# **4D-Var Project**

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# Derivation of Strong-Constraint 4D-Var Algorithm for Lorenz-63 model

### Strong-Constraint 4D-Var Overview

The idea of strong-constraint 4D-Var is that we want to find an *actual solution* to the ODE system given above that "best fits" the data. This approach carries the inherent assumption that the model is perfect if we have the perfect initial condition, which is a decent assumption in our case since we are running a synthetic trial with perturbed observations from this model to begin with.

In other words, we look to find the appropriate initial condition  $\mathbf{x}_0 = \mathbf{x}(t_0)$  that produces a solution  $\mathbf{x}(t)$  that is "closest" to the data over the entire time interval (where we cannot neglect system dynamics on each of these intervals, due to nonlinearity).

Formally, we look to find an initial condition  $x_0$  that minimizes the cost functional

$$J(\mathbf{x}_0) = rac{1}{2} (\mathbf{x}_0 - \mathbf{x}_0^b)^T (P_0^b)^{-1} (\mathbf{x}_0 - \mathbf{x}_0^b) + rac{1}{2} \sum_{k=0}^K (\mathscr{H}_k(\mathbf{x}_k) - \mathbf{y}_k)^T R_k^{-1} (\mathscr{H}_k(\mathbf{x}_k) - \mathbf{y}_k)$$

subject to

$$\mathbf{x}_{k+1} = \mathscr{M}_{k+1}(\mathbf{x}_k), \ \ k = 0, 1, \dots, K-1$$

where  $\mathcal{M}_k$  and  $\mathcal{H}_k$  are the full, nonlinear model ODE and observation operators,  $P_0^b$  is the (known) covariance of the background guess for the initial state, and  $R_k$  is the (known) covariance of the observation operator.

To minimize this cost functional, we take the gradient of J with respect to the initial condition,  $\mathbf{x}_0$ .

Note that, given a perturbation  $\delta \mathbf{x}_0$  of the initial condition, the first variation is going to have the form

$$\delta J = (
abla_{\mathbf{x}_0} J)^T \delta \mathbf{x}_0$$

Note that we can linearize our model about this perturbation to get a Tangent Linear Model (TLM). This will simplify our analysis by ignoring higher-order behavior of the perturbations, which is often unneccessary to get good results.

The perturbation of the initial condition is propagated through our model using the tangent linear equation

$$\delta \mathbf{x}_{k+1} = M_{k+1} \delta \mathbf{x}_k$$

where  $M_{k+1}$  is the Jacobian matrix of  $\mathscr{M}_{k+1}$ , with partial derivatives of  $\mathbf{x}_{k+1}$  with respect to  $\mathbf{x}_k$ .

Taking the first variation of the cost functional gives

$$\delta J = (\mathbf{x} - \mathbf{x}_0^b)^T (P_0^B)^{-1} \delta \mathbf{x}_0 + \sum_{k=0}^K (\mathscr{H}_k(\mathbf{x}_k) - \mathbf{y}_k)^T R_k^{-1} H_k \delta x_k$$

where  $H_k$  is the Jacobian of  $\mathscr{H}_k$ , the nonlinear observation model at timestep k.

Note that we want to explicitly determine the perturbation of the cost functional with respect to the perturbation of the *initial condition*, and that  $\delta \mathbf{x}_k$  depends on the perturbation of the initial condition by the relationship

$$\delta \mathbf{x}_{k+1} = M_{k+1} M_k \dots M_1 M_0 \delta \mathbf{x}_0$$

Also note that, if we do not have an observation at every timestep that we increment our state, then the linearizations used in the cost functional must be propogated through time appropriately (as in the equation above, multiplying the correct number of Jacobian matrices at each timestep that we have incremented our state vector and not recieved an observation).

Putting this all together, we are going to solve for the minimizer of this cost functional by introducing adjoint state vectors  $\mathbf{p}_k$  at each observation timestep  $k=0,1,\ldots,K$ . Since, by our Tangent Linear Model equation,  $\delta\mathbf{x}_k=M_k\delta\mathbf{x}_{k-1}$ , we have that  $\mathbf{p}_k^T(\delta\mathbf{x}_k-M_k\delta\mathbf{x}_{k-1})=0$ , so adding terms like this to the cost functional does not change its value.

