VDP lorenzx

October 11, 2023

```
[]: import numpy as np
     from scipy.integrate import solve_ivp
     import pandas as pd
     import matplotlib.pyplot as plt
     from reservoirpy.observables import nrmse, rsquare
     import reservoirpy as rpy
     # just a little tweak to center the plots, nothing to worry about
     from IPython.core.display import HTML
     HTML("""
     <style>
     .img-center {
        display: block;
         margin-left: auto;
         margin-right: auto;
         }
     .output_png {
        display: table-cell;
         text-align: center;
         vertical-align: middle;
         }
     </style>
     """)
     rpy.set_seed(42)
     rpy.verbosity(0)
     from reservoirpy.nodes import Reservoir, Ridge
     from reservoirpy.datasets import mackey_glass
```

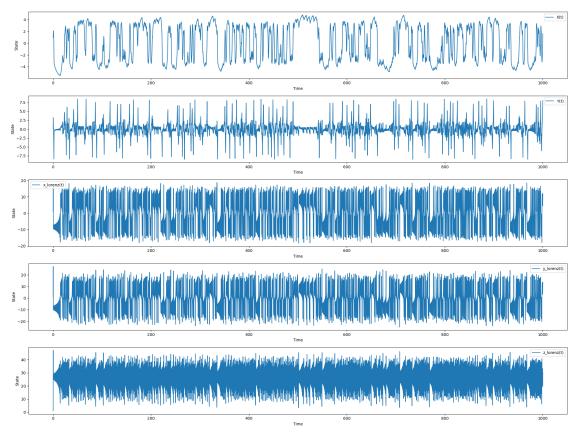
```
[]: import numpy as np
from scipy.integrate import solve_ivp
import matplotlib.pyplot as plt

def coupled_system(t, state, mu, sigma, rho, beta):
    X, Y, x_lorenz, y_lorenz, z_lorenz = state
```

```
dXdt = Y
    dYdt = mu * (1 - X**2) * Y - X + x_lorenz
    dx_lorenz_dt = sigma * (y_lorenz - x_lorenz)
    dy_lorenz_dt = x_lorenz * (rho - z_lorenz) - y_lorenz
    dz_lorenz_dt = x_lorenz * y_lorenz - beta * z_lorenz
    return [dXdt, dYdt, dx_lorenz_dt, dy_lorenz_dt, dz_lorenz_dt]
mu = 1.0
sigma = 10.0
rho = 28.0
beta = 8/3
initial_state = [1.0, 0.0, 1.0, 1.0, 1.0] # Initial states for [X, Y, _____
 \rightarrow x_lorenz, y_lorenz, z_lorenz]
t span = (0, 1000)
sol = solve_ivp(coupled_system, t_span, initial_state, args=(mu, sigma, rho, __
 ⇒beta), t_eval=np.linspace(t_span[0], t_span[1], 10000))
# Plot the results
plt.figure(figsize=(20, 15))
plt.subplot(5, 1, 1)
plt.plot(sol.t, sol.y[0], label='X(t)')
plt.xlabel('Time')
plt.ylabel('State')
plt.legend()
plt.subplot(5, 1, 2)
plt.plot(sol.t, sol.y[1], label='Y(t)')
plt.xlabel('Time')
plt.ylabel('State')
plt.legend()
plt.subplot(5, 1, 3)
plt.plot(sol.t, sol.y[2], label='x_lorenz(t)')
plt.xlabel('Time')
plt.ylabel('State')
plt.legend()
plt.subplot(5, 1, 4)
plt.plot(sol.t, sol.y[3], label='y_lorenz(t)')
plt.xlabel('Time')
plt.ylabel('State')
plt.legend()
```

```
plt.subplot(5, 1, 5)
plt.plot(sol.t, sol.y[4], label='z_lorenz(t)')
plt.xlabel('Time')
plt.ylabel('State')
plt.legend()

plt.tight_layout()
plt.show()
```

