Abstract
Kalman filtering (KF) is a popular form of data assimilation, especially in real-time applications. It combines observations with an equation that describes dynamic evolution to produce an estimate of the present time state of a system. Although KF does not use future information in producing an estimate of the state vector, later reanalysis of the archival data set can produce an improved estimate, in which all data, past, present and future, contribute. We examine the case in which the reanalysis is performed using generalized least squares (GLS), and establish the relationship between the real-time Kalman estimate and the GLS reanalysis. We show that the KF solution at a given time is equal to the GLS solution that one would obtain if data excluded future times. Furthermore, we show that the recursive procedure in KF is exactly equivalent to the solution of the GLS problem via Thomas’ algorithm for solving the block-tridiagonal matrix that arises in the reanalysis problem. This connection suggests that GLS reanalysis is better considered the final step of a single process, rather than a “different method” arbitrarily being applied, post factor. The connection also allows the concept of resolution, so important in other areas of inverse theory, to be applied to KF formulations.

Introduction
In this paper, we compare two data assimilation methods that are routinely applied to monitor time-dependent linear systems, one based on Generalized Least Squares (GLS) and the other on Kalman Filtering (KF). The purpose of the analysis is to enumerate the similarities and differences between the methods, and to provide pathways for applying GLS concepts, and especially resolution analysis, to KF.

At any time, $t_i$, $1 \leq i \leq K$, a linear system is described by a state vector (model parameter vector), $m^{(i)}$, say of length, $M$. This state vector evolves away from the initial condition

$$m^{(1)} = m^{(1)}_A$$

according to the dynamical equation:

$$m^{(i)} = Dm^{(i-1)} + s^{(i-1)}$$

Here, $D$ is the dynamics matrix and $s$ is the source vector. Neither the initial conditions nor the source is known exactly, but rather have uncertainty described by their respective covariance matrices, $C_A$ and $C_S$.

This formulation well-approximates the behavior of systems described by linear partial differential equations that are first order in time. For example, let $m_n^{(i)} = m(x_n, t_i)$ be temperature at time, $t_i = i\Delta t$, and position, $x_n = n\Delta x$, where $\Delta t$ and $\Delta x$ are small increments, and suppose that $m(x, t)$ satisfies the thermal diffusion equation, $\partial m / \partial t = c \partial^2 m / \partial x^2 + q$ (with zero boundary conditions). This partial differential equation can be approximated by:
\[
\frac{\mathbf{m}^{(i)} - \mathbf{m}^{(i-1)}}{\Delta t} = \frac{c}{(\Delta x)^2} \Delta_2 \mathbf{m}^{(i-1)} + \mathbf{q}^{(i-1)} \quad \text{or} \quad \mathbf{m}^{(i)} = \mathbf{D} \mathbf{m}^{(i-1)} + \mathbf{s}^{(i-1)}
\]

\[
\mathbf{D} = \left( \frac{c\Delta t}{(\Delta x)^2} \Delta_2 + \mathbf{I} \right) \mathbf{m}^{(i-1)} \quad \text{and} \quad \mathbf{s}^{(i-1)} = \Delta t \mathbf{q}^{(i-1)}
\]

which has the form of the dynamical equation. Here, the choices:

\[
\Delta_2 = \begin{bmatrix}
1 & 0 & 1 & 0 & \cdots \\
1 & -2 & 1 & 0 & \cdots \\
1 & -2 & 1 & & \cdots \\
1 & & & & \cdots
\end{bmatrix}
\]

and

\[
\mathbf{q}^{(i-1)} = \begin{bmatrix}
q_2^{(i-1)} \\
\vdots \\
q_{M-1}^{(i-1)} \\
0
\end{bmatrix}
\]

encode both the differential equation and the boundary conditions. The matrix, \(\mathbf{D}\), is sparse in this example, as well as in many other cases in which it approximates a differential operator.

The data equation expresses the relationship between the state vector and the observables:

\[
\mathbf{G}^{(i)} \mathbf{m}^{(i)} = \mathbf{d}^{(i)} \quad \text{with covariance} \quad \mathbf{C}_d
\]

In the simplest case, the observations may be of selected elements of the state vector, itself, in which case, each row of \(\mathbf{G}\) is zero, except for a single element, say in column, \(k\), that is unity. Here, \(k(i, n)\) is a function that associates \(d_n^{(i)}\) with \(m_k^{(i)}\). Other more complicated relationships are possible. For instance, in tomography, the data is a line integral through \(m(x, y, t)\) (with \(y\) another spatial dimension). The data kernel, \(\mathbf{G}\), is sparse in these two cases. In other cases, it may not be sparse.

The data assimilation problem is to estimate the state vector, \(\mathbf{m}^{(i)}\), \(1 < i \leq K\), using the dynamical equation supplemented with the data equation, together with the initial condition. One possible approach is based on Generalized Least Squares (GLS); another upon Kalman Filtering (KF). In this paper we explore the differences between the two and discuss the pros and cons of each.

In order to simplify notation, we concatenate the state vectors for times into an overall vector, \(\mathbf{m} = (\mathbf{m}^{(1)}, \ldots, \mathbf{m}^{(K)})\), and data vectors into an overall vector, \(\mathbf{d} = (\mathbf{d}^{(2)}, \ldots, \mathbf{d}^{(N)})\). By assumption, no observations are made at time, \(i = 1\).

**Generalized Least Squares Applied to the Data Assimilation Problem**

Generalized Least Squares (GLS) is a technique used to estimate \(\mathbf{m}\) when two types of information are available: prior information and data. By prior information, we mean expectations about the properties of the behavior of \(\mathbf{m}\), that are based on past experience or general physical considerations. The dynamical equation and initial condition discussed in the previous section are examples of prior information. By data we mean direct observations, as typified by the data equation discussed in the previous section.

Prior information can be represented by the linear equation, \(\mathbf{Hm} = \mathbf{h}\) (with covariance \(\mathbf{C}_h\)) and observations can be represented by the linear equation, \(\mathbf{Gm} = \mathbf{d}\) (with covariance \(\mathbf{C}_o\)). The Bayesian
principle leads to the optimal solution, which we denote the Generalize Least Squares (GLS) solution, \( \mathbf{m}_G \). It minimizes a combination of the weighted \( L_2 \) error in prior information and the weighted \( L_2 \) error in the data (where the weighting depends upon the covariances).

