Time Series Analysis in Python: User Manual

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1 Introduction

Time series analysis comprises methods for analyzing time series data in order to extract meaningful statistics and other characteristics of the data. It is widely used in signal processing, pattern recognition, mathematical finance, weather forecasting, earthquake prediction, control engineering, and largely in any domain of applied science and engineering which involves temporal measurements.

In this project, we will play a game with time series in finance. It has gained its popularity in Wall Street recently, since it is fundamental to most promising quantitative investment strategies. We develop a system that can predict future prices of stocks using various kinds of methods for time series analysis.

This manual is a brief introduction of the functionality of our package. In specific, we illustrate how to use the provided functions to fit the model, predict the stock price, trade, cluster etc. Introduction of the main source files and key functions is also included.

We utilize standard packages from python including numpy, scipy, and matplotlib. The integration of C code and Python code is based on the Cython package.

2 Program Structure

The high-level program structure is shown in Figure 1. The division of work is pretty even, and there are some minor work that are too trivial to mention extensively here. Overall, the program consists of two parts. The first part deals with a single time series, and exploits the time correlation within the time series. There is data preprocessing module that takes the raw data and get the right data format for later analysis. Optimization, model, and solver are used all together to fit the models using maximum likelihood estimation. Finally, there is a post processing module that makes use of the information from model. The post processing module includes parameter inference, trading strategy, and option pricing. The second part deals with a collection of time series of data, basically it exploits the correlation between different time series. In this way, we can gain more insights into the financial market. While these insights are impossible to obtain from a single time series.

3 Functionality

To start with, we will give an description of the source files.

• basemodel.py: Provided the basemodel of time series.

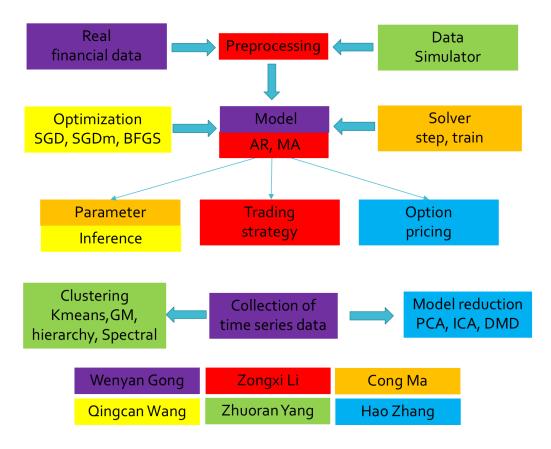


Figure 1: Program structure and division of work

- cluster.py: Provided the Cluster class for clustering. It includes four clustering methods: k-means, hierarchy clustering, Gaussian mixture modeling, and spectral clustering.
- Data_prossessor.py: Provided the transforming function between stock price and return, the function getting the maximum drawdown and the function returning the indicator of peak and trough.
- Gradient_check.py: Provided the function for deriving numerical gradient.
- Inference.py: Provided functions for computing the auto-covariance of AR model and testing significance of parameters.
- Model.py: Provided model class AR and MA, including functions for computing the loglikelihood and gradient given the data and an AR or MA model and the function for future price prediction.
- Optim.py: Provided several optimization methods: stochastic gradient descent, stochastic gradient descent with momentum update and Broyden–Fletcher–Goldfarb–Shanno algorithm
- Option_pricing.py: Provided functions for option pricing.
- Reduction.py: Provided functions for dimensional reduction for data visualization.
- Solver.py: Provided functions for model fitting using maximum likelihood estimation given the data and the specified model.
- Trading.py: Provided functions for deciding the transaction time point based on future price prediction and computing profit over the time.

• Ts_gen.pyx & c_ts_gen.c: Provided functions to generate simulation data.

3.1 Data Preparation and Preprocessing

We provide both a real dataset and methods to generate simulated data from various time series models. For data simulation, we provide functions that samples from the AR(p), MA(q), ARMA (p,q), GARCH(p,q) models, which is written in C code. See c_ts_gen.c for the source code. To integrate C code into Python, we write a Cython wrapper ts_gen.pyx, which generates a Python module named ts_gen. This module consists of five functions. The inputs of these functions all consist of four elements:

- 1. param: a numpy Nd array or a few float numbers the model parameters.
- 2. time: an integer the total time of the time series.
- 3. num: an integer number of independent sample paths,
- 4. burnin: an integer the length of burnin period. This parameter has default value 2000.

These functions all output a numpy Nd array with dimension (num \times time) which stores num independent paths of the time series data until T =time. Functions that generate simulated data are listed as follows.