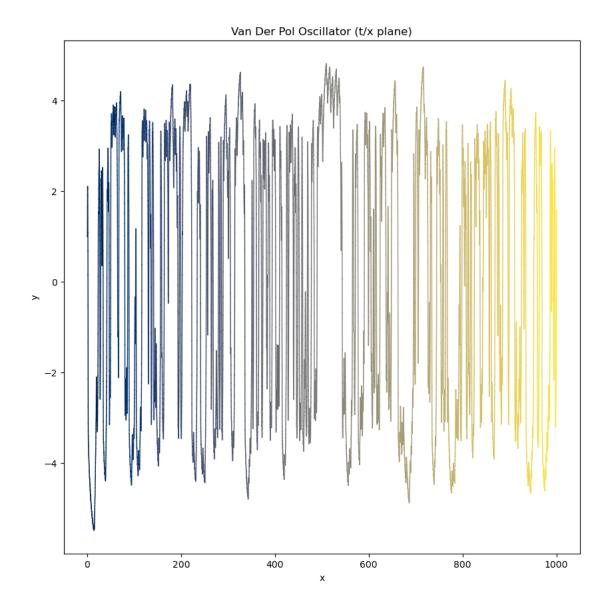


```
[]: #Time, Van Der Pol Oscillator X, lorenz x

data = pd.DataFrame({
    'Time': sol.t,
    'X': sol.y[0],
    'x_lorenz': sol.y[2]
})

# DataFrame CSV
data.to_csv('coupled_system_dataset.csv', index=False)
```

```
[]: data_loaded = pd.read_csv('forced_vanderpol_dataset.csv')
[]: X = data.values
[]: X.shape
[]: (10000, 3)
[]: def plot_vanderpol_2d(X, timesteps):
        # 2D
        fig, ax = plt.subplots(figsize=(10, 10))
        ax.set_title("Van Der Pol Oscillator (t/x plane)")
        ax.set_xlabel("x")
        ax.set_ylabel("y")
        #
        for i in range(timesteps-1):
            ax.plot(X[i:i+2, 0], X[i:i+2, 1], color=plt.cm.cividis(255*i//
      →timesteps), lw=1.0)
        plt.show()
[]: plot_vanderpol_2d(X, 10000)
```



1 esn fitting

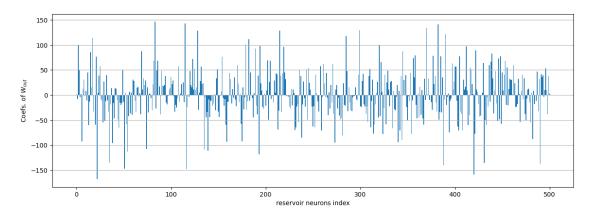
```
[]: X = X[:, 1]
#z-score normalization
X = (X - np.mean(X)) / np.std(X)

[]: X = X.reshape(-1, 1)
X.shape
[]: (10000, 1)
```

```
[ ]: X
[]: array([[0.32261709],
            [0.3229254],
            [0.32770749],
            [0.03927255],
            [0.26074807],
            [0.51432419]])
[]: from reservoirpy.datasets import to_forecasting
     #1
     X, Y = to_forecasting(X, forecast=1)
     X_train1, Y_train1 = X[:1000], Y[:1000]
     X_test1, Y_test1 = X[1000:], Y[1000:]
[]: #
     units = 500
     leak_rate = 0.17113764962939737
     spectral_radius = 1.182082887824895
     input_scaling = 0.9697295578201843
     connectivity = 0.10407566959031667
                                             # - density of reservoir internal matrix
     input_connectivity = 0.2 # and of reservoir input matrix
     regularization = 5.602814683095597e-09
     seed = 1234
                             # for reproducibility
[]: def reset_esn():
         from reservoirpy.nodes import Reservoir, Ridge
         reservoir = Reservoir(units, input_scaling=input_scaling,__
      ⇔sr=spectral_radius,
                               lr=leak_rate, rc_connectivity=connectivity,
                               input_connectivity=input_connectivity, seed=seed)
                   = Ridge(1, ridge=regularization)
         return reservoir >> readout
[]: from reservoirpy.nodes import Reservoir, Ridge
     reservoir = Reservoir(units, input_scaling=input_scaling, sr=spectral_radius,
                           lr=leak_rate, rc_connectivity=connectivity,
                           input_connectivity=input_connectivity, seed=seed)
               = Ridge(1, ridge=regularization)
     readout
     esn = reservoir >> readout
```

```
[]: | y = esn(X[0]) # initialisation
     reservoir.Win is not None, reservoir.W is not None, readout.Wout is not None
[]: (True, True, True)
[]: np.all(readout.Wout == 0.0)
[]: True
[]: esn = esn.fit(X_train1, Y_train1, warmup=100)
[]: def plot_readout(readout):
         Wout = readout.Wout
         bias = readout.bias
         Wout = np.r_[bias, Wout]
         fig = plt.figure(figsize=(15, 5))
         ax = fig.add_subplot(111)
         ax.grid(axis="y")
         ax.set_ylabel("Coefs. of $W_{out}$")
         ax.set_xlabel("reservoir neurons index")
         ax.bar(np.arange(Wout.size), Wout.ravel()[::-1])
         plt.show()
```