Several equivalent forms of the GLS solution, \( \mathbf{m}_G \), and its posterior variance, \( \mathbf{C}_m \), are common in the literature. We enumerate a few of the more commonly-used forms here:

Form 1 groups the prior information and data equations into a single equation, \( \mathbf{Fm} = \mathbf{h} \):

\[
\mathbf{m}_G = [\mathbf{F}^T\mathbf{F}]^{-1}\mathbf{F}^T\mathbf{f} \quad \text{with} \quad \mathbf{F} = \begin{bmatrix} \mathbf{C}_h^{-\frac{1}{2}} \mathbf{H} \\ \mathbf{C}_o^{-\frac{1}{2}} \mathbf{G} \end{bmatrix} \quad \text{and} \quad \mathbf{f} = \begin{bmatrix} \mathbf{C}_h^{-\frac{1}{2}} \mathbf{h} \\ \mathbf{C}_o^{-\frac{1}{2}} \mathbf{d} \end{bmatrix}
\]

\[
\mathbf{C}_m = [\mathbf{F}^T\mathbf{F}]^{-1}
\]  

(6)

Form 2 introduces generalized inverses, \( \mathbf{G}^{-g} \) and \( \mathbf{H}^{-g} \):

\[
\mathbf{m}_G = \mathbf{G}^{-g} \mathbf{d} + \mathbf{H}^{-g} \mathbf{h}
\]

with \( \mathbf{G}^{-g} \equiv \mathbf{A}^{-1}\mathbf{G}^T\mathbf{C}_o^{-1} \) and \( \mathbf{H}^{-g} \equiv \mathbf{A}^{-1}\mathbf{H}^T\mathbf{C}_h^{-1} \) and \( \mathbf{A} \equiv \left[ \mathbf{G}^T\mathbf{C}_o^{-1}\mathbf{G} + \mathbf{H}^T\mathbf{C}_h^{-1}\mathbf{H} \right]^{-1} \)

\[
\mathbf{C}_m = \mathbf{A}^{-1}
\]

(7)

Form 3 organizes the solution in terms of the prior state vector, \( \mathbf{m}_A \); that is, the state vector implied by the prior information, acting alone:

\[
\mathbf{m}_G = \mathbf{G}^{-g} \mathbf{d} + \mathbf{P}_G \mathbf{m}_A \quad \text{with} \quad \mathbf{P}_G \equiv (\mathbf{I} - \mathbf{G}^{-g} \mathbf{G})
\]

and with \( \mathbf{m}_A \equiv \begin{cases} \mathbf{H}^{-1} \mathbf{h} & \exists \mathbf{H}^{-1} \\ ([\mathbf{H}^T\mathbf{C}_h^{-1}\mathbf{H}]^{-1} \mathbf{H}^T\mathbf{C}_h^{-1}\mathbf{h}) & \not\exists \mathbf{H}^{-1} \end{cases} \quad \text{and} \quad \mathbf{C}_A = [\mathbf{H}^T\mathbf{C}_h^{-1}\mathbf{H}]^{-1}
\]

\[
\mathbf{C}_m = \mathbf{P}_G \mathbf{C}_A
\]

(8)

The matrix, \( \mathbf{P}_G \), plays the role of a projection matrix. See Appendix A.1 for a deviation of the covariance equation.

Form 4 introduces the deviation, \( \Delta \mathbf{m} \), of the solution from the prior state vector, and the corresponding deviation, \( \Delta \mathbf{d} \), of the data from that predicted by the prior state vector:

\[
\Delta \mathbf{m} = \mathbf{G}^{-g} \Delta \mathbf{d} \quad \text{and} \quad \Delta \mathbf{m} \equiv \mathbf{m}_G - \mathbf{m}_A \quad \text{and} \quad \Delta \mathbf{d} \equiv \mathbf{d} - \mathbf{G} \mathbf{m}_A
\]

(9)

Finally, Form 5: uses as alternate form of the generalized inverse:

\[
\Delta \mathbf{m} = \mathbf{G}'^{-g} \Delta \mathbf{d} \quad \text{with} \quad \mathbf{G}'^{-g} \equiv \mathbf{C}_A \mathbf{G}^T \mathbf{A}'^{-1} \quad \text{and} \quad \mathbf{A}' \equiv [\mathbf{C}_o + \mathbf{G} \mathbf{C}_A \mathbf{G}^T]
\]

(10)

Tarantola and Valette (1982) prove the equality of the two forms using a matrix identity that we denote TV82-A (see Appendix A.2). Because \( \mathbf{A} \) is \( M \times M \) and \( \mathbf{A}' \) is \( N \times N \), the first form is most useful when
$M < N$; the second when $M > N$. However, a decision to use one or the other must also take in consideration the sparsity of the various matrix products.

Form 4 is derived from Form 3 by subtracting $A\mathbf{m}_A$ from both sides of the Gram equation,

$$A(\mathbf{m}_G - \mathbf{m}_A) = \mathbf{a} - A\mathbf{m}_A$$

$$\left[G^TC_o^{-1}G + H^TC_h^{-1}H\right](\mathbf{m}_G - \mathbf{m}_A) = G^TC_o^{-1}(\mathbf{d} - G\mathbf{m}_A) + H^TC_h^{-1}(\mathbf{h} - H\mathbf{m}_A)$$

(11)

and then by requiring that the second term on the right-hand side vanish, which leads to:

$$A\Delta\mathbf{m} = G^TC_o^{-1}\Delta\mathbf{d} \quad \text{with} \quad \Delta\mathbf{d} = \mathbf{d} - G\mathbf{m}_A \quad \text{and} \quad \mathbf{m}_A = [H^TC_h^{-1}H]^{-1}H^TC_h^{-1}\mathbf{h}$$

(12)

That is, $\mathbf{m}_A$ is due to the prior information acting along. The deviatoric manipulation is completely general; alternately the first term could have been made to vanish, in which leads to:

$$A\Delta\mathbf{m} = H^TC_h^{-1}\Delta\mathbf{h} \quad \text{with} \quad \Delta\mathbf{h} = \mathbf{h} - H\mathbf{m}_G \quad \text{and} \quad \mathbf{m}_G = [G^TC_o^{-1}G]^{-1}G^TC_o^{-1}\mathbf{d}$$

(13)

Here, $\mathbf{m}_G$ is due to the data acting alone. Note that deviatoric manipulations of this type never change the form of the matrix, $A$. We will apply this principle later in the paper.