For an example of usage, see the following example.

```
from tsap.ts_gen import arma1_gen
import numpy as np
Y = ar1_gen(0.5, sigma = 1.0, time = 200, num = 10, burnin = 2000)
print(Y.shape)
```

The output, as expected, is >>(10, 200)

3.2 Model

The model related functions are written in model.py, which contains two model classes: AR and MA. They are used to fit an AR or MA model by given parameters, to compute the log-likelihood and gradients given an AR or MA model and the data and to give predictions. The functions are introduced below. Firstly, you can fit an AR or MA model by given parameters. The inputs are as follows:

- 1. lag: the lag parameter for the fitted model, a number;
- 2. phi: the coefficients for the fitted model, a numpy array whose dimension is the lag given above;
- 3. sigma: the variance of noise for the fitted model, a number;
- 4. intercept: the intercept of noise for the fitted model, a number.

The output is a model object. To fit the model, we can run

```
AR_model = AR(lag=3, phi=np.array([[1],[1],[1]]), sigma=1, intercept=0.1)
MA_model = MA(lag=3, phi=np.array([[1],[1],[1]]), sigma=1, intercept=0.1)
```

Secondly, you can compute the log-likelihood and gradients given an AR or MA model and the data. The input is as follows:

1. data: One or several time series. The input data should be a 2-dimensional array. Each row of data represents a time series over the time, e.g. the stock return over the year of a single stock. The number of rows should be the number of time series.

The output is a number for the computed loglikelihood and a tuple for the computed gradients. This function is public but mostly it serve for the Solver class. To compute the log-likelihood and the gradients, we can run

```
AR_11h, AR_grads= AR_model.loss(data)
2 MA_11h, MA_grads= MA_model.loss(data)
```

Finally, you can make prediction based on a fitted model. The input is as follows:

- 1. data: the time series before the first prediction. The number of columns (the length of the time series) should exceed the given lag for a reasonable prediction;
- 2. nstep: the number of predictions.

To make predictions, we can run

```
AR_future=AR_model.predict(data,nstep=5)

2 MA_future=MA_model.predict(data,nstep=5)
```

3.3 Optimization

The optimization part optim.py implements various optimization update rules: stochastic gradient descent (sgd), stochastic gradient descent with momentum update (sgd_momentum) and Broyden-Fletcher-Goldfarb-Shanno algorithm (bfgs).

Each update rule accepts current iteration point and the gradient of the object function and produces the next iteration point. Each update rule has the same interface:

```
def update(x, dx, config=None):
```

The inputs are as follows:

1. x: A numpy array giving the current iteration point.

- 2. dx: A numpy array of the same shape as x giving the gradient of the object function with respect to x.
- 3. config: A dictionary containing hyper-parameter values such as learning rate, momentum, etc. If the update rule requires caching values over many iterations, then config will also hold these cached values.

The outputs are as follows:

- 1. next_x: The next point after the update.
- 2. config: The config dictionary to be passed to the next iteration of the update rule.

The update rules are called in Solver class like this:

```
objfcn, grads = self.model.objfcn(self.X)
for p, x in self.model.params.iteritems():
    dx = grads[p]
    config = self.optim_configs[p]
    next_x, next_config = self.update_rule(x, dx, config)
    self.model.params[p] = next_x
    self.optim_configs[p] = next_config
```

3.4 Solver

We provide a class named Solver which bridges the model class and the optimization methods. The solver class is used to solve the maximum likelihood estimation of various time series models. To construct an instance of the solver class, you need the following inputs:

- 1. **solver**: an instance of the model class. Most importantly, given parameters of the model and the data, it can return the negative log-likelihood of the data and the gradients with respect to the parameters.
- 2. data: an numpy 2d array which specifies the data we obtain. The size of the numpy 2d array is $N \times T$, where N is the number of time series we get and T is the length of each time series.
- 3. kwargs: a list of command line arguments. They include
 - (a) update_rule: a string used to specify the optimization method to use. It could be either sgd or sgd_momentum.
 - (b) optim_config: a dictionary used to specify the hyper-parameters for the optimization methods. For example, the key can be a string like learning_rate and the value can be 1e-5.
 - (c) num_epochs: an integer number used to specify the number of epochs for the optimization process
 - (d) batch_size: an integer number used to specify the number of examples for one step of optimization process.
 - (e) print_every: an integer number used to specify how often the program displays the information of the loss value.

