

q3

May 24, 2020

```
[1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.metrics import mean_squared_error
from sklearn.preprocessing import StandardScaler
from skfda.representation.basis import BSpline
import rpy2.robjects as robjects
from multiprocessing import Process, Queue
from IPython.core.display import display, HTML
display(HTML("<style>.container { width:100% !important; }</style>"))
```

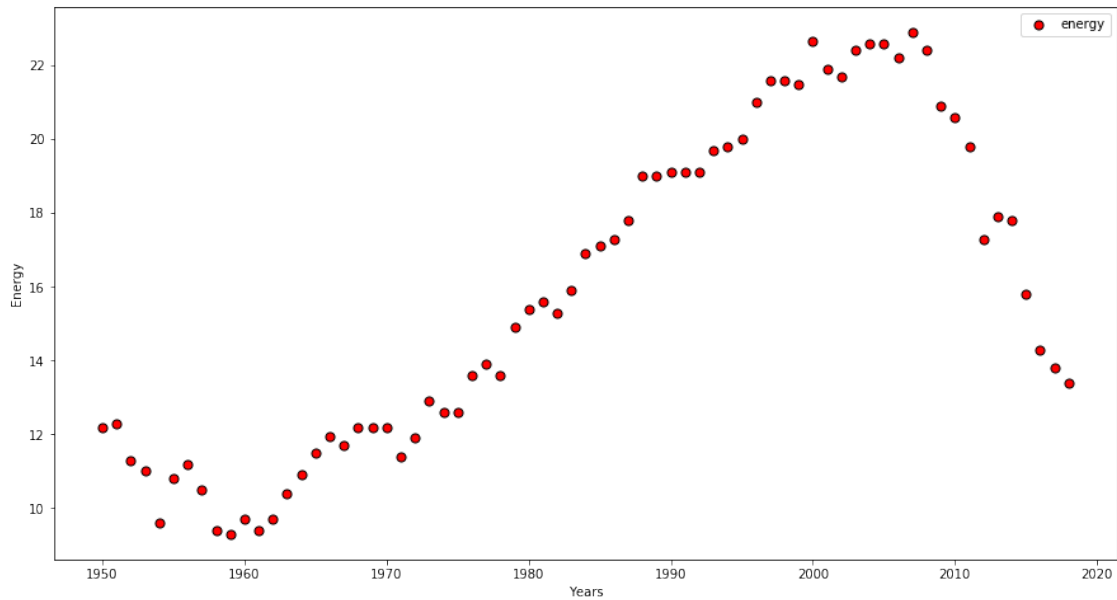
<IPython.core.display.HTML object>

0.0.1 Read in the Data

```
[2]: df = pd.read_csv('P04.csv', header=None)
```

0.0.2 View the Data

```
[3]: fig, axes = plt.subplots(figsize=(15,8))
ax=plt.subplot()
df.plot(kind='scatter', x=0, y=1, ax=ax, color='r', edgecolor='k', s=50,
        label='energy')
plt.xlabel('Years')
plt.ylabel('Energy')
plt.show()
```



0.0.3 Separate x and y vectors

```
[4]: X = df[0].values
     y = df[1].values
```

0.0.4 Cubic Splines Implementation

This implementation is a python recreation of the R example code. It utilizes OLS to find the weight vector

```
[5]: def cubic_splines(x, y, **kwargs):
      """
      inputs
      -----
      x      -> Domain vector (independent variables)
      y      -> Response vector (dependent variables)
      kwargs -> should have any of the following options
                  knots --> [list]: list of knots
                  num_knots --> [int]: number of desired knots. These knots will
      ↳ be equidistant and calculated from input x

      outputs
      -----
      B      -> Weight matrix found from OLS
      yhat   -> Fitted data
```

```

makeH -> Function needed to transform new unseen data
'''
if 'knots' in kwargs.keys():
    assert isinstance(kwargs['knots'], list), 'knots should be a list'
    knots = np.array(kwargs['knots'])

elif 'num_knots' in kwargs.keys():
    assert isinstance(kwargs['num_knots'], int), 'num_knots should be an
→integer'
    knots = np.linspace(min(x), max(x), kwargs['num_knots'])

'''
This is a closure function that is a returned object from this function so
→a user can transform new unseen independent variables
after initially fitting the training data with the supplied knots
'''
def wrapper(knots):
    def inner(x):
        basis = []

        h1 = 1
        h2 = x
        h3 = x**2
        h4 = x**3
        basis = [h1, h2, h3, h4]
        for i in range(len(knots)):
            h = (x - knots[i])**3
            if h <= 0:
                h = 0
            basis.append(h)

        H = np.array(basis)
        return H
    return inner

#create a transform function to be returned
makeH = wrapper(knots)

x = np.array(x)
y = np.array(y)
knots = np.array(knots)

#1st basis function is all ones