In the subsequent analysis, we will focus on the Gram equations:

$$A \Delta\mathbf{m} = G^TC_o^{-1}\Delta\mathbf{d} \equiv \mathbf{a}$$

(14)

The initial condition and the dynamical equation can be into a single prior information equation of the form, $H\mathbf{m} = \mathbf{h}$:

$$\begin{bmatrix}
I & -D & I \\
-D & I & -D \\
\vdots & \ddots & \ddots \\
-D & I & -D
\end{bmatrix}
\begin{bmatrix}
\mathbf{m}^{(1)} \\
\mathbf{m}^{(2)} \\
\vdots \\
\mathbf{m}^{(K)}
\end{bmatrix} =
\begin{bmatrix}
\mathbf{m}^{(1)}_A \\
\mathbf{s}^{(1)} \\
\vdots \\
\mathbf{s}^{(K-1)}
\end{bmatrix} \quad \text{with covariance} \quad C_h$$

(15)

Here, $C_h \equiv \text{diag}(C_A, C_s, C_s, \ldots C_s)$. Several quantities derived from $H$, and which we will use later, are:

$$H^T = \begin{bmatrix}
I & -D^T & -D^T & \cdots \\
I & -D^T & \cdots & \ddots \\
I & \cdots & -D^T & \ddots \\
I & \cdots & \cdots & -D^T
\end{bmatrix} \quad \text{and} \quad H^{-1} = \begin{bmatrix}
I & D & D & D \\
D & I & D & D \\
D & D & I & D \\
D & D & D & I
\end{bmatrix}$$
The data equation expresses the relationship between the state vector and the observables, and assuming that no data are available for time, \( i = 1 \), as the form:

\[
\begin{bmatrix}
0 & G^{(2)} \\
& \ddots & \ddots \\
& & 0
\end{bmatrix}
\begin{bmatrix}
m^{(1)} \\
m^{(2)} \\
\vdots \\
m^{(M)}
\end{bmatrix}
= \begin{bmatrix}
d^{(2)} \\
d^{(3)} \\
\vdots \\
d^{(N)}
\end{bmatrix}
\]

with covariance \( C_o \)

Here, \( C_o \equiv \text{diag}(C_d, C_d, C_d, \ldots, C_d) \) is a summary covariance matrix. We note that:

\[
G^T C_d^{-1} = \begin{bmatrix}
0 & G^{(2)T} C_d^{-1} G^{(2)T} \\
& \ddots & \ddots \\
& & 0
\end{bmatrix}
\text{ and } G^T C_d^{-1} d = \begin{bmatrix}
0 \\
& \ddots \\
& & 0
\end{bmatrix}
\]

The matrix, \( A \), in the first form of the Gram equation is block-triangular and symmetric:

\[
\begin{bmatrix}
A_1 & B^T \\
B & A_2 & B^T \\
& \ddots & \ddots \\
& & B^T \\
B & A_N
\end{bmatrix}
\begin{bmatrix}
m^{(1)} \\
m^{(2)} \\
\vdots \\
m^{(K)}
\end{bmatrix}
= \begin{bmatrix}
a_1 \\
a_2 \\
\vdots \\
a_K
\end{bmatrix}
\]

with elements:

\[
A_i = \begin{cases}
[D^T C_s^{-1} D + C_A^{-1}] & (i = 1) \\
D^T C_s^{-1} D + C_s^{-1} + G^{(i)} C_d^{-1} G^{(i)T} & (1 < i < K) \\
C_s^{-1} + G^{(K)} C_d^{-1} G^{(K)T} & (i = K)
\end{cases}
\]
\[ \mathbf{B} = -\mathbf{C}_s^{-1} \mathbf{D} \]

The vector, \( \mathbf{a} \), on the right-hand side of the Gram equation, is:

\[
\mathbf{a}_i = \begin{cases} 
-\mathbf{D}^T \mathbf{C}_s^{-1} \mathbf{s}^{(1)} + \mathbf{C}_A^{-1} \mathbf{m}_A^{(1)} & (i = 1) \\
\left[ -\mathbf{D}^T \mathbf{C}_s^{-1} \mathbf{s}^{(i)} + \mathbf{C}_s^{-1} \mathbf{s}^{(i-1)} + \mathbf{G}^{(i)} \mathbf{C}_d^{-1} \mathbf{d}^{(i)} \right] & (i > 1)
\end{cases}
\]

(21)

Here, we define \( \mathbf{s}^{(i)} \) to be zero.

**Recursive Solution Using the Thomas method**

Insight into the behavior of the solution can be gained by applying the Thomas (1949) method (see Appendix A.3). It consists of a forward-in-time pass through the system that recursively calculates two quantities, \( \mathbf{A}_i \) and \( \mathbf{a}_i \):

\[
\mathbf{A}_i^{-1} \equiv \begin{cases} 
\mathbf{A}_1^{-1} & (i = 1) \\
[\mathbf{A}_i - \mathbf{B} \mathbf{A}_{i-1} \mathbf{B}^T]^{-1} & (i > 1)
\end{cases}
\]

\[
\mathbf{a}_i \equiv \begin{cases} 
\mathbf{a}_1 & (i = 1) \\
[\mathbf{a}_i - \mathbf{B} \mathbf{a}_{i-1}] & (i > 1)
\end{cases}
\]

(22)

After the forward recursion, the system is block-bidiagonal with row \( i \) having elements \( \mathbf{A}_i \) and \( \mathbf{B}^T \) (except for the last row, which lacks the \( \mathbf{B}^T \)) and the modified right-hand side is \( \mathbf{a}_i \). The solution, \( \mathbf{m}_G^{(i)} \), is achieved through a backward recursion:

\[
\mathbf{m}_G^{(i)} = \begin{cases} 
\mathbf{A}_K^{-1} \mathbf{a}_K & (i = K) \\
\mathbf{A}_i^{-1} [\mathbf{a}_i - \mathbf{B}^T \mathbf{m}^{(i+1)}] & (i < K)
\end{cases}
\]

(23)

It is evident that information is propagated both forward and backward in time during the solution process. Furthermore, computation time grows no faster than the number of steps, \( K \), in the recursion.