For an example of usage, see the following.

```
from tsap.solver import Solver
    from tsap.model import AR, MA
    from tsap.ts_gen import ar1_gen
3
    lag = 1
    sigma = 2.0
    intercept = 0.1
    phi = np.random.randn(lag, 1)
    AR_model = AR(lag=lag, phi=phi, sigma=sigma, intercept=intercept)
    # use the Solver to solve the AR model
    AR_solver = Solver(AR_model, Y,
11
    update_rule='sgd',
    optim_config={
13
    'learning_rate': 3e-5,},
   num_epochs=3000, batch_size=1,
15
    print_every=100)
    AR_solver.train()
    AR_model.params
  The output, as expected, is
    the loss is 424.110945
    the loss is 399.237624
    the loss is 378.012857
    the loss is 357.765501
    the loss is 337.557537
    the loss is 317.542124
    the loss is 299.271549
    the loss is 285.752707
    the loss is 278.916407
    the loss is 276.705150
    the loss is 276.152286
    the loss is 276.012074
    the loss is 275.972341
    {'intercept': array([-0.12022489]),
14
    'phi': array([[ 0.47215311]]),
    'sigma': array([ 0.96874162])}
16
```

For more examples. please refer to the demos.

3.5 Inference

The methods that perform statistical inference is written in inference.py, which consists of five functions, i.e., acovf, acf, BL_stat, yule_walker, and ar_select. In the following, we give a detailed introduction of the usages of these functions. Firstly, acovf and acf returns the auto-covariance and auto-correlations of one path of time series. The input include

- 1. x: an one dimensional numpy array.
- 2. demean: a boolean variable that tells whether to subtract the mean.
- 3. nlags: maximum order of lags in autocorrelation.
- 4. fft: boolean, whether use fast Fourier transform.

The output is an array of auto-covariance or auto-correlation. To use these function, we can run

```
auto_cov = acovf(x, demean=True, fft=False)
2 auto_cor = acf(x, nlags = 40, demean=True, fft=False)
```

Secondly, BL_stat performs Box-Ljung test that tests if the time series data are correlated. The input are acf_array, which is an array of auto-correlation coefficients (ACF), order, which is the maximum order of ACF used, and nobs, which is the time length that is used to compute the ACF. The outputs are the test statistic, a float number, the p-value, also a float number, and the test result, which a boolean. To use this function, an example is

```
test_stat, pval, test_result = BL_stat(acf_array, order, nobs):
```

Finally, yule_walker and ar_select are specified for the AR(p) model. In specific, ar_select takes an array of time series as input and outputs the estimated order of the autoregresive model. Whereas yule_walker takes the time series and the order as input and outputs the estimated parameters using Yule Walker method. The usage will be clear from the following example.

```
import tsap.inference as inference
from tsap.ts_gen import ar1_gen

ts = ar1_gen(0.3, sigma = 1.0, time = 500, num = 1, burnin = 2000)
order_est = inference.ar_select(ts)
print("The estimated order of AR model is "+ str(order_est))
rho, sigma = inference.yule_walker(ts, order = 1)
print("The estimated model parameter is "+ str(rho))

The output is
>>The estimated order of AR model is 2
>>The estimated model parameter is [ 0.2973149]
```

3.6 Trading Strategy

The functions for trading is written in trading.py, which consists of four functions, i.e., signal_generation, profit_loss, trade, and rolltrade. In the following, we give a detailed introduction of the usage of these functions.

Firstly, signal_generation returns the trading signal, like what time to buy and what time to sell. The input includes

- 1. X: a one dimensional numpy array, which is the price series.
- 2. window: a positive integer that specifies the largest length of one trade. If window=5, that means there's at most one trade activity within 5 days. The default number is 2.

The output is a one dimensional numpy array, whose elements are 1 (buying signal), -1 (selling signal), 0 (no signal). To use these function, we can run

```
import trading
2 signal = trading.signal_generation(X, window = 5)
```

Secondly, signal_generation returns the profit based on given trading signals. The input includes

- 1. X: a one dimensional numpy array, which is the price series.
- 2. signal: a one dimensional numpy array that contains trading signals. Usually, it takes the output of signal_generation.
- 3. money: a positive float number, which is the initial wealth. The default number is 1.