[]: plot_readout(readout)

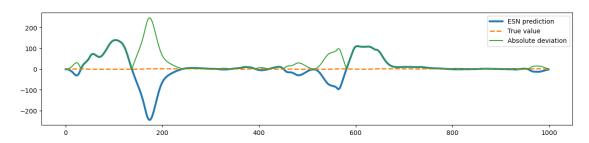


```
[]: def plot_results(y_pred, y_test, sample=1000):
    fig = plt.figure(figsize=(15, 7))
    plt.subplot(211)
    plt.plot(np.arange(sample), y_pred[:sample], lw=3, label="ESN prediction")
```

[]: Y_pred1 = esn.run(X_test1)

$m 2 \qquad \qquad RC$

[]: plot_results(Y_pred1, Y_test1)



```
[ ]: rsquare(Y_test1, Y_pred1), nrmse(Y_test1, Y_pred1)
```

[]: (-2102.3865272720186, 14.545452155873795)

3 nrmse 14.545452155873795

3.1 spectral_radius, leak_rate, input_scaling, regularization, connectivity.

```
def objective(dataset, config, *, cell_number, leak_rate, spectral_radius, input_connectivity, input_scaling, connectivity, regularization, seed):

X, Y = to_forecasting(dataset, forecast=10)
X_train1, Y_train1 = X[:1000], Y[:1000]
X_test1, Y_test1 = X[1000:], Y[1000:]

instances = config["instances_per_trial"]

variable_seed = seed

losses = []; r2s = [];
```