The Thomas method has a disadvantage in the common case where the covariances matrices, \( \mathbf{C}_A \), \( \mathbf{C}_s \) and \( \mathbf{C}_d \) are diagonal and when \( \mathbf{D} \) and \( \mathbf{G}^{(i)} \) are sparse, because although \( \mathbf{F} \) is then also sparse, the matrices, \( \mathbf{A}_i^{-1} \), are in general not sparse, so the effort needed to compute them scales with \( M^2 \). Other direct methods share this limitation, too. Consequently, the overall calculation scales with \( KM^3 \). The conjugate gradient method, applied to the Gram equation, \( \mathbf{F}^T \mathbf{F} \mathbf{m} = \mathbf{F}^T \mathbf{h} \), is usually a better choice. This method requires that the quantity, \( \mathbf{u} = (\mathbf{F}^T \mathbf{F}) \mathbf{v} \), be calculated for an arbitrary vector, \( \mathbf{v} \), and this quantity can be very efficiently calculated as \( \mathbf{u} = \mathbf{F}^T (\mathbf{F} \mathbf{v}) \). In cases in which the dynamical equation approximates a partial differential equation, the number of non-zero elements in the matrix, \( \mathbf{F} \), scale with \( KM \). The conjugate gradient algorithm requires no more than \( KM \) iterations (and often much fewer), each requiring \( KM \) multiplications. Thus, the overall solution time scales with \( K^2 M^2 \). Consequently, the conjugate gradient method has a speed advantage when \( M > K \).
**Present Time Solution**

Suppose that the analysis focuses on the “present time”, $i$, in the sense that only the solution, $m_p^{(i)}$, determined using data up to and including time, $i$, is of interest. One can assemble a sequence of present-time solutions during the forward recursion, using the fact that the $i$th solution can always be considered to be the final one, and no backwards recursion is needed to compute the solution for the final time. However, the forms of the “final” $\hat{A}_i$ and $\hat{a}_i$ differ from that of the previous $A$s in the recursion, so a separate computation is needed:

$$m_p^{(i)} = \hat{A}_i^{-1} \hat{a}_i$$

$$\hat{A}_i' = \hat{A}_i - D^T C_s^{-1} D = C_{st}^{-1} + G(i) C_d^{-1} G(i)^T$$

and

$$\hat{a}_i' = \hat{a}_i + D^T C_s^{-1} s^{(i)}$$

(24)

Consequently, in order to create a sequence of present-time solution, the two matrix inverses – not one – must be calculated at each step in the forward recursion. The present-time solution is the same as the reference solution, $m_b^{(i)}$, defined in the previous section.

**Kalman Filtering**

Kalman Filtering (KF) is a solution method with an algorithm that, like the Thomas present-time solution, is forward-in-time, only. It consists of four steps, the final three of which are iterated.

Step 1 assigns the $i = 1$ solution, $m_K^{(1)}$, its covariance, $C_m^{(1)}$, at the 1st time step:

$$m_K^{(1)} = m_A^{(1)} \text{ and } C_m^{(1)} = C_A$$

(25)

Step 2 propagates the solution and its covariance forward in time using the dynamical equation, and considers it to be prior information.

$$m_A^{(i)} = (D m_K^{(i-1)} + s^{(i-1)}) \text{ and } C_A^{(i)} = D C_m^{(i-1)} D^T + C_s$$

(26)

Step 3 uses GLS to combine the prior information, $m_A^{(i)}$, with covariance $C_A^{(i)}$, and data, $d^{(i)}$, with covariance $C_d$, into a solution, $m_K^{(i)}$, with covariance, $C_m$. Any of the (equivalent) GLS solutions described above can be used in this step.

Step 4, increments $i$ and returns to Step 2, creating a recursion.

Most often, the GLS solution and its variance are written as:

$$m_K^{(i)} = m_A^{(i)} + G_i^{-g} \Delta d^{(i)} \text{ with } \Delta d^{(i)} = d^{(i)} - G(i) m_A^{(i)}$$

with

$$G_i^{-g} = C_A^{(i)} G(i)^T \left[ C_d^{(i)} + G(i) C_A^{(i)} G(i)^T \right]^{-1}$$

and

$$C_A^{(i)} = D C_m^{(i-1)} D^T + C_s$$

and

$$C_m^{(i)} = \left[ I - G_i^{-g} G(i) \right] C_A^{(i)} = P^{(i)} C_A^{(i)} \text{ and } P_g(i) \equiv \left[ I - G_i^{-g} G(i) \right]$$
However, any of the equivalent forms described above can substitute, such as:

\[ m_K^{(i)} = \tilde{A}_i^{-1} \tilde{a}_i \]

\[ c_A^{(i)} = DC_m^{(i-1)}D' + C_s \quad \text{and} \quad \tilde{A}_i^{-1} = [G^{(i)}rC_{dA}^{-1}G^{(i)} + [c_A^{(i)}]^{-1}]^{-1} = C_m \]

\[ \tilde{a}_i = G^{(i)}C_{dA}^{-1}d + [c_A^{(i)}]^{-1}Dm_K^{(i-1)} + [c_A^{(i)}]^{-1}s^{(i-1)} \]

(28)

**Is Kalman Filtering “filtering”?**

A standard Infinite Impulse Response (IIR) filter has the form \( v * m = u * z \), where \( z \) is the input timeseries, \( m \) is the “output” timeseries, \( u \) and \( v \) are filters (with \( v_1 = 1 \)) and \(*\) signifies convolution. Key to this formulation is that the filter coefficients are constants; that is, they are not a function of time.

If the generalized inverse in KF was time-independent, so that \( G_i^{-g} = G^{-g} \), then KF could be put into form of an IIR filter:

\[
\begin{bmatrix}
[I] \\
[G^{-g}Gd - I]
\end{bmatrix}
\begin{bmatrix}
m^{(i-1)} \\
m^{(i)} \\
\vdots
\end{bmatrix} = \begin{bmatrix}
G^{-g} & -G^{-g}G
\end{bmatrix}
\begin{bmatrix}
d^{(i-2)} \\
s^{(i-1)} \\
\vdots
\end{bmatrix}
\]

(29)

because the convolution reproduces the KF solution:

\[
m^{(i)} = m^{(i-1)} + G^{-g} \left( d^{(i)} - G(Dm^{(i-1)} + s^{(i-1)}) \right)
\]

(30)

So, from this point of view, the KF has a \( v \) of length 2 (each element of which is a matrix), and a \( u \) of length 1 (each element of which is a row vector of two matrices). However, this formulation does not really correspond to a standard IIR filter, because the filter coefficients, which depend upon the generalized inverse, \( G_i^{-g} \), depend upon time, \( i \). Hence, the word “filter”, though generally indicative of the process, oversimplifies the actual operation being performed. It is not filtering in the strict sense.