Hence, we now look to minimize the augmented cost functional

$$\delta J = (\mathbf{x} - \mathbf{x}_0^b)^T (P_0^B)^{-1} \delta \mathbf{x}_0 + \sum_{k=0}^K (\mathscr{H}_k(\mathbf{x}_k) - \mathbf{y}_k)^T R_k^{-1} H_k \delta x_k - \sum_{k=0}^K \mathbf{p}_k^T (\delta \mathbf{x}_k - M_k \delta \mathbf{x}_{k-1})$$

Rearranging this equation, exploiting the symmetry of the covariance matrices, and regrouping in terms of each  $\delta \mathbf{x}_k$ , we have that

$$egin{aligned} \delta J &= \left[ (P_0^B)^{-1} (\mathbf{x}_0 - \mathbf{x}_0^b) + H_0^T R_0^{-1} (\mathscr{H}_0(\mathbf{x}_0) - \mathbf{y}_0) + M_0^T \mathbf{p}_1 
ight] \delta \mathbf{x}_0 \ &+ \left[ \sum_{k=1}^{K-1} H_k^T R_k^{-1} (\mathscr{H}_k(\mathbf{x}_k) - \mathbf{y}_k) - \mathbf{p}_k + M_k^T \mathbf{p}_{k+1} 
ight] \delta \mathbf{x}_k \ &+ \left[ H_K^T R_k^{-1} (\mathscr{H}_K(\mathbf{x}_K) - \mathbf{y}_K) - \mathbf{p}_K 
ight] \delta \mathbf{x}_K \end{aligned}$$

which is valid for any choice of the adjoint states  $\mathbf{p}_k$ .

So we are going to choose these states such that the only term that survives is  $\delta \mathbf{x}_0$ .

This means simply that

$$egin{aligned} \mathbf{p}_K &= H_K^T \mathbf{R}_k^{-1} (\mathscr{H}_K(\mathbf{x}_k) - \mathbf{y}_k) \ \mathbf{p}_k &= H_k^T R_k^{-1} (\mathscr{H}_k(\mathbf{x}_k) - \mathbf{y}_k) + M_k^T \mathbf{p}_{k+1}, \qquad k = K-1, K-2, \dots, 1 \ \mathbf{p}_0 &= (P_0^B)^{-1} (\mathbf{x}_0 - \mathbf{x}_0^b) + H_0^T R_0^{-1} (\mathscr{H}_0(\mathbf{x}_0) - \mathbf{y}_0) + M_0^T \mathbf{p}_1 \end{aligned}$$

Where we have found then that

$$\delta J = \mathbf{p}_0^T \delta \mathbf{x}_0$$

as desired.

So  $\mathbf{p}_0$  is the desired gradient that we will use in an iterative gradient descent algorithm.

Note that  $\mathbf{p}_0$  depends on all of the other  $\mathbf{p}_k$  in an iterative way backwards from  $\mathbf{p}_K$ .

We define the weighted innovation vector as

$$d_k = H_k^T R_k^{-1} (\mathscr{H}_k(\mathbf{x}_k) - \mathbf{y}_k)$$

Using this notation, we can rewrite this gradient in terms of a gradient on the initial state/background and initial observation ( $\delta J^b$ ), and a gradient on the remaining observations ( $\delta J^o$ ) by fixing

$$\delta J = \delta J^b + \delta J^o$$

where

$$\delta J^b = (P_0^B)^{-1}(\mathbf{x}_0 - \mathbf{x}_0^b)$$

and

$$\delta J^o = d_0 + M_0^T (d_1 + M_1^T (d_2 + M_2^T (\dots M_{K-1}^T (d_K))))$$

Hence, the algorithm for 4D-Var amounts to iteratively doing the following:

- 1. Given a guess for the initial condition, integrate the solution to the ODE forward in time using an ODE solver (like solve\_ivp provided in scipy) over a given set of time gridpoints. This gives us our  $\mathbf{x}_k$ .
- 2. Determine the needed Jacobian matrices  $M_k$  and  $H_k$  at each observation timestep, as well as the (weighted, nonlinear) innovations terms  $d_k$  that depend on each of these.

(Note that if we only have observations at some of the timesteps that we have states at, we need to take matrix products over the time interval between observations to get the correct Jacobians used in our cost functional)

3. Construct the gradient  $\delta J^o$  by working backward in time from the last observation timestep to the first observation timestep, multiplying the right sums by the right Jacobians (working

inside-out of the equation for  $\delta J^o$  given above)

- 4. Construct the full gradient  $\delta J$  by adding  $\delta J_o$  to  $\delta J^b$ .
- 5. Perform some sort of gradient descent algorithm to find the analyzed initial condition  $\mathbf{x}_0^a$  that minimizes the cost functional at this iteration.
- 6. Repeat steps 1-5 with this new initial condition. Repeat until some sort of convergence is achieved.