The output is a one dimensional numpy array, which is the time series of profit at different time points. To use these function, we can run

```
import trading
2 signal = trading.signal_generation(X, window = 5)
profit = trading.profit_loss(Y, signal, money = 100)
```

Next, trade returns the profit in certain time period. Specifically, given a fitted model, it uses this model to forecast future returns based on historical returns, which are obtained by transfering historical prices to historical returns. After that, it again transfers the returns to prices and then generates trading signal based on predicted prices. At last it calculates the profit based on the real price series and the trading signals. The input includes

- 1. X: a one dimensional numpy array, which is historical price.
- 2. Y: a one dimensional numpy array, which is future prices.
- 3. M: a class from model.py, which is a fitted model.
- 4. nstep: an positive integer, which specifies how many steps to predict.
- 5. window: an positive integer that specifies the largest length of one trade. If window=5, that means there's at most one trade activity within 5 days.
- 6. money: a positive float number, which is the initial wealth. The default number is 1.0.

The output is a one dimensional numpy array, which is the time series of profit at different time points. To use these function, we can run

```
import trading
import model

AR_model = model.AR(lag=3, phi=np.array([[1],[1],[1]]), sigma=1, intercept
=0.1)
profit = trading.trade(X, Y, M = AR_model, nstep = 10, window = 5, money =
100)
```

Lastly, rolltrade returns the profit, trading signal, and the predicted prices by rolling trading in the whole time period, which is an extension of trade. It will repeat the following procedure until the data is used up. Specifically, given a fitted model, it uses this model to forecast future returns based on historical returns, which are obtained by transfering historical prices to historical returns. After that, it again transfers the returns to prices and then generates trading signal based on predicted prices. At last it calculates the profit based on the real price series and the trading signals. The input includes

- 1. X: a one dimensional numpy array, which is historical price.
- 2. M: a class from model.py, which is a fitted model.

- 3. 1: an integer number, which specifies how many data points are used to train for each trading.
- 4. nstep: an positive integer, which specifies how many steps to predict.
- 5. window: an positive integer that specifies the largest length of one trade. If window=5, that means there's at most one trade activity within 5 days.
- 6. money: a positive float number, which is the initial wealth. The default number is 1.0.

The output is three one-dimensional numpy array, which are the time series of profit, trading signal, and the predicted prices at different time points. To use these function, we can run

```
# import library
2 import trading
  import model
4 import numpy as np
  import data_processor as dp
  # read data and do the preprocessing
8 data = np.loadtxt("../data/GOOG.csv", delimiter=',')
 X = np.array([data[0:100]])
10 Y = dp.get_return(X)
12 # initialize the model
  lag = 5
14 sigma = 1.0
  intercept = 0.1
16 phi = np.array([[ 0.04560256],[ 0.0535601 ],[-0.78190871],[ 1.30062633],[
     0.46616754]])
  AR_model = AR(lag=lag, phi=phi, sigma=sigma, intercept=intercept)
  # solve the model
20 _, grads = AR_model.loss(Y)
  solver = Solver(AR_model, Y, update_rule='sgd_momentum', optim_config={'
     learning_rate': 1e-6,}, num_epochs=10000, batch_size=1,print_every=10)
22 solver.train()
24 # get the trading profit, signal and predicted price
  profit, signal, out_pred_price = trading.rolltrade(X, M = AR_model, l = 100,
      nstep = 20, window = 5, money = 100)
```

3.7 Option Pricing

The class OptionPricing calculates the call option price of an underlying stock based on the Black-Scholes model.

The following parameters are needed to construct an instance of the OptionPricing class:

- 1. sigma: the volatility of the underlying stock price, which is the standard deviation of the stock's returns.
- 2. K: the strike price of the option.
- 3. T: the expiry time of the option.
- 4. r: the risk-free interest rate.

5. Smax: the maximum stock price we want to consider.

The method solve_black_scholes calculates the option pricing of an instance, given the grid size of price and time (nS and nt) as input parameters. The option price as an function of stock price and time will be stored in the instance.

After running solve_black_scholes, we can use get_option_price method to get the option price of given underlying stock price S and time t.

Following is a usage example of the OptionPricing class:

```
goog = np.genfromtxt("../data/GOOG.csv", delimiter=",")
sigma = np.std((goog[1:] - goog[:-1]) / goog[:-1])

option_price = OptionPricing(sigma=sigma, T=90, K=800, r=0.005, Smax=1200)
option_price.solve_black_scholes(nS=100, nt=300)
print(option_price.get_option_price(S=810, t=30))
```

3.8 Clustering

We also provide realizations of four clustering methods: K-means, spectral clustering, hierarchical clustering, and Gaussian mixture modeling. These methods are organized by the Cluster class, which is initialized by an input data matrix of dimension (nsample × nfeature) where nsample is the number of data points and nfeature is the number of features. An object in the class consists of three attributes: X, which stores the data matrix, _nsample, and _nfeature. All clustering functions in this class is has the form

```
centroid, labels, clusters =Clustering_method(self, nClusters, maxIter)
```