**The Present-Time Thomas and Kalman Filtering Solutions are Equal**

We will now demonstrate that the present-time Thomas solution, \( m_p^{(i)} \), and the Kalman filtering solution, \( m_K^{(i)} \), are equal. We will make use of an identity, abbreviated TV82-B, that is due to Tarantola and Valette (1982), which shows that for invertible symmetric matrices \( C_1 \) and \( C_2 \) and arbitrary matrix, \( M \):

\[
C_2 - C_2M^T[M^2M^T + C_1]^{-1}MC_2 = [M^T C_1^{-1}M + C_2^{-1}]^{-1}
\]

(31)
Thus, for instance, when $C_1^{-1} = C_2^{(i)}$, $C_2^{-1} = C_s$, and $M = D^T$:

$$
\left[ C_2^{(i)} \right]^{-1} = \left[ DC_m^{(i-1)} D^T + C_s \right]^{-1} = C_s^{-1} - C_s^{-1} D \left[ D^T C_s^{-1} D + \left[ C_m^{(i-1)} \right]^{-1} \right] D^T C_s^{-1}
$$

(32)

The PT and KF recursions both start with $m_1^{(1)} = m_{PT1}^{(1)} = m_A$ and $C_1^{(1)} = C_{PTA}^{(1)} = C_A$. The $(i = 2)$ case in irregular, and must be examine separately. The KF solution is:

$$
m_2^{(2)} = \tilde{A}_2^{-1} \tilde{a}_2
$$

with $\tilde{A}_2 = G^{(2)} C_d^{-1} G^{(2)} + \left[ C_A^{(2)} \right]^{-1}$ with $C_A^{(2)} = DC_A D^T + C_s$

and $\tilde{a}_2 = G^{(2)} C_d^{-1} d^{(2)} + \left[ C_A^{(2)} \right]^{-1} D m_1^{(1)} + \left[ C_A^{(2)} \right]^{-1} s^{(1)}$

(33)

This can be compared with the present-time Thomas solution:

$$
m_2^{(2)} = \tilde{A}_2^{-1} \tilde{a}_2
$$

with $\tilde{A}_2 = G^{(2)} C_d^{-1} G^{(2)} + [DC_A D^T + C_s]^{-1}$

$$
\tilde{a}_2 = G^{(2)} C_d^{-1} d^{(2)} + C_s^{-1} D \tilde{A}_2^{-1} D^T C_s^{-1} s^{(1)} + C_s^{-1} D \tilde{A}_2^{-1} C_s^{-1} D C_A^{-1} m_A
$$

(34)

Note that we used TV82-B to simplify the expression for $\tilde{A}_2$. By inspection, $\tilde{A}_2 = \tilde{A}_2$. Thus, the two solutions are equal if $\tilde{a}_2 = \tilde{a}_2$. The terms involving $d^{(2)}$ match. The terms involving $s^{(1)}$ would match if it could be shown that:

$$
C_s^{-1} - C_s^{-1} D [D^T C_s^{-1} D + C_A^{-1}]^{-1} D^T C_s^{-1} = [DC_A D^T + C_s]^{-1}
$$

(35)

But this equation is true by TV82-B. The terms $m_A$ also match, because of the matrix identity

$$
[DC_A D^T + C_s]^{-1} D = C_s^{-1} D [D^T C_s^{-1} D + C_A^{-1}]^{-1} D C_A^{-1}
$$

(36)

proved in Appendix A.4. Consequently, the solutions, $m_2^{(2)} = m_2^{(2)}$, and their posterior covariances, $C_{km}^{(2)} = \tilde{A}_2^{-1} = C_{pm}^{(1)} = \tilde{A}_2^{-1}$ are equal. Applying $C_2^{(i)} = \tilde{A}_2^{-1}$ to the Karman recursion, and TV82-B and $\tilde{A}_i = \tilde{A}_i + D^T C_s^{-1} D^T$ to present-time Thomas recursion, leads to:

$$
\tilde{A}_{i+1} = G^{(i+1)} C_d^{-1} G^{(i+1)} + [D \tilde{A}_i^{-1} D^T + C_s]^{-1}
$$

$$
\tilde{A}_{i+1} = G^{(i+1)} C_d^{-1} G^{(i+1)} + C_s^{-1} - C_s^{-1} D [D^T C_s^{-1} D + \tilde{A}_i^{-1}]^{-1} D^T C_s^{-1}
$$
\[ \dot{\mathbf{A}}_i = \mathbf{G}(i+1)\mathbf{C}_d^{-1}\mathbf{G}(i+1)^T + \left[\mathbf{D}\tilde{\mathbf{A}}_i^{-1}\mathbf{D}^T + \mathbf{C}_s\right]^{-1} \]

(37)

Thus, \( \tilde{\mathbf{A}}_{i+1} = \mathbf{A}'_{i+1} \) as long as \( \tilde{\mathbf{A}}_i^{-1} = \mathbf{A}'_i^{-1} \). Because the latter is true for \( i = 2 \), so the formula can be successively applied to show \( \tilde{\mathbf{A}}_{i+1} = \mathbf{A}'_{i+1} \) for all \( i > 2 \). Similarly, the procedure that demonstrated the equality of \( \tilde{\mathbf{a}}_2 \) and \( \mathbf{A}_2 \) can be extended to

\[
\tilde{\mathbf{a}}_{i+1} = \mathbf{G}(i+1)^T\mathbf{C}_d^{-1}\mathbf{d}^{(i+1)} + \left[\mathbf{D}\tilde{\mathbf{A}}_i^{-1}\mathbf{D}^T + \mathbf{C}_s\right]^{-1}\mathbf{s}^{(i)} + \left[\mathbf{D}\tilde{\mathbf{A}}_i^{-1}\mathbf{D}^T + \mathbf{C}_s\right]^{-1}\mathbf{Dm}_K^{(i)}
\]

and

\[
\tilde{\mathbf{a}}_i = \mathbf{C}_s^{-1}\mathbf{s}^{(i)} + \mathbf{G}(i+1)^T\mathbf{C}_d^{-1}\mathbf{d}^{(i+1)} + \mathbf{C}_s^{-1}\mathbf{D}\tilde{\mathbf{A}}^{-1}_i[\mathbf{a}'_i - \mathbf{D}^T\mathbf{C}_s^{-1}\mathbf{s}^{(i)}]
\]

\[= \mathbf{G}(i+1)^T\mathbf{C}_d^{-1}\mathbf{d}^{(i+1)} + \left[\mathbf{C}_s^{-1} - \mathbf{C}_s^{-1}\mathbf{D}\tilde{\mathbf{A}}^{-1}_i\mathbf{D}^T\mathbf{C}_s^{-1}\mathbf{s}^{(i)}\right] + \mathbf{C}_s^{-1}\mathbf{D}\tilde{\mathbf{A}}^{-1}_i\tilde{\mathbf{a}}'_i
\]

\[= \mathbf{G}(i+1)^T\mathbf{C}_d^{-1}\mathbf{d}^{(i+1)} + \left[\mathbf{C}_s + \mathbf{D}\tilde{\mathbf{A}}^{-1}_i\mathbf{D}^T\right]^{-1}\mathbf{s}^{(i)} + \mathbf{C}_s^{-1}\mathbf{D}\tilde{\mathbf{A}}^{-1}_i\tilde{\mathbf{a}}'_i\mathbf{m}_K^{(i)}
\]