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# Introduction to the Physical Model

We are trying to assimilate data into the Lorenz-63 Model, which is a toy model from meterorology. The model is a first-order ODE system of the form

We wish to assimilate the Lorenz-63 equations by standard 4D-Var, where the Lorenz-63 equations are given by

$$rac{dx}{dt} = -\sigma(x - y)$$
 $rac{dy}{dt} = 
ho x - y - xz$ 
 $rac{dz}{dt} = xy - eta z$ 

where 
$$\sigma=10$$
,  $\rho=28$ , and  $\beta=8/3$ .

This system is known to be chaotic, which makes it a good candidate for testing how well our 4D-Var algorithm performs on a set of data.

# Data Source for Experimentation

Since this is a toy model, it is difficult to assimilate real-life datasets with this model. So we will use what the book calls a "twin experiment" or "synthetic run" to evaluate the performance of our algorithm.

In other words, the following process will be used to generate the observation data:

- Create a "true" solution by giving an ODE solver a "background" initial condition, and then integrate the solution over a time interval. This will be the reference for how well our algorithm behaves.
- 2. Create "observations" by taking the model solution at various gridpoints (not necessarily at every one), and perturbing it by some mean-zero gaussian value with a given covariance matrix. (This assumes that the observation operator is the identity operator)

This process will give us a set of "true" solutions at each time gridpoint, and then a set of "observations" at a selection of these gridpoints.

# Application to the Lorenz-63 system

We have that the Tangent Linear Model/Jacobian for the Lorenz-63 system for going from state timestep k to state timestep k+1 (assuming that the length of time between these timesteps is the constant  $\Delta t$ ) is given by

$$M_{k:k+1} = egin{pmatrix} 1 - \sigma \Delta t & \sigma \Delta t & 0 \ (
ho - z_k) \Delta t & 1 - \Delta t & -x_k \Delta t \ y_k \Delta t & x_k \Delta t & 1 - eta \Delta t \end{pmatrix}$$

Our data assimilation method only has observed data at every other timestep where we run our model. So we actually need to multiply two of these matrices together to get the  $M_k$  values used in the cost functional.

Also, by the data generation algorithm given above, we have that the observation operator is given by

$$\mathscr{H}_k(x_k,y_k,z_k)=(x_k,y_k,z_k)$$

which is just the identity opertaor, and thus,

$$H_k = egin{pmatrix} 1 & 0 & 0 \ 0 & 1 & 0 \ 0 & 0 & 1 \end{pmatrix}$$

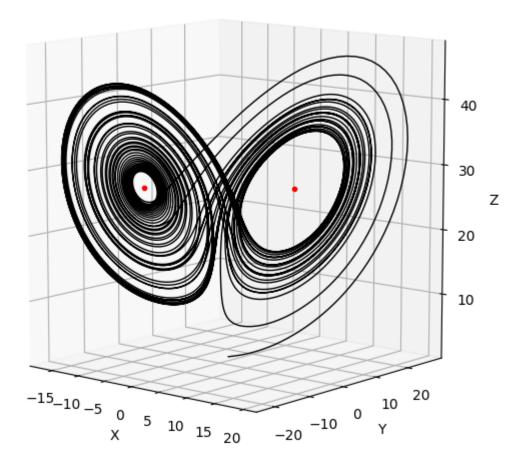
is the desired Jacobian used in our algorithm.

```
In []: # Imports
   import numpy as np
   from matplotlib import pyplot as plt
   from scipy.integrate import solve_ivp
   from functools import reduce
   from ast import literal_eval
In []: # Parameters of Lorenz-63 system
```

```
t_{span} = (0, 50)
t = np.linspace(0, 50, 10000)
x0 = np.ones(3)
example_sol = solve_ivp(lorenz, t_span, x0, t_eval=t)
# Plot the example solution
fig = plt.figure(figsize=(7,7))
ax = fig.add subplot(projection='3d')
ax.plot(example_sol.y[0,:], example_sol.y[1,:], example_sol.y[2,:], color='k', lin
ax.plot(np.sqrt(beta*(rho - 1)), np.sqrt(beta*(rho - 1)), rho-1, 'o', markersize=3
ax.plot(-np.sqrt(beta*(rho - 1)), -np.sqrt(beta*(rho - 1)), rho-1, 'o', markersize
ax.set_xlabel('X')
ax.set_ylabel('Y')
ax.set zlabel('Z')
ax.set_box_aspect(aspect=(1,1,1.1), zoom=0.95)
ax.view_init(10, -50, 0)
ax.set_title("Lorenz Attractor")
ax.legend()
plt.show()
```