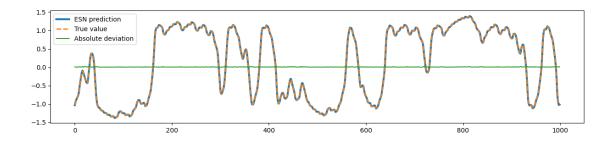
```
for n in range(instances):
             reservoir = Reservoir(cell_number, input_scaling=input_scaling,__
      ⇔sr=spectral_radius,
                             lr=leak rate, rc connectivity=connectivity,
                             input_connectivity=input_connectivity, seed=seed)
             readout
                       = Ridge(1, ridge=regularization)
             esn = reservoir >> readout
             esn = esn.fit(X_train1, Y_train1, warmup=100)
             Y_pred1 = esn.run(X_test1)
             loss = nrmse(Y_test1, Y_pred1)
             r2 = rsquare(Y_test1, Y_pred1)
             losses.append(loss)
             r2s.append(r2)
         return {'loss': np.mean(losses),
                 'r2': np.mean(r2s)}
[]: hyperopt_config = {
         "exp": f"hyperopt-MG-onestep-prediction", # the experimentation name
         "hp_max_evals": 200,
                                          # the number of differents sets of
      →parameters hyperopt has to try
         "hp method": "random",
                                       # the method used by hyperopt to chose
      ⇔those sets (see below)
         "seed": 42,
                                          # the random state seed, to ensure
      \hookrightarrow reproducibility
         "instances_per_trial": 3,
                                          # how many random ESN will be tried with
      ⇔each sets of parameters
         "hp_space": {
                                          # what are the ranges of parameters
      \rightarrow explored
                                                        # the number of neurons is_{\sqcup}
             "cell_number": ["choice", 500],
      → fixed to 500
             "spectral_radius": ["loguniform", 1e-2, 10], # the spectral radius is_{\sqcup}
      ⇔log-uniformly distributed between 1e-2 and 10
             "leak rate": ["loguniform", 1e-3, 1], # idem with the leaking rate, __
      \hookrightarrow from 1e-3 to 1
             "input_scaling": ["uniform", 0, 1],
                                                          # the input scaling
      →uniformly distributed between 0 and 1
             "input_connectivity": ["choice", 0.2], # the number of neurons is fixed_
      →to 0.2
```

```
"regularization": ["loguniform", 1e-10, 1], # the regularization⊔
      ⇒parameter is log-uniformly distributed between 1e-10 and 1.
             "connectivity": ["uniform", 0, 1], # the connectivity of the randomu
      →network in the hidden layer uniformly distributed between 0 and 1.
             "seed": ["choice", 1234]
                                               # an other random seed for the ESN_
      \hookrightarrow initialization
         }
     }
     import json
     # we precautionously save the configuration in a JSON file
     \# each file will begin with a number corresponding to the current
      ⇔experimentation run number.
     with open(f"{hyperopt_config['exp']}.config.json", "w+") as f:
         json.dump(hyperopt_config, f)
[]: from reservoirpy.hyper import research
     best = research(objective, X, f"{hyperopt_config['exp']}.config.json", ".")
               | 200/200 [20:52<00:00, 6.26s/trial, best loss:
    0.11718737488530441
[]: best
[]: ({'cell_number': 0,
       'connectivity': 0.9548228482666989,
       'input connectivity': 0,
       'input_scaling': 0.5933563673798359,
       'leak_rate': 0.8382221591273349,
       'regularization': 2.236024179842572e-06,
       'seed': 0,
       'spectral_radius': 0.7323231724059845},
      <hyperopt.base.Trials at 0x297971b50>)
            hyperparameters
    4
                                     esn fitting
```

```
[]: #
    units = 500
    leak_rate = 0.8382221591273349
    spectral_radius = 0.7323231724059845
    input_scaling = 0.5933563673798359
    connectivity = 0.9548228482666989 # - density of reservoir internal matrix
    input_connectivity = 0.2 # and of reservoir input matrix
    regularization = 2.236024179842572e-06
```

```
seed = 1234
                              # for reproducibility
[]: from reservoirpy.nodes import Reservoir, Ridge
     reservoir = Reservoir(units, input_scaling=input_scaling, sr=spectral_radius,
                            lr=leak_rate, rc_connectivity=connectivity,
                            input_connectivity=input_connectivity, seed=seed)
     readout
               = Ridge(1, ridge=regularization)
     esn = reservoir >> readout
[]: y = esn(X[0]) # initialisation
     reservoir.Win is not None, reservoir.W is not None, readout.Wout is not None
[]: (True, True, True)
[]: np.all(readout.Wout == 0.0)
[]: True
[]: esn = esn.fit(X_train1, Y_train1, warmup=100)
[ ]:  #W_out
     plot_readout(readout)
         Coefs. of Wout
          -15
                                            reservoir neurons index
[]: Y_pred1 = esn.run(X_test1)
```

[]: plot_results(Y_pred1, Y_test1)



```
[]: rsquare(Y_test1, Y_pred1), nrmse(Y_test1, Y_pred1)
[]: (0.9999515001268733, 0.0022087063689438216)
      nrmse\ 0.0022087063689438216
       14.545452155873795
[]: X_test1.shape
[]: (8999, 1)
            X Y_pred1
    6
[]: Y_pred1[-10:]
[]: array([[-0.75039814],
            [-0.69751703],
            [-0.63477833],
            [-0.54839908],
            [-0.43259336],
            [-0.29977342],
            [-0.15150833],
            [ 0.03041797],
            [ 0.24860936],
            [ 0.5010843 ]])
[]: X[-9:]
[]: array([[-0.74533189],
            [-0.69160057],
            [-0.6266132],
            [-0.54082037],
            [-0.43050355],
            [-0.29857988],
```

[-0.14443869],

[0.03927255], [0.26074807]])