(38)

Here we have used TVS2-B and \( \mathbf{m}_K^{(i)} = \tilde{\mathbf{A}}^{-1}_i\tilde{\mathbf{a}}'_i \). The terms ending in \( \mathbf{d}^{(i+1)} \) match. The terms ending in \( \mathbf{s}^{(i)} \) also match, since it has been established previously that \( \tilde{\mathbf{A}}^{-1}_i = \tilde{\mathbf{A}}^{-1}_i \). In order for \( \tilde{\mathbf{a}}_{i+1} \) to equal \( \tilde{\mathbf{a}}_{i+1} \), we must have \( \mathbf{m}_K^{(i)} = \mathbf{m}_K^{(i)} \) and:

\[
\left[\mathbf{D}\tilde{\mathbf{A}}^{-1}_i\mathbf{D}^T + \mathbf{C}_s\right]^{-1}\mathbf{D} = \mathbf{C}_s^{-1}\mathbf{D}[\tilde{\mathbf{A}}^{-1}_i + \mathbf{D}^T\mathbf{C}_s^{-1}\mathbf{D}]\tilde{\mathbf{A}}'_i
\]

(39)

However, this equation has the same form a the identity in (36), where the equality was demonstrated. Starting with \( i = 2 \), we have \( \mathbf{m}_K^{(2)} = \mathbf{m}_K^{(2)} \) and \( \tilde{\mathbf{A}}'_2 = \tilde{\mathbf{A}}_2 \), which implies \( \tilde{\mathbf{A}}'_3 = \tilde{\mathbf{A}}_3 \) and \( \tilde{\mathbf{a}}_3 = \tilde{\mathbf{a}}_3 \), which implies \( \mathbf{m}_K^{(3)} = \mathbf{m}_K^{(3)} \). This process can be iterated indefinitely, establishing that the present-time Thomas and Kalman solutions, and their posterior variance, are equal.

**Comparison between the Present Time Solution and GLS**

The present time solutions at time, \( j \), depends on information available for times, \( (i \leq j) \) but not upon information that subsequently becomes available (that is, for times \( (i > j) \). While this limitation is necessary real-time scenarios, wher. However, the lack of future data leads to solution that is poorer estimate of the true solution, than a GLS solution in which the state vectors at all time are globally adjusted to best-fit all the prior information and data.

An outlier that occurs at, or immediately before, the present moment can cause large error in the present time solution. The global solution is less affected because measurements in the near future may compensate (Fig. 1).
Fig. 1. Hypothetical data assimilation scenario, with $K = 4$, $M = N = 1$, $D = 1$, $s = 0.25$. The initial condition (red box) satisfies $m_a^{(0)} = 0$. The dynamical equation requires that the slope (dotted red lines) be about $1/4$. The data, $d^{(i)}$, are noisy versions of state $m^{(i)}$, which is expected to linearly increase with time with slope, $s$. When the “present time” is $i = 3$, the present time solution (grey), $m_P^{(3)}$, is pulled down by the noisy datum, $d^{(i)}$, leading to a poor fit to the dynamics at that time. The GLS solution, $m_G^{(3)}$, is less affected by the outlier, because the datum at time, $i = 4$, better defines the linear trend, leading to a solution at $i = 3$ that better matched the dynamics.

Having established links between KF and GLS, we are now able to apply several useful inverse theory concepts, and especially resolution (Backus and Gilbert, 1968; Wiggins 1972). In a GLS problem, model resolution refers to the ability of the data assimilation process to reconstruct deviations of the true model from the one predicted by the prior information, alone (Menke, 2014). Data resolution refers to its ability to reconstruct deviations of the data from the one predicted by the prior information (Menke and Creel, 2021). Model resolution is quantified by a $(KM) \times (KM)$ matrix, $R = GG^{-\beta}$, and data resolution by a $(KN) \times (KN)$ matrix, $N = GG^{-\beta}$, that satisfy.

\[
(m - m_A) = R(m_{true} - m_A) \quad \text{and} \quad (d - Gm_A) = N(m_{true} - Gm_A)
\]

Resolution is perfect when $R = N = I$. When $R \neq I$, an element of the reconstructed state vector is a weighted average of all elements of the true state vector. When $N \neq I$, an element of the reconstructed data vector is a weighted average of all elements of the true data vector. The resolution matrices quantify resolution in both time and space. As we will show in the example, below, the model resolution (or data resolution) can be temporally poor, even when it is spatially good.

Another important quantity is the full $(KM) \times (KM)$ posterior covariance matrix, $C_m = [F^T F]^{-1}$. In addition to correlations between elements of the state vector at a given time, it contains the correlations between elements of the state vectors at different times. These coefficients are needed for computing confidence intervals of quantities that depend of the state vectors at two or more times. Although $R$, $N$ and $C_m$ are large matrices, methods are available for efficiently computing selected elements of them using the conjugate gradient method (Menke, 2014).

**Example**

We consider a data assimilation problem based on the heat diffusion equation (3), with $\Delta t = \Delta x = 1$ and $c_0 = 0.4$. The state vector, $m^{(i)}$, represents temperature and is of length $M = 31$. The source Gaussian in space and impulsive in time, according to

\[s_j^{(i)} = \exp \left[ -\frac{s}{2} s_x^{-2} (x_j - \bar{x})^2 \right] \delta_{i1} + n_s^{(j)}\]
with scale length, \( s_x = 5 \), and peak position, \( \bar{x} = \frac{1}{2} M \Delta x \). Here, \( n_s(j) \) is a Normally-distributed random variable with zero mean and variance, \( \sigma_s^2 = 0.05 \). The initial condition is:

\[
\begin{bmatrix} m_A^{(0)} \end{bmatrix}_i = m_0 + n_A^{(j)}
\]

where \( m_0 = 0.1 \) and \( n_A^{(j)} \) is a Normally-distributed random variable with zero mean and variance, \( \sigma_d^2 = 0.05 \). The dynamical equation (2) is iterated for \( K = 61 \) time steps, to provide the “true” state vector. As expected, the solution has a Gaussian shape with a width that increases, and an amplitude that decreased, with time (Fig. 2A). The data are a total of \( N = 10 \) temperature measurement at each time, \( i \geq 2 \), made at randomly-selected positions (without duplications) and perturbed with Normally-distributed random noise with zero mean and variance, \( \sigma_d^2 = 0.07 \).