#### Lorenz Attractor

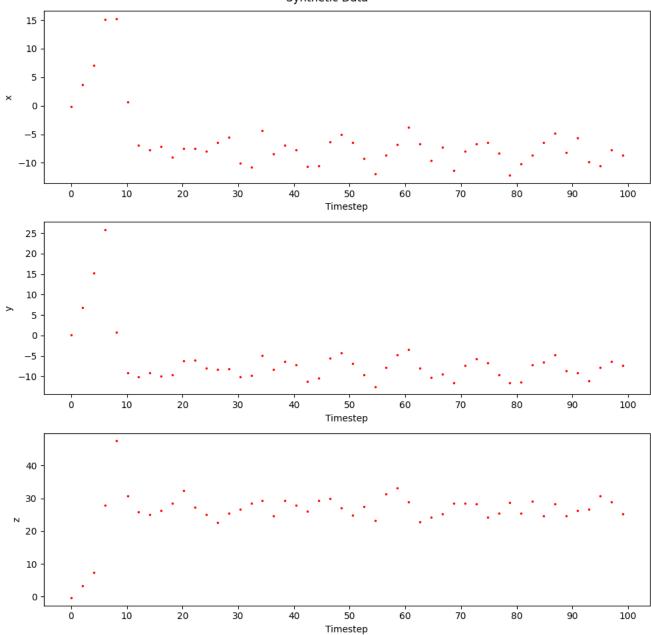
— Actual Solution



```
In [ ]: # Generate toy data for assimilation
        def get_observed_data(actual_data, covar):
            """Get observed datapoints given actual datapoints and
            an amount of noise/covariance.
            Parameters:
                actual_data ((n,t) ndarray) - Actual datapoints at
                  each timestep
                covar ((n,n) ndarray) - The covariance to use for
                  noise of observations
            Returns:
                observed_data ((n,t) ndarray) - Perturbed datapoints
                  at each timestep
                data_slice (Index Slice) - The slice of indexes we kept
            n, t = actual data.shape
            mu = np.zeros(n)
            perturbations = np.random.multivariate_normal(mu, covar, t).T
            all observations = actual data + perturbations
            return np.delete(all_observations, np.s_[1::2], axis=1), np.s_[::2] # we only
        def get_Li(state: np.ndarray,
                     dt: float) -> np.ndarray:
            """Gets the TLM of the Lorenz equations.
            Parameters:
                state ((3,) ndarray): The model-integrated/forecast state
                    at the current timestep.
                dt (float): The size of the timestep in our evaluation.
            Returns:
                ((3,3) ndarray): The TLM for the Lorenz model.
            x,y,z = state
            L = np.array([
                [1 - sigma * dt, sigma * dt, 0],
                [(rho - z) * dt, 1 - dt, -x * dt],
                [dt * y, dt * x, 1 - beta * dt]
            ]) # TLM derived from perturbing ODE and ignoring higher order terms
            return L
        def get_di(state: np.ndarray,
                   obs: np.ndarray,
                   H: callable,
                   H_linear: np.ndarray,
                   R_inv: np.ndarray,) -> np.ndarray:
            """Get the weighted innovation vector at a specific timestep.
            Currently not vectorized, could be improved by vectorizing over
            entire time interval.
```

```
Parameters:
        state ((n,) ndarray): The (forecast) state (resulting from
            a forward integration of the nonlinear model)
        obs ((m,) ndarray): The observations at each timestep.
        H (callable): The (potentially nonlinear) observation operator,
            vectorized to handle an entire vector of observations.
        H_linear ((n,m) ndarray): The linearized version of the observation
            operator (partial H/partial x i).
        R_inv ((m,m) ndarray): The inverse of the background covariance
            matrix.
    Returns:
        d_i (np.ndarray): The weighted innovation vectors at each
            timestep.
    unweighted_innovation = H(state) - obs
    return -H_linear.T @ R_inv @ unweighted_innovation
# Get actaul solution (to compare 4D-Var against)
t_{span} = (0, 5)
t = np.linspace(0, 5, 100) # 5 seconds with time steps of 0.05
x0 \text{ true} = np.ones(3)
                               # true/background initial condition
sol = solve_ivp(lorenz, t_span, x0_true, t_eval=t)
# Get observed data for the above actual solution (for assimilation)
# NOTE: Observed data starts at timestep 1 (not 0), and only is
# found at every other timestep of the solution sol.y
# (aka, indexes 1, 3, 5, ..., or in other words, sliced by [1::2])
R = np.array([
   [3,2,1],
    [2,2,2],
   [1,2,4]
])
observed data, data slice = get observed data(sol.y, R)
# Plot this all on the same axis
fig, axs = plt.subplots(3, figsize=(10,10))
plt.setp(axs, xticks=np.arange(0, 10.5, 0.5), xticklabels=np.arange(0, 210, 10))
for i, (ax, label) in enumerate(zip(axs, ['x', 'y', 'z'])):
   ax.set_ylabel(label)
    ax.set_xlabel('Timestep')
    ax.plot(t[data_slice], observed_data[i,:], 'r.', ms=3, label='Observations')
fig.suptitle("Synthetic Data")
fig.tight_layout()
plt.savefig("good_example_2.jpg")
fig.show()
```