```

```

h1 = np.ones(len(x))

#2nd basis function is linear
h2 = x

#3rd basis function is quadratic
h3 = x**2

#4th basis function is cubic
h4 = x**3

basis = [h1, h2, h3, h4]

#get the truncated power basis for  $\sigma_{ki} < x < \sigma_{ki+1}$ 
for i in range(len(knots)):
    h = (x - knots[i])**3
    h[h <= 0] = 0

    basis.append(h)

#transformation matrix
H = np.array(basis)

#weight vectors found from OLS
B = np.dot(np.linalg.pinv(np.dot(H, H.T)), np.dot(H,y))

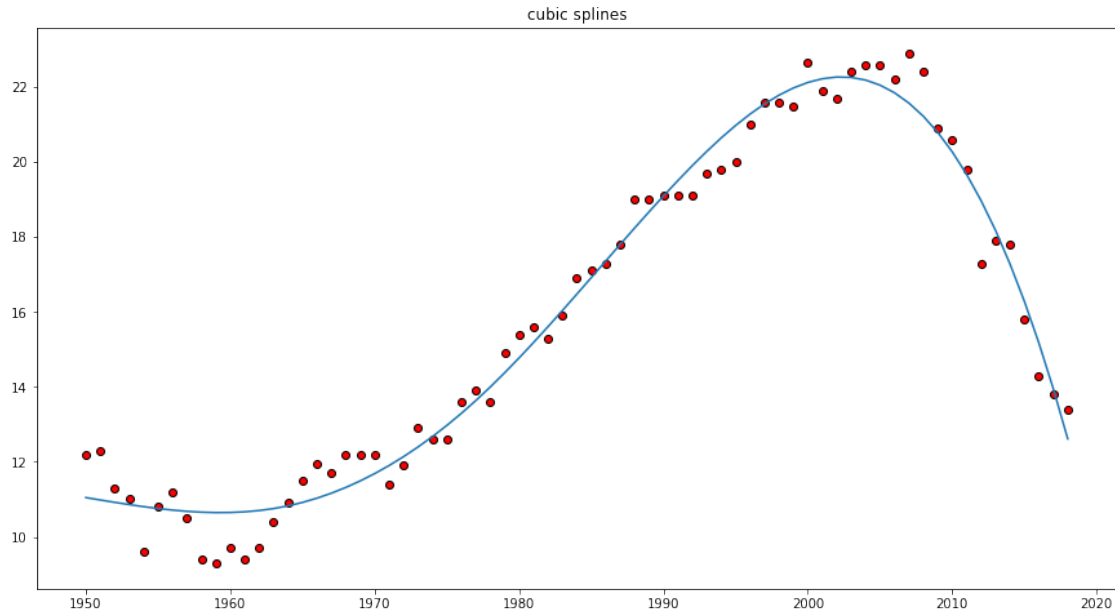
#return weight vector, the fitted data, and the transformation function to
→transform new unseen data
return B , np.dot(H.T,B), makeH

```

0.0.5 Test out cubic splines on the data with 6 knots

```
[6]: b, yhatcs, transformfunc = cubic_splines(X,y,num_knots=6)
```

```
[7]: plt.figure(figsize=(15,8))
plt.scatter(X, y , color='r', edgecolor='k')
plt.plot(X, yhatcs)
plt.title('cubic splines')
plt.show()
```