GLS solutions were computed by both the full Thomas algorithm (Fig. 2B) and by solving the \([F^T F]m = F^T f\) system by the conjugate gradient method (not shown). They were found to be identical to machine precision. Present-time solutions were computed for both the present time Thomas (Fig. 2C) and Kalman Filtering algorithms (Fig. 2D). They were also found to be identical to machine precision.

In general, the both the GLS and present time solutions fit the data well. However, the present time solution matches the true model more poorly than does the GLS solution (Fig. 3). In this numerical experiment, the present time solution is about 10% poorer than the GLS solution, quantified with the root mean squared deviation from the true solution. However, the percentage varies considerably when the underlying parameters are changed.
Fig. 3. Histogram of ratios of the root mean squared error between the present time estimate, $m_P^{(i)}$, $i = 1, \ldots, K$, and the true state and the root mean squared error between GLS estimate, $m_G^{(i)}$, $i = 1, \ldots, K$, and the true state, for 1000 realizations of the exemplary data assimilation problem.

Both the Thomas and Kalman versions of the present time algorithm are well suited for providing ongoing diagnostic information, such as posterior covariance and root mean squared data fitting error (Fig. 4), which can provide quality control in real time applications.

Fig. 4. Kalman Filtering solution of the exemplary data assimilation problem. (A) the data, $d^{(i)}$, $i = 1, \ldots, N$. (B) The estimated state $m_K^{(i)}$, $i = 1, \ldots, K$. (C) The root mean square data prediction as a function of time, $i$. (D) The posterior variance as a function of time, $i$. 
The model resolution matrix, $\mathbf{R}$, (Fig. 5A) for this exemplary problem has a poorly-populated central diagonal, meaning that some elements of the state vector, $m_j^{(i)}$, are well-resolved from their spatial neighbors, $m_{j \pm 1}^{(i)}$, while others are very poorly resolved. The matrix large elements along other diagonals, corresponding to, offset from the main diagonal by $M$ rows, indicating that some elements are not well resolved from their temporal neighbors, $m_j^{(i \pm 1)}$. The data resolution matrix, $\mathbf{N}$, (Fig. 5A) has a well-populated central diagonal, meaning that most elements of the predicted data vector, $d_{j}^{\text{pre}(i)}$, are well-resolved from their spatial neighbors, $d_{j \pm 1}^{(i)}$. Like $\mathbf{R}$, it also has elements along other diagonals, corresponding to, offset from the main diagonal by $M$ rows, indicating that some elements are not well resolved from their temporal neighbors, $d_{j}^{\text{pre}(i \pm 1)}$.

Fig. 5. Central portion of the resolution matrices for the exemplary data assimilation problem. (A) The model resolution matrix, $\mathbf{R}$. (B) The data resolution matrix, $\mathbf{N}$. The portion shown corresponds time slices $20 \leq i \leq 23$.

**Summary**

In this paper, we examine a data processing scenario in which real-time data assimilation is performed using Kalman Filtering, and then reanalysis is performed using generalized least squares (GLS). In this problem, spatial characteristics of the system are described by a state vector (mode parameter vector), and its temporal characteristics by the evolution of the state-vector with time. We explore the relationship between the real-time Kalman Filter estimate and the GLS reanalysis estimate of the state vector. We show that the KF solution at a given time is equal to the GLS solution that one would obtain if it excluded data for future times. Furthermore, we show that the recursive procedure in KF is exactly equivalent to the solution of the GLS problem via Thomas’ algorithm for solving the block-tridiagonal matrix that arises in the reanalysis problem. This connection indicates that GLS reanalysis is better considered the final step of a single process, rather than a “different method” arbitrarily being applied, post factor. Now that this connection between KF and GLS has been identified, the familiar GLS concepts of model and data resolution can be applied to KF. We provide an exemplary problem, based on thermal diffusion. In addition to showcasing our result, the example demonstrates that the state vector and vector of predicted data can be poorly-resolved in time, even when they are well-resolved in space.

**References**


Appendix

A.1. Proof that $C_m = [I - G^{-g}G]C_A \equiv P_G C_A$. Because $d$ and $m_A$ are independent of one another, $m_G = G^{-g}(d - Gm_A) + m_A = G^{-g}d + (I - G^{-g}G)m_A$, the normal rules of error propagation apply. The posterior covariance, $C_m$, is:

$$C_m = G^{-g}C_d G^{-g^T} + (I - G^{-g}G)C_A (I - G^T G^{-g^T}) =$$

$$C_A + G^{-g}C_d G^{-g^T} + G^{-g}G C_A G^T G^{-g^T} - C_A G^T G^{-g^T} - G^{-g}G C_A =$$

$$C_A + G^{-g} [G C_A G^T + C_d] G^{-g^T} - C_A G^T G^{-g^T} - G^{-g}G C_A =$$

$$= C_A + C_A G^T A^{-1} AA^{-1} G C_A - C_A G^T A^{-1} G C_A - C_A G^T A^{-1} G C_A =$$

$$= C_A + C_A G^T A^{-1} G C_A - C_A G^T A^{-1} G C_A - C_A G^T A^{-1} G C_A =$$

$$= C_A - C_A G^T A^{-1} G C_A = C_A - G^{-g} G C_A = [I - G^{-g}G] C_A \equiv P_G C_A$$

Here, we have used the fact that $G^{-g} = C_A G^T A^{-1}$, with $A = [G C_A G^T + C_d]$. Although the matrix, $P_G \equiv [I - G^{-g}G]$, has the form of a projection operator, it is a function of $C_A$, and has deceptive properties. Consider the case in which $C_A = \varepsilon S$, where $S$ is an invertible symmetric matrix. In the limit of the parameter, $\varepsilon$, becoming indefinitely large, $C_m$ does not also become indefinitely large, but rather tends to a constant:

$$\lim_{\varepsilon \to \infty} C_m = \lim_{\varepsilon \to \infty} [I - \varepsilon SG[T][\varepsilon GSG^T + C_d]^{-1} G] \varepsilon S =$$

$$\lim_{\varepsilon \to \infty} [I - \varepsilon SG[T][\varepsilon^{-1} G^{-1} S^{-1} G^{-1} - \varepsilon^{-2} G^{-1} S^{-1} G^{-1} C_d G^{-1} S^{-1} G^{-1}] G] \varepsilon S =$$

$$G^{-1} C_d G^{-1} = [G^T C_d^{-1} G]^{-1}$$
This expression can be recognized as the posterior variance that arises from the data, only. The zero limit is:

$$\lim_{\varepsilon \to 0} C_m = \lim_{\varepsilon \to 0} [I - \varepsilon S G T [\varepsilon G S G T + C_d]^{-1} G] G S = \varepsilon S = 0$$

That is, when \(m_A\) is known very accurately, so is \(m_G\).