/var/folders/x\_/vtkqq9c562v45n61xlf3t5cr0000gq/T/ipykernel\_45818/2758749215.py:103:
UserWarning: Matplotlib is currently using module://matplotlib\_inline.backend\_inlin
e, which is a non-GUI backend, so cannot show the figure.
 fig.show()



```
# Code up one iteration of inner loop to try this out
def analyze_4dvar_lorenz(
    t: np.ndarray,
   x0_b: np.ndarray,
    data: np.ndarray,
    B: np.ndarray,
    R: np.ndarray,
    H: callable,
   H_linear: np.ndarray,
    tol: float=1e-6,
   maxiter: int=100,
    data_slice: np.lib.index_tricks.IndexExpression=np.s_[:],
    verbose: bool=False,
    alpha: float = 0.01
    )-> np.ndarray:
    """Get the analyzed solution to a Lorenz-63 DA problem using 4D-Var.
```

```
Parameters:
    t ((t,) ndarray): The timesteps to evaluate the solution at.
    x0_b ((3,) ndarray): The background for the initial condition.
    data ((m,t*) ndarray): The data for assimilation with the Lorenz-63
        model (must have at least 2 observations)
    B ((3,3) ndarray): The covariance matrix for the background
        initial condition.
    R ((m,m) ndarray): The covariance matrix for the observations.
    H (callable): The (potentially nonlinear) observation operator,
        transforming states to observations.
    H linear ((3,m) ndarray): The linearized observation operator
        (must be the same at each timestep for this function)
    data_slice (IndexExpression): If provided, the slices for the timesteps
        at which data are provided (if not provided, assumed to be at
        all timesteps)
    verbose (boolean): Whether or not to print out status reports during
        runtime (default is False)
    alpha (float): The gradient descent step length parameter (default
        is 0.01)
Returns:
   analysis ((t,3) ndarray): The analyzed state after using 4DVar.
# Setup inverse matrices and other needed parameters
B inv = np.linalg.inv(B)
R_{inv} = np.linalg.inv(R)
t_span = (np.min(t), np.max(t)) # Min and max time on interval
dt = t[1] - t[0]
                                # Size of timestep
# Set up cost function
J = lambda x: 0.5 * (x[0] - x0_b).T @ B_inv @ (x[0] - x0_b) \
             + 0.5 * np.sum([(H(x) - y).T @ R_inv @ (H(x) - y)])
              for x,y in zip(x[data_slice], data.T)])
# Get indexes of timesteps we have data at
data_times = np.arange(len(t))[data_slice]
padded data times = list(data times)  # Used for calculating adjoint
if min(data_times) != 0:
    padded_data_times.insert(0, 0)
if max(data_times) != len(t) - 1:
    padded_data_times.append(len(t) - 1)
# ----- 4D-Var Loop -----
norm_grad_J = np.inf
best_J = np.inf
numiter = 0
x0_a = x0_b.copy() # Analyzed initial condition (start at background)
success = False
while norm grad J > tol and numiter < maxiter:
    if verbose and numiter%500==0:
        print(f"ITERATION: {numiter}")
   ## Step 1: Run full nonlinear model to get forecast state
   ## (AKA forward integration)
    sol = solve_ivp(lorenz, t_span, x0_a, t_eval=t)
```