Here Clustering_method is one of Kmeans, H_clustering, Spectral, and Gaussian_mixture. In addition, nClusters is the number of clusters and maxIter is the maximum number of iterations, which has default value 300. The output consists of three items. centroid, a nClusters × _nfeatures matrix storing the centers of clusters; labels is the predicted labels; clusters stores the index of data points in each cluster. The following example shows how to use our package to cluster the S&P 500 dataset. First we import and standardize the data.

```
import numpy as np
# read SP500 data

SP500 = np.genfromtxt('../data/SP500array.csv', delimiter=',')
SP500 = SP500.T

nStock = len(SP500[:,0])
nTime = len(SP500[0,:])

# preprocessing, standardize data

X = np.copy(SP500)
for i in range(nStock):
    X[i,:] = (X[i,:] - np.mean(X[i,:]))/np.std(X[i,:])
```

The we import our package and fit an K-means clustering model on the data with 3 clusters.

```
1 from src.cluster import Cluster
  model = Cluster(X)
3 # run K-means
5 import time
```

```
7 start = time.time()
  centroid, labels, clusters = model.kMeans(nClusters = 5)
9 end = time.time()

11 print("K-means takes "+str(end-start)+" seconds")

The output is
1 K-means takes 0.873986959457 seconds
```

3.9 Model Reduction

When multiple time series are available, we can analysis the data with modell reduciton methods. The model reduction function is provided in reduction.py. In this file, we provide a class Reduction.

```
class Reduction(object):
    """Callable modal reduction object.

Example usage:
    xreduction = Reduction(X), X shape [n_features, n_samples], make sure X is zero-mean

xmean, ux, at, energy_content = xreduction.PCA(n_components=3)
"""
```

The member functions of class Reduction consists of Principal Component Analysis (PCA)

- [2], Independent Component Analysis (ICA) [1], and Dynamic Mode Decomposition (DMD)
- [3] [4]. The definitions of three functions are declared below:

```
def PCA(self, n_components=None):
2
          Principal component analysis (PCA) of data in matrix
          n_components: integer, number of principal components
          Returns:
          ux: principal components
          at: principal components coefficients
          energy_content: energy content percentage in the principal
     components
10
      def ICA(self, n_components, gfunc='logcosh', tol=1e-4, max_iter=200):
12
          Independent component analysis(ICA) of data in matrix X
          n_components: integer, number of independent components
16
          gfunc: string, 'logcosh' or 'exp', default 'logcosh', Non-gaussian
     function
          tol: float, tolerance of iteration, default 1e-4
18
          max_iter: integer, maximum iteration steps, default 200
          Returns:
          Ex: array, mean of data
          T: array [n_features, n_features], whitening matrix, st, xtilde = Tx
          A: array [n_features, n_components], mixing matrix, st, xtilde = As
          W: array [n_components, n_features], orthogonal rows, unmixing
24
     matrix, st, W = inv(A), s = W*xtilde
```

```
S: array, [n_components, n_samples], source data, st, S = W*Xtilde

"""

def DMD(self, n_components=None):

"""

Dynamic mode decomposition(DMD) of time series data x(k), find

square

matrix A such that x(k+1) = Ax(k). Find eigendecomposition of A, and

corresponding DMD modes, and DMD eigenvalues.

"""
```

To get a better sense of how to analyse real data with this class. Now we will give an example using S&P 500.

```
1 import numpy as np
  import reduction
  # preprocessing, subtract each stock price mean and normalize by std
5 x = np.copy(SP500)
  for i in range(nStock):
      x[i,:] = (x[i,:] - np.mean(x[i,:]))/np.std(x[i,:])
9 # read SP500 data
  SP500 = np.genfromtxt('../data/SP500array.csv', delimiter=',')
11 \text{ SP500} = \text{SP500.T}
_{13} # call and test PCA
  n_{components} = 5
15 SPreduction = reduction.Reduction(x)
  ux, at, energy_content = SPreduction.PCA(n_components)
  # call and test ICA
19 \text{ n\_components} = 10
  SPreduction = reduction.Reduction(x)
21 Ex, T, A, W, S = SPreduction.ICA(n_components)
_{23} # call and test DMD
  n_{components} = 20
25 SPreduction = reduction.Reduction(x)
  evals, modes, energy_content = SPreduction.DMD(n_components)
```

References

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