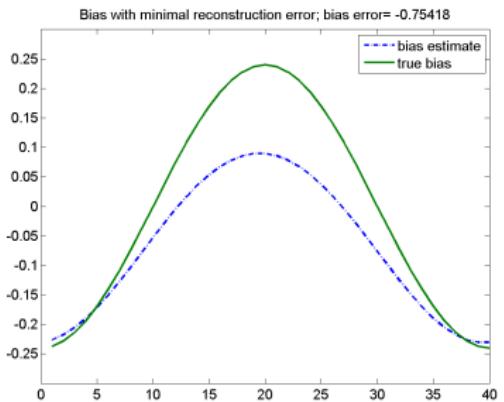
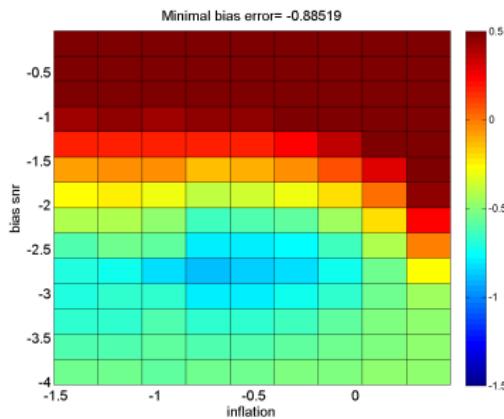
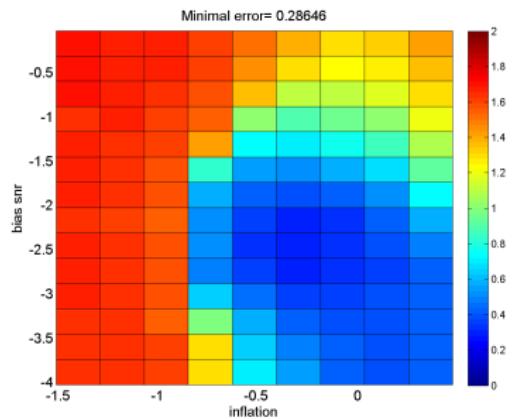
Lorenz-96 with added bias. $N = 40$ members, Model start from random initialization and then 10000 step. From there assimilation every step and we look after 20000 assimilation cycles. Initial members were constructed by adding random Gaussian noise of variance 1.3 in each location.

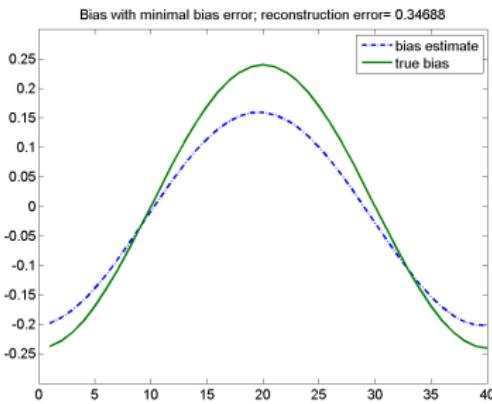
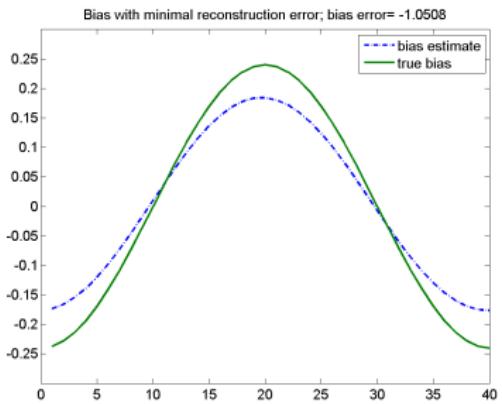
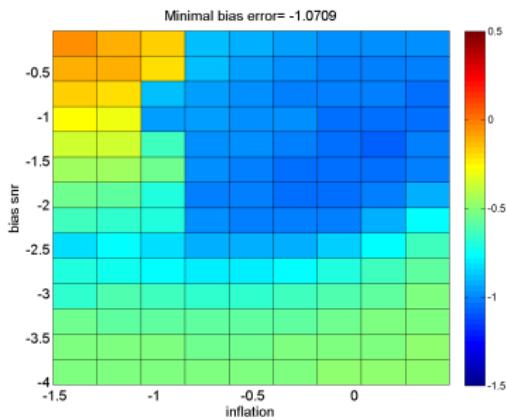
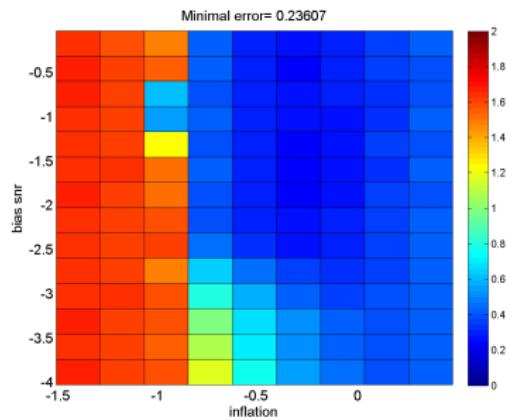
Additive bias $A \sin\left(\frac{2\pi(i-1)}{N}\right)$ all variables observed with $= \sigma^2 \mathbf{I}$, $\sigma^2 = 0.09$. Metric is rms of analysis averaged over time (over 30000 time steps).