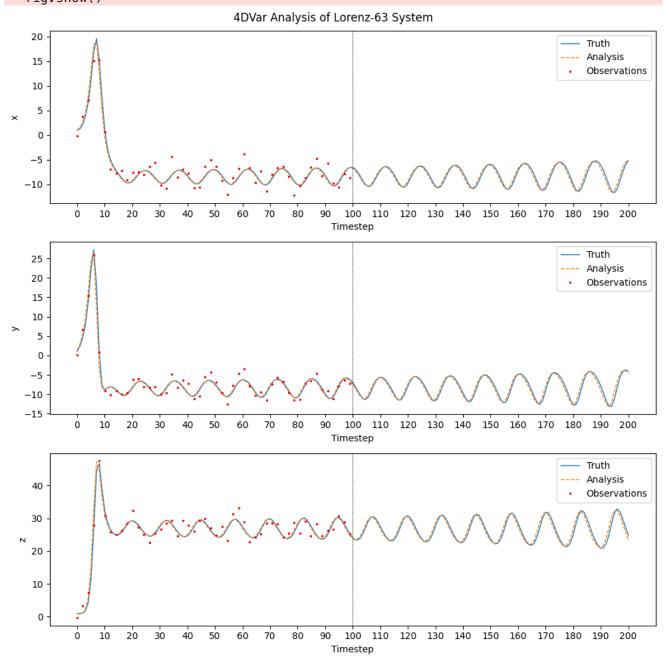
```
state_a = sol.y # Shape is (3,t)
    ## Step 2: Get adjoint/innovation information using this state
    # Get (non-transposed) TLM using the current states and timesteps
    Li raw = []
    for state in state a.T:
        Li_raw.append(get_Li(state, dt))
    # Combine (transposed) adjoints at correct timesteps
    # to align correctly with states
    adjoints = [
        (reduce(np.dot, Li raw[start:end])).T
        for start, end
        in zip(padded_data_times[:-1], padded_data_times[1:])
    # Get weighted innovations
    innovations = []
    for state, obs in zip(state a.T[data slice], data.T):
        innovations.append(get_di(state, obs, H, H_linear, R_inv))
    ## Step 3: Work "backwards in time" to get gradient of J o
    grad_J_o = innovations[-1]
    for adjoint, innovation in zip(adjoints[::-1], innovations[:-1:-1]):
        grad J o = innovation + adjoint @ grad J o
    ## Step 4: Get full gradient
    grad J b = B inv @ (x0 a - x0 b)
    grad_J = grad_J_b + grad_J_o
    norm_grad_J = np.linalg.norm(grad_J, ord=2)
    if verbose and numiter%500 == 0:
        print(f" GRADIENT: {grad_J}")
        print(f" L2-NORM OF GRADIENT: {norm_grad_J}")
    ## Step 5: If we have converged, break.
    ## Otherwise, run full gradient descet algorithm
    # Check for convergence (gradient being "small enough")
    if norm grad J < tol:</pre>
        success = True
        break
    # Gradient descent that works (fixed rate)
    alpha used = alpha
    if norm grad J < 1: # Smaller step size for smaller gradient</pre>
        alpha used *= 0.1
    if norm_grad_J < 0.1:</pre>
        alpha used *= 0.05
    x0_a -= alpha_used * grad_J
    if verbose and numiter%500 == 0:
        print(f" NEW x0 a: {x0 a}")
    # Set up for next iteration
    numiter += 1
# End of algorithm. Check for successful convergence
```

```
if not success:
                print(f"WARNING: Algorithm did not converge in {maxiter} iterations")
            # Get analyzed state by integrating with new, analyzed initial condition,
            # and return it
            print(f"FINAL x0 a: {x0 a}")
            print(f"FINAL GRADIENT NORM: {norm grad J}")
            sol = solve_ivp(lorenz, t_span, x0_a, t_eval=t)
            state a = sol.y # Shape is (3,t)
            return state_a
In [ ]: # Get model covariance matrix (overestimate is good since we don't know
        # how much to trust the model)
        B = np.eye(3)
        R_{perturbed} = np.eye(3) * 5
        # Get observation operators (full and linearized)
        H = lambda x: x
        H_{linear} = np.eye(3)
        # Get the analysis
        alpha size = 0.01
        x0_b = np.array([0.7, 1.2, 0.9])
        analysis = analyze_4dvar_lorenz(t, x0_b, observed_data, B, R_perturbed, H, H_linea
                                        data_slice=data_slice, maxiter=20000, tol=5e-2,
                                        verbose=False, alpha=alpha_size)
        # Forecast both actual and anlyzed solutions to double the time
        t span long = (0, 10)
        t long = np.linspace(0, 10, 200)
        # Actual solution
        x0 b = np.ones(3)
                                   # true/background initial condition
        actual_sol = solve_ivp(lorenz, t_span_long, x0_b, t_eval=t_long)
        # Analyzed solution forecast
        x0 a = analysis[:,0]
        analyzed_forecast_sol = solve_ivp(lorenz, t_span_long, x0_a, t_eval=t_long)
        # Plot this all on the same axis
        fig, axs = plt.subplots(3, figsize=(10,10))
        plt.setp(axs, xticks=np.arange(0, 10.5, 0.5), xticklabels=np.arange(0, 210, 10))
        for i, (ax, label) in enumerate(zip(axs, ['x', 'y', 'z'])):
            ax.set_ylabel(label)
            ax.set_xlabel('Timestep')
            ax.plot(t_long, actual_sol.y[i,:], label='Truth', lw=1)
            ax.plot(t_long, analyzed_forecast_sol.y[i,:], '--', label='Analysis', lw=1)
            ax.plot(t[data_slice], observed_data[i,:], 'r.', ms=3, label='Observations')
            ax.axvline(5, linestyle='--', color='k', lw=0.5)
            ax.legend()
        fig.suptitle("4DVar Analysis of Lorenz-63 System")
        fig.tight layout()
        plt.savefig("good_example_2.jpg")
        fig.show()
```