0.0.6 Cubic Splines MSE Function

The `cubic_mse` function can be ran as a single process but also supports a multiprocessing Queue object for multiprocessing of a hyperparameter gridsearch. It utilizes leave one out cross validation and aggregates the error for each left out independent variable predicted response. The errors from each LOO is then used to calculate the overall MSE. If a multiprocessing Queue object is supplied, the result is put on the queue and returned to the main process, otherwise the MSE is simply returned

```
[8]: def cubic_mse(x,y,num_knots, q=None):
    """
    inputs
    -----
    x          -> the independent variable vector
    y          -> the dependent variable vector
    num_knots -> the number of knots to be used for the MSE calculation
    q          -> multiporcessing Queue object

    returns
    -----
    mse -> the mse for all errors for LOOCV
    """
    #initialize empty array to hold all errors
    preds = np.zeros(len(y))
    y_true = np.zeros(len(y))
```

```

#perform LOOCV
for i in range(len(y)):
    X = np.delete(x, i)
    Y = np.delete(y, i)

    #fit data on all independent variables except the value left out
    weights, fit, transform_func = cubic_splines(X, Y, num_knots=num_knots)

    #predict on the left out value
    xtest = x[i]
    xtest = transform_func(xtest)
    pred = np.dot(weights, xtest)

    #calculate error
    preds[i] = pred
    y_true[i] = y[i]

#calculate mse
mse = mean_squared_error(y_true, preds)

#return mse
if q == None:
    return mse
else:
    q.put(mse)

```

0.0.7 b-splines implementation

The b-splines implementation was created by using the `skfda.representation.basis` library and the `BSpline` object was utilized. The function takes in the independent and dependent variables along with the desired polynomial degree and returns the weight vector, the fitted data, and the object needed to transform new unseen data

```

[9]: def bspline(x, y, order, **kwargs):
    """
    inputs
    -----
    x      -> The independent variable vector
    y      -> the dependent variable vector
    order  -> the degree of polynomia
    kwargs -> should have any of the following options
               knots --> [list]: list of knots
               num_knots --> [int]: number of desired knots. These knots will be
    ↪ equidistant and calculated from input x

```

```

outputs
-----
B      -> Weight matrix found from OLS
yhat   -> Fitted data
bss    -> Object needed to transform new unseen data into an H tensor
'''

if 'knots' in kwargs.keys():
    assert isinstance(kwargs['knots'], (list, np.ndarray)), 'knots should be a list'
    knots = np.array(kwargs['knots'])

elif 'num_knots' in kwargs.keys():
    assert isinstance(kwargs['num_knots'], int), 'num_knots should be an integer'
    knots = np.linspace(min(x), max(x), kwargs['num_knots'])

#instantiate a b-splines object
bss = BSpline(knots=knots, order=order)

#create the b-spline transformed tensors
H = bss.evaluate(x)

#find the weight vectors using OLS
B = np.dot(np.linalg.pinv(np.dot(H, H.T)), np.dot(H,y))

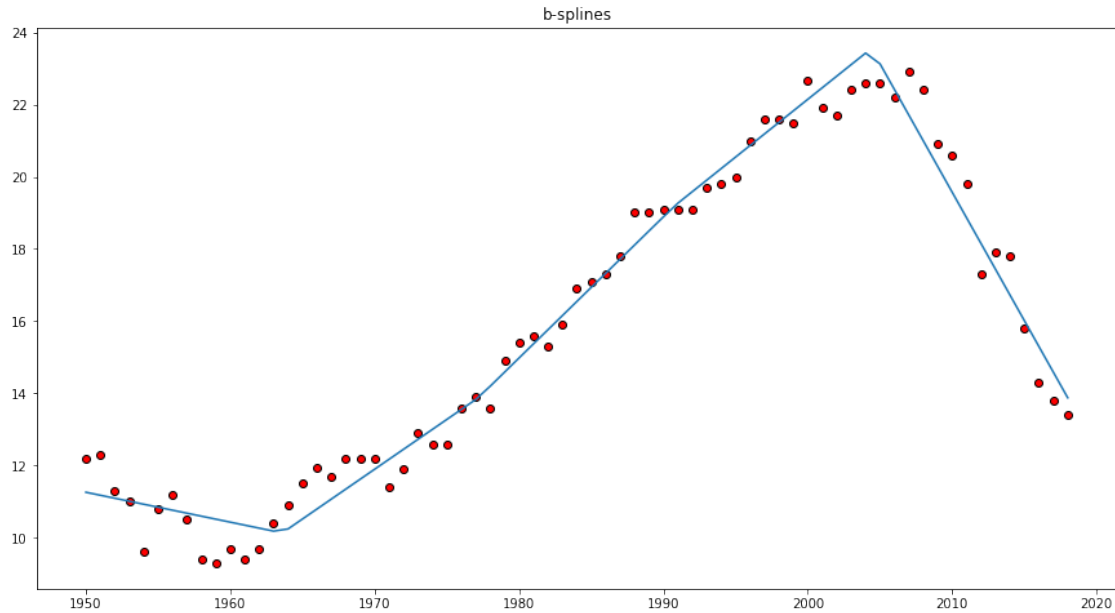
return B, np.dot(H.T,B), bss