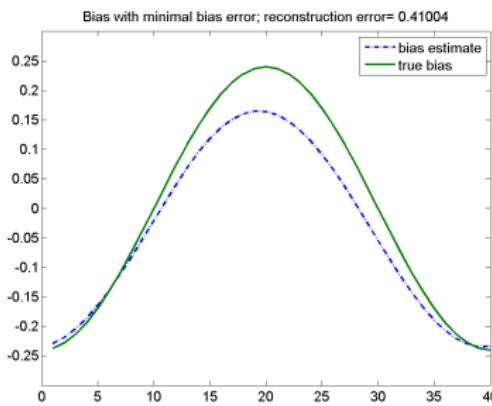
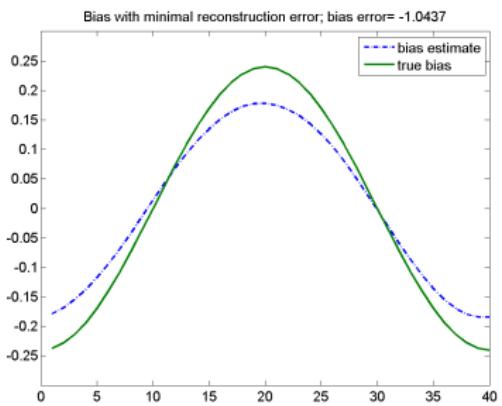
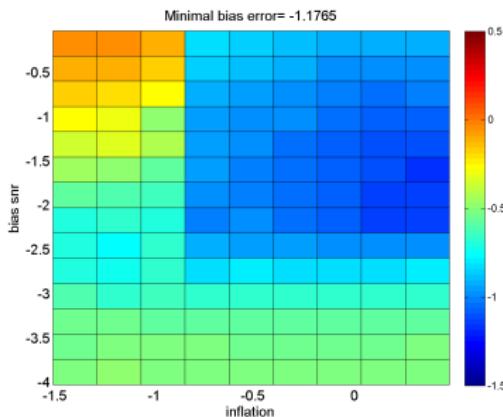
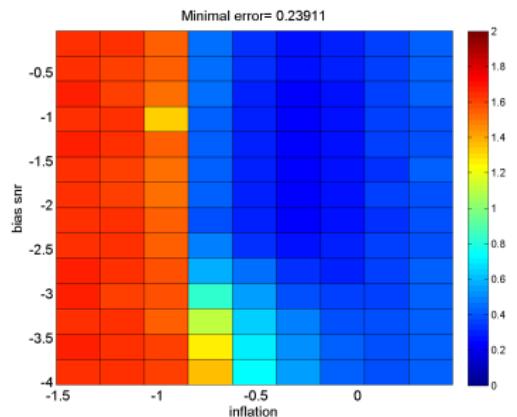
Different strategies

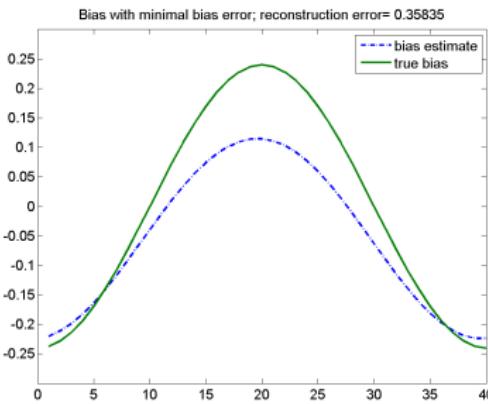
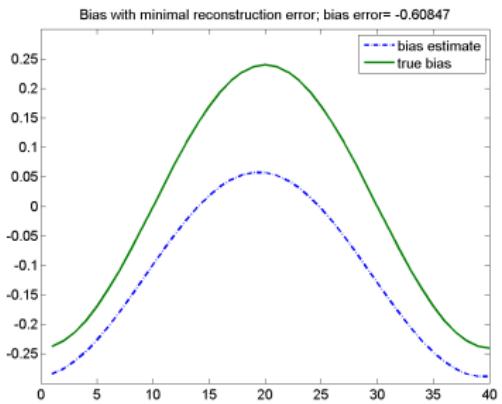
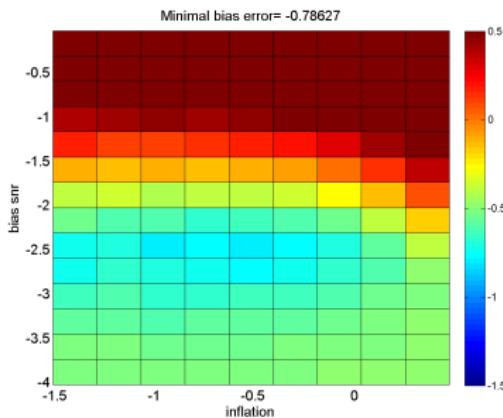
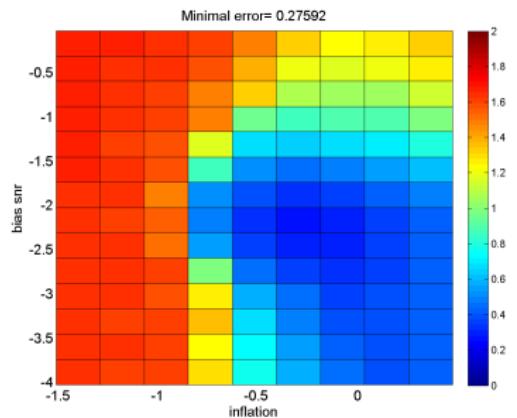
Bias forecast model is persistence in time. Spatial covariance specified by OI with empirical covariance function

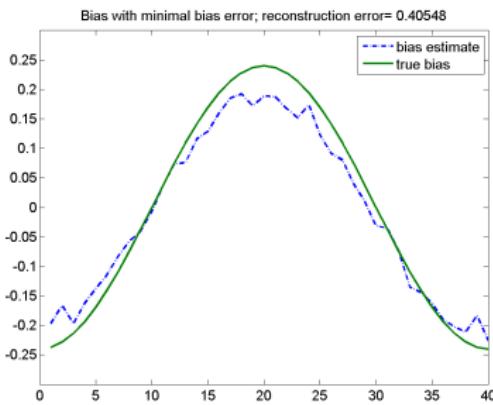
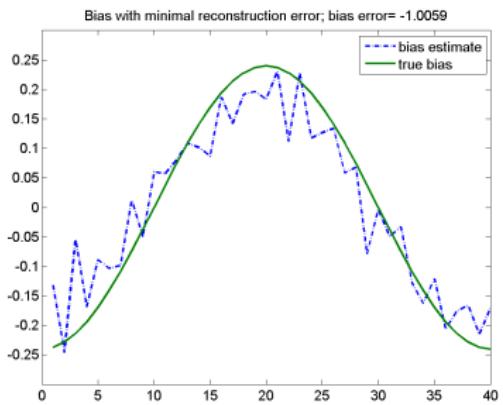
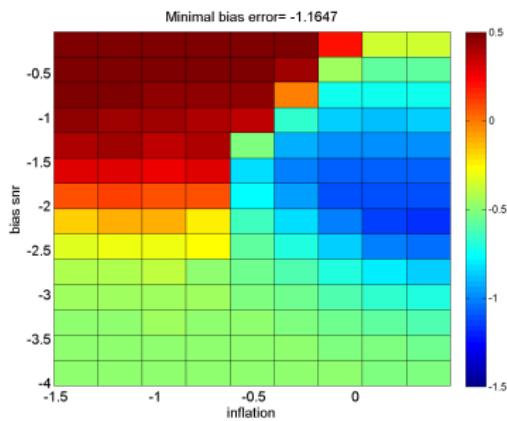
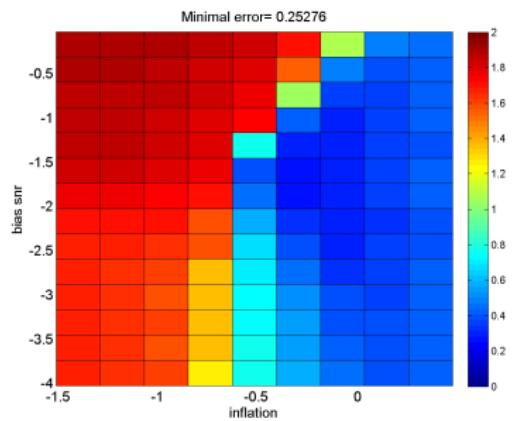
- EnKF for model.
- EnKS for model and constant bias over the smoother window (can be detangled as in EnKS approaches)
- First bias and then model estimate or inverse
- Iterations or not in scale separation
- Full observation or only partial observation
- Different spatial covariance model for bias