FINAL x0\_a: [1.08145789 1.35124336 0.82493907]

FINAL GRADIENT NORM: 0.049998819617316614

/var/folders/x\_/vtkqq9c562v45n61xlf3t5cr0000gq/T/ipykernel\_45818/3234203672.py:43:
UserWarning: Matplotlib is currently using module://matplotlib\_inline.backend\_inline, which is a non-GUI backend, so cannot show the figure.
 fig.show()



## **Climate Data Assimilation**

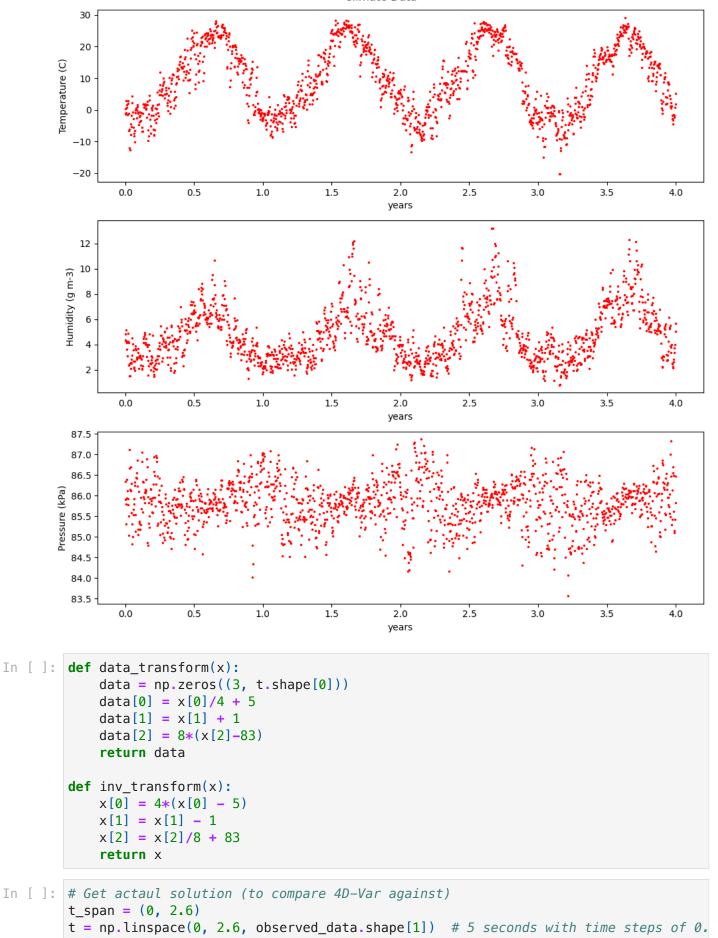
Lorenz-63 model is a simplified version of an atmospheric model used for weather forcast. In this context, each of the variables x, y, z represents a specific state of the atmosphere. For example, x can represent temperature, y can represent humidity, and z can represent pressure. Although the model is overly simplified and shouldn't really be used for weather forcast, we found some real world data that could be used for data assimilation using our model. The data

was obtained from https://caas.usu.edu/weather/ and contains 4-year temperature, humidity, and barometric pressure data.