```

0.0.8 Test out b-splines with 6 knots and 2nd order polynomial

```
[10]: weights, yhatbs, transform_object = bspline(X, y, 2, num_knots=6)
```

```
[11]: plt.figure(figsize=(15,8))
plt.scatter(X,y, color='r', edgecolor='k')
plt.plot(X, yhatbs)
plt.title('b-splines')
plt.show()
```



0.0.9 b-Spline MSE Function

The `bspline_mse` function can be ran as a single process but also supports a multiprocessing Queue object for multiprocessing of a hyperparameter gridsearch. It utilizes leave one out cross validation and aggregates the error for each left out independent variable predicted response. The errors from each LOO is then used to calculate the overall MSE. If a multiprocessing Queue object is supplied, the result is put on the queue and returned to the main process, otherwise the MSE is simply returned

```
[12]: def bspline_mse(x, y, order, num_knots, q= None):
    """
    inputs
    -----
    x          -> the independent variable vector
    y          -> the dependent variable vector
    order      -> polynomial degree
    num_knots  -> the number of knots to be used for the MSE calculation
    q          -> multiporcessing Queue object

    returns
    -----
    mse -> the mse for all errors for LOOCV
    """
    #initialize error array
    preds = np.zeros(len(y))
    y_true = np.zeros(len(y))
```



```

#perform LOOCV
for i in range(len(y)):
    X = np.delete(x, i)
    Y = np.delete(y, i)

    #fit the data on all independent variables except the left out value
    weights, fit, transform_object = bspline(X, Y, order=order,
    ↪ num_knots=num_knots)

    xtest = x[i]
    xtest = transform_object.evaluate([xtest])
    pred = np.dot(weights, xtest)

    preds[i] = pred
    y_true[i] = y[i]

#calculate mse
mse = mean_squared_error(y_true, preds)

if q == None:
    return mse
else:
    q.put(mse)

```

0.0.10 Smoothing Spline implementation

The smoothing spline implmentation using the rpy2 library which allows for the direct usage of R function in python. The code is a Python implementation of the R example code provided. A closure function is created so new predictions can be made on unseen data with respect to the fit

```

[13]: def smoothing_splines(x, y, spar):
    '''
    inputs
    -----
    x -> independent variables
    y -> dependent variables
    spar -> smoothing parameter

    outputs
    -----
    yhat -> the fitted values
    pred_obj -> function used to predict on newe unseen data

    '''