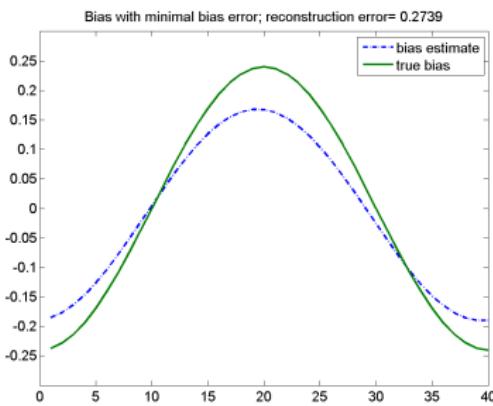
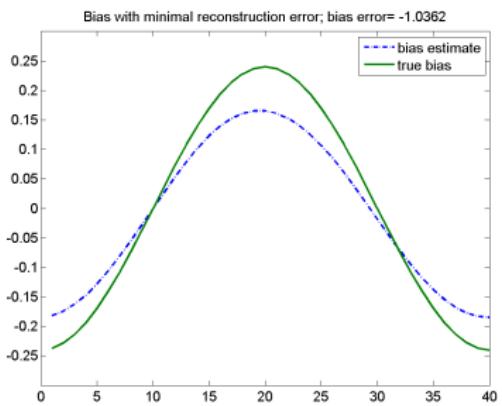
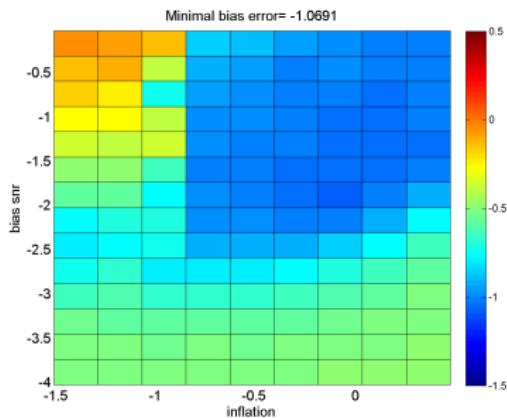
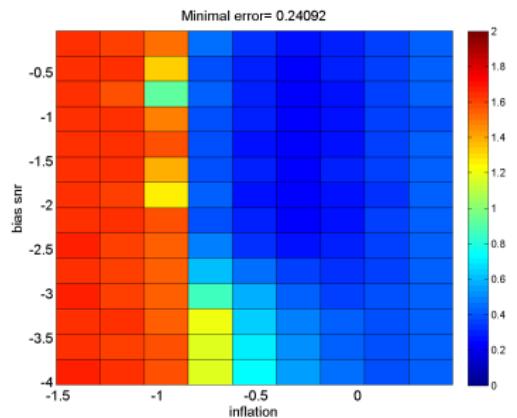












Discussion

- ➊ Results with bias correction in the middle between results with unbiased model and result with biased model uncorrected for bias.
- ➋ Only small differences between EnKF and EnKS? In theory bias at the end is anyway the best estimate ?
- ➌ Iterations on scale separation does (slightly) improve results
- ➍ Method seems more robust than Dee's (in particular with partial observations)
- ➎ Worth continuing ?

References

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Dee, D. P.: Bias and data assimilation, *Quarterly Journal of the Royal Meteorological Society*, 131, 33233343, doi:10.1256/qj.05.137, <http://dx.doi.org/10.1256/qj.05.137>, 2005.