The data is first transformed to match the expected values of the Lorenz-64 model output with the specified parameter values as before. The transformed data is then used to fit the model, and the resulting model output is inversely transformed to represent the values of the original observed data. This transformation can be viewed as our observation model  $\mathcal{H}_k$ .

```
In [ ]: # Load data
        f = open("pressure.txt")
        pressure = np.array(literal_eval(f.read()))
        f.close()
        f = open("abs humidity.txt")
        abs_humidity = np.array(literal_eval(f.read()))
        f.close()
        f = open("temperature.txt")
        temperature = np.array(literal eval(f.read()))
        f.close()
        # There were some missing temperature data.
        # The average value surrounding the data
        # was used to fill in the missing values.
        temperature[890:902, 1] = (temperature[889, 1] + temperature[902, 1]) / 2
        temperature [1029:1037, 1] = (temperature [1028, 1] + temperature [1037, 1]) / 2
        observed_data = np.zeros((3, temperature.shape[0]))
        observed data[0] = temperature[:, 1]
        observed data[1] = abs humidity[:, 1]
        observed data[2] = pressure[:, 1]
        t = np.linspace(0, 4, temperature.shape[0])
        # Plot this all on the same axis
        fig, axs = plt.subplots(3, figsize=(10,10))
        for i, (ax, label) in enumerate(zip(axs, ['Temperature (C)', 'Humidity (g m-3)', '
            ax.set ylabel(label)
            ax.set xlabel('years')
            ax.plot(t, observed_data[i,:], 'r.', ms=3, label='Observations')
        fig.suptitle("Climate Data")
        fig.tight layout()
        fig.show()
```

/var/folders/x\_/vtkqq9c562v45n61xlf3t5cr0000gq/T/ipykernel\_45818/2301448032.py:34:
UserWarning: Matplotlib is currently using module://matplotlib\_inline.backend\_inlin
e, which is a non-GUI backend, so cannot show the figure.
 fig.show()



```
x0_{true} = np.ones(3) # true/background initial condition
 sol = solve_ivp(lorenz, t_span, x0_true, t_eval=t)
 # Get observed data for the above actual solution (for assimilation)
 # NOTE: Observed data starts at timestep 1 (not 0), and only is
 # found at every other timestep of the solution sol.y
 # (aka, indexes 1, 3, 5, ..., or in other words, sliced by [1::2])
 transformed data = data transform(observed data)
 # Get model covariance matrix (overestimate is good since we don't know
 # how much to trust the model)
 B = 2*np.eye(3)
 R perturbed = np.eye(3) * 5
 # Get observation operators (full and linearized)
 H = lambda \times : \times
 H_{linear} = np.eye(3)
 # Get the analysis
 alpha size = 0.01
 x0_b = np.array([5.1, 5.3, 22.8])
 analysis = analyze_4dvar_lorenz(t, x0_b, transformed_data, B, R_perturbed, H, H_li
                                 maxiter=50000, tol=5e-2,
                                 verbose=False, alpha=alpha_size)
FINAL x0 a: [ 5.71278877 3.11436604 25.03484356]
FINAL GRADIENT NORM: 0.04999953611187953
 t_{long} = np.linspace(0, 3.5, 200)
 # Analyzed solution forecast
```

In  $[ ]: t_span_long = (0, 3.5)$  $x0_a = analysis[:,0]$ analyzed forecast sol = solve ivp(lorenz, t span long, x0 a, t eval=t long) y = inv transform(analyzed forecast sol.y) data\_orig = np.zeros((3, temperature.shape[0])) data orig[0] = temperature[:, 1] data orig[1] = abs humidity[:, 1] data\_orig[2] = pressure[:, 1] # Plot this all on the same axis fig, axs = plt.subplots(3, figsize=(10,10)) plt.setp(axs, xticks=np.arange(0, 10.5, 0.5), xticklabels=np.arange(0, 210, 10)) for i, (ax, label) in enumerate(zip(axs, ['Temperature (C)', 'Humidity (g m-3)', ax.set ylabel(label) ax.set\_xlabel('Timestep') ax.plot(t\_long, y[i], '--', label='Analysis', lw=1) ax.plot(t, data\_orig[i,:], 'r.', ms=3, label='Observations') ax.axvline(2.6, linestyle='--', color='k', lw=0.5) ax.legend() fig.suptitle("4DVar Analysis of Lorenz-63 System fitted on Climate data") fig.tight\_layout() fig.show()

/var/folders/x\_/vtkqq9c562v45n61xlf3t5cr0000gq/T/ipykernel\_45818/3443905760.py:25:
UserWarning: Matplotlib is currently using module://matplotlib\_inline.backend\_inline, which is a non-GUI backend, so cannot show the figure.
 fig.show()