**A.2.** Derivation of the TV82-A and TV82-B identities, following Taantola and Valette (1982). Consider invertible symmetric matrices, \(C_1\) and \(C_2\), and arbitrary matrix, \(M\). The expression \(M^T + M^T C_1^{-1} MC_2 M^T\) can alternately be factored:

\[
M^T C_1^{-1} [C_1 + MC_2 M^T] = [C_2^{-1} + M^T C_1^{-1} M] C_2 M^T
\]

Multiplying by the inverses yields identity TV82-A:

\[
C_2 M^T [C_1 + MC_2 M^T]^{-1} = [C_2^{-1} + M^T C_1^{-1} M]^{-1} M^T C_1^{-1}
\]

Now consider the expression \(C_2 - C_2 M^T [C_1 + MC_2 M^T]^{-1} MC_2\), which by the above identity equals \(C_2 - [C_2 + M^T C_1^{-1} M]^{-1} M^T C_1^{-1} MC_2\). Factoring out the term in brackets

\[
[C_2^{-1} + M^T C_1^{-1} M]^{-1} \left[ [C_2^{-1} + M^T C_1^{-1} M] C_2 - M^T C_1^{-1} MC_2 \right]
\]

cancelling terms yields identity TV82-B:

\[
C_2 - C_2 M^T [C_1 + MC_2 M^T]^{-1} MC_2, = [C_2^{-1} + M^T C_1^{-1} M]^{-1}
\]

**A.3.** Thomas (1949) algorithm for a symmetric block-diagonal matrix is well-known; we reproduce it here for completeness. The \(i\)th row of the matrix has elements \(B, A_i, B^T\) and the right-hand size is \(a_i\). Consider the step in the upper-triangularization process when rows \((i - 1)\) and above have been triangularized, but rows \((i - 1)\) and below have not:

\[
\tilde{A}_{i-1} m^{(i-1)} + B^T m^{(i)} = \hat{a}_{i-1}
\]

\[
Bm^{(i-1)} + A_i m^{(i)} + B^T m^{(i+1)} = a_i
\]

The second row is modified by multiply the top row by \(-B \tilde{A}_{i-1}^{-1}\) and adding the result to the second, which eliminates the first term, yielding:

\[
[A_i - B \tilde{A}_{i-1}^{-1} B^T] m^{(i)} + B^T m^{(i+1)} = a_i - B \tilde{A}_{i-1}^{-1} \hat{a}_{i-1}
\]

Note that the new row has two terms, and that the coefficient of the second is always \(B^T\), which is the same pattern as the first row. Thus, the bottom row becomes a new top row, and the recursion is

\[
\tilde{A}_1 = A_1 \quad \text{followed by} \quad \tilde{A}_i = [A_i - B \tilde{A}_{i-1}^{-1} B^T]
\]

\[
\tilde{a}_1 = a_1 \quad \text{followed by} \quad \tilde{a}_i = a_i - B \tilde{A}_{i-1}^{-1} \hat{a}_{i-1}
\]

or equivalently

\[
\tilde{A}_1^{-1} = A_1^{-1} \quad \text{followed by} \quad \tilde{A}_i^{-1} = [A_i - B \tilde{A}_{i-1}^{-1} B^T]^{-1}
\]

\[
\tilde{b}_1 \equiv A_1^{-1} \tilde{a}_1 = A_1^{-1} a_1 \quad \text{followed by} \quad \tilde{b}_i \equiv A_i^{-1} \tilde{a}_i = A_i^{-1} [a_i - B \tilde{A}_{i-1}^{-1} \hat{a}_{i-1}] = A_i^{-1} [b_i - \tilde{b}_{i-1}]
\]
After the recursion, the matrix upper-bidiagonal with diagonals, $\mathbf{A}_i$ and $\mathbf{B}^T$ and the right-hand side is $\mathbf{d}_i$. It is back-solved as:

$$ \mathbf{m}^{(K)} = \mathbf{A}_K^{-1} \mathbf{a}_K \quad \text{followed by} \quad \mathbf{m}^{(i)} = \mathbf{A}_i^{-1} [\mathbf{a}_i - \mathbf{B}^T \mathbf{m}^{(i+1)}] $$

or equivalently

$$ \mathbf{m}^{(K)} = \mathbf{b}_K = \mathbf{A}_K^{-1} \mathbf{a}_K \quad \text{followed by} \quad \mathbf{m}^{(i)} = \mathbf{A}_i^{-1} [\mathbf{a}_i - \mathbf{B}^T \mathbf{m}^{(i+1)}] = \mathbf{b}_i - \mathbf{A}_i^{-1} \mathbf{B}^T \mathbf{m}^{(i-1)} $$

A.4. Verification of the identity in (36)

$$ [\mathbf{D} \mathbf{C}_A \mathbf{D}^T + \mathbf{C}_s]^{-1} \mathbf{D} = \mathbf{C}_s^{-1} \mathbf{D} [\mathbf{D}^T \mathbf{C}_s^{-1} \mathbf{D} + \mathbf{C}_A^{-1}]^{-1} \mathbf{D} \mathbf{C}_A $$

$$ [\mathbf{D} \mathbf{C}_A \mathbf{D}^T + \mathbf{C}_s]^{-1} \mathbf{D} \mathbf{C}_A = \mathbf{C}_s^{-1} \mathbf{D} [\mathbf{D}^T \mathbf{C}_s^{-1} \mathbf{D} + \mathbf{C}_A^{-1}]^{-1} \mathbf{D} \mathbf{C}_A $$

$$ \mathbf{D} \mathbf{C}_A [\mathbf{D}^T \mathbf{C}_s^{-1} \mathbf{D} + \mathbf{C}_A^{-1}] = [\mathbf{D} \mathbf{C}_A \mathbf{D}^T + \mathbf{C}_s] \mathbf{C}_s^{-1} \mathbf{D} $$

$$ [\mathbf{D} \mathbf{C}_A \mathbf{D}^T \mathbf{C}_s^{-1} \mathbf{D} + \mathbf{D}] = [\mathbf{D} \mathbf{C}_A \mathbf{D}^T \mathbf{C}_s^{-1} \mathbf{D} + \mathbf{D}] $$