```

```

#get the smooth.spline function used by r
smooth_spline = robjects.r['smooth.spline']

#conver x and y to r objects
x = robjects.FloatVector(list(x))
y = robjects.FloatVector(list(y))

#fit the data
spline_obj = smooth_spline(x=x, y=y, spar=float(spar))

'''
This closure function is returned so predictions can be made on new unseen
data.
'''
def wrapper(spline_obj):
    def inner(x):
        x = robjects.FloatVector(list(x))
        yspline = robjects.r['predict'](spline_obj,x).rx2('y')
        return np.array(yspline)
    return inner

#create the prediction function
predict_obj = wrapper(spline_obj)

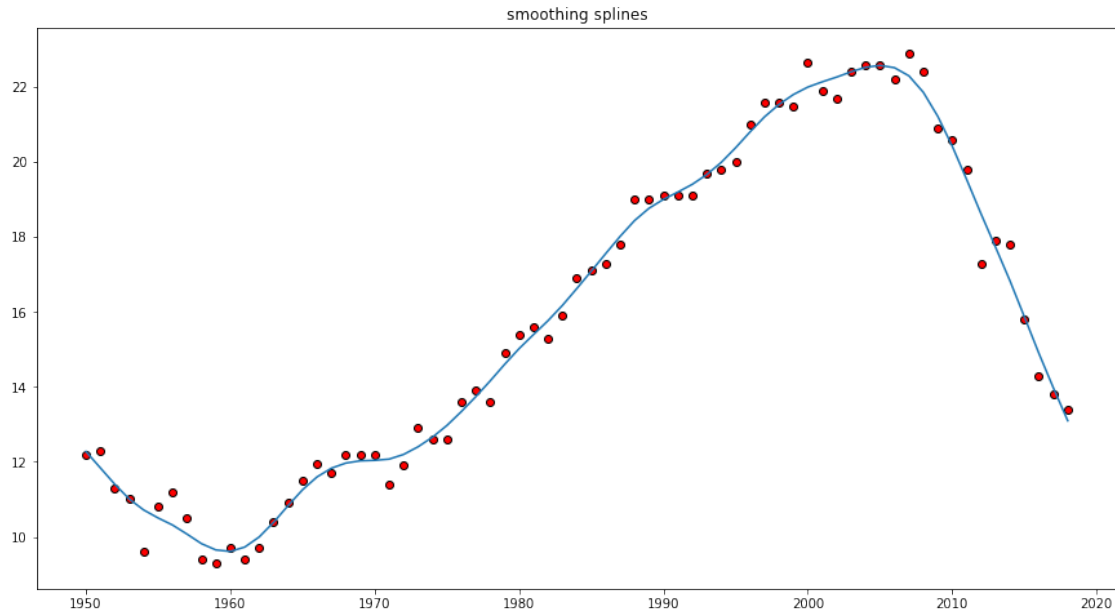
#return yhat and prediction function
return np.array(spline_obj.rx2('y')) , predict_obj

```

0.0.11 test out smoothing splines with a smoothing parameter of 0.5

```
[14]: yhatss, predict_obj = smoothing_splines(x=X, y=y, spar=0.5)
```

```
[15]: plt.figure(figsize=(15,8))
plt.scatter(X,y,color='r', edgecolor='k')
plt.plot(X,yhatss)
plt.title('smoothing splines')
plt.show()
```



0.0.12 Smoothing Spline MSE Function

The `smoothing_mse` function can be ran as a single process but also supports a multiprocessing Queue object for multiprocessing of a hyperparameter gridsearch. It utilizes leave one out cross validation and aggregates the error for each left out independent variable predicted response. The errors from each LOO is then used to calculate the overall MSE. If a multiprocessing Queue object is supplied, the result is put on the queue and returned to the main process, otherwise the MSE is simply returned

```
[16]: def smoothing_mse(x, y, spar, q=None):
    """
    inputs
    -----
    x          -> the independent variable vector
    y          -> the dependent variable vector
    spar       -> smoothing parameter
    q          -> multiporcessing Queue object

    returns
    -----
    mse -> the mse for all errors for LOOCV
    """
    #initialize error array
    preds = np.zeros(len(y))
    y_true = np.zeros(len(y))
```

```

#perform LOOCV
for i in range(len(y)):
    X = np.delete(x, i)
    Y = np.delete(y, i)

    #fit data except left out independent variable
    fit, predict_object = smoothing_splines(X, Y, spar)

    #predict on left out value
    xtest = x[i]
    pred = predict_object([xtest])

    preds[i] = pred
    y_true[i] = y[i]

#calculate mse
mse = mean_squared_error(y_true, preds)

if q == None:
    return mse
else:
    q.put(mse)

```

0.0.13 Gaussian kernel implementation

The gaussian kernel function is a Python refactor of the R example code provided - It creates a closure function use for real time evaluation of independent variables

```

[17]: #kernel function
def kerf(z):
    kernel = np.exp(-z*z / 2) / np.sqrt(2*np.pi)
    return kernel

def gaussian_kernel(x,y,lambd_):
    '''
    inputs
    -----
    x          -> the independent variable vector
    y          -> the dependent variable vector
    lambda_    -> lambda value for kernel function

    returns
    -----
    func -> evaluation function for unseen independent variables
    '''

```

```

'''

def wrapper(x,y,lambda_):
    def inner(xtest):
        if not isinstance(xtest, (list, np.ndarray)):
            xtest = [xtest]
        N = len(xtest)
        f = np.zeros(N)

        for k in range(N):
            z = kerf((xtest[k] - x) / lambda_)
            f[k] = np.sum(z * y) / np.sum(z)

        return f
    return inner

func = wrapper(x, y, lambda_)

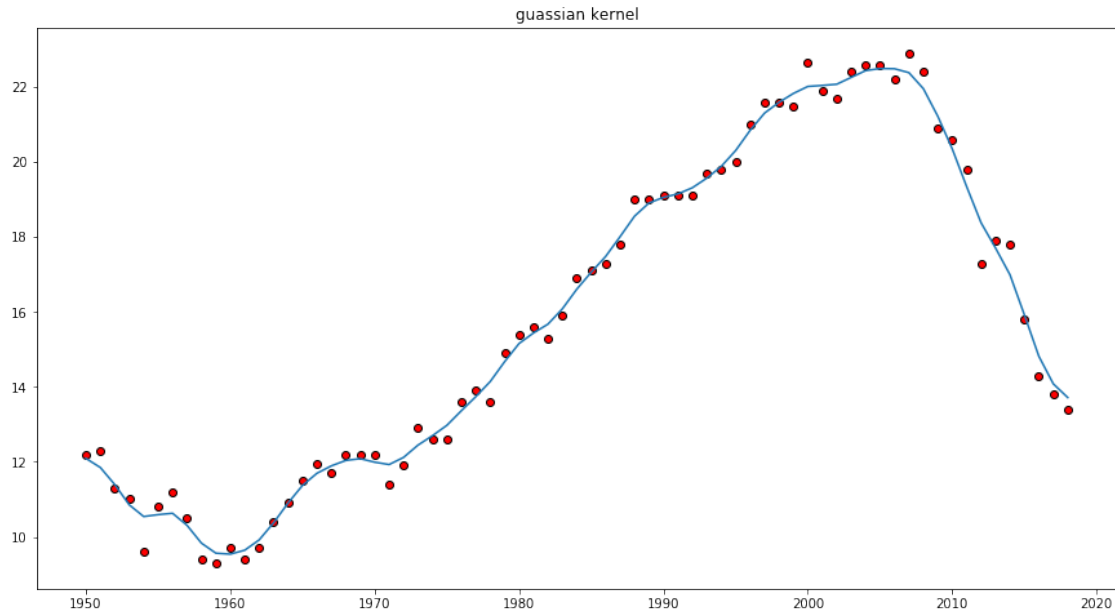
return func

```

0.0.14 Test out gaussian kernel with a lambda value of 1.2

```
[18]: gk = gaussian_kernel(X,y,1.2)
```

```
[19]: plt.figure(figsize=(15,8))
plt.scatter(X,y,color='r', edgecolor='k')
plt.plot(X, gk(X))
plt.title('gaussian kernel')
plt.show()
```



0.0.15 Gaussian Kernel MSE Function

The `gaussian_mse` function can be ran as a single process but also supports a multiprocessing Queue object for multiprocessing of a hyperparameter gridsearch. It utilizes leave one out cross validation and aggregates the error for each left out independent variable predicted response. The errors from each LOO is then used to calculate the overall MSE. If a multiprocessing Queue object is supplied, the result is put on the queue and returned to the main process, otherwise the MSE is simply returned

```
[20]: def gaussian_mse(x, y, lambda_, q=None):
    """
    inputs
    -----
    x          -> the independent variable vector
    y          -> the dependent variable vector
    lambda_    -> lambda parameter
    q          -> multiporcessing Queue object

    returns
    -----
    mse -> the mse for all errors for LOOCV
    """
    preds = np.zeros(len(y))
    y_true = np.zeros(len(y))

    for i in range(len(y)):
```

```

X = np.delete(x, i)
Y = np.delete(y, i)

xtest = x[i]
pred_obj = gaussian_kernel(x,y,lambda_)

pred = pred_obj(xtest)

preds[i] = pred
y_true[i] = y[i]

mse = mean_squared_error(y_true, preds)

if q == None:
    return mse
else:
    q.put(mse)

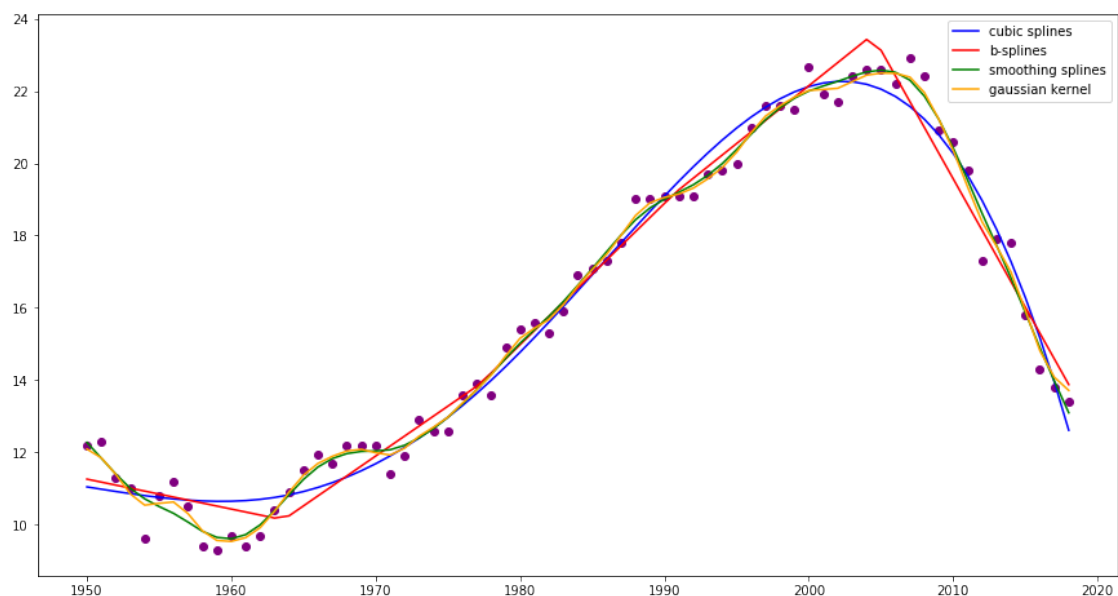
```

0.0.16 Viewing all tests together

```

[21]: plt.figure(figsize=(15,8))
plt.scatter(X,y, color='purple')
plt.plot(X,yhatcs, color='blue', label='cubic splines')
plt.plot(X,yhatbs, color='red', label='b-splines')
plt.plot(X,yhatss, color='green', label='smoothing splines')
plt.plot(X,gk(X), color='orange', label='gaussian kernel')
plt.legend()
plt.show()

```



0.1 LOOCV Implementation

This function utilizes multiprocessing and performs grid search across an array of hyperparameters for each algorithm. It returns a dictionary for each algorithm containing the best hyperparameter and the subsequent MSE value from the LOOCV

```
[22]: def LOOCV(x,y):  
    '''  
    inputs  
    -----  
    x -> independent variable  
    y -> dependent variable  
  
    outputs  
    -----  
    results -> dictionary containing results for each algorithm  
    '''  
  
    #range of knots to grid search over (knots low = 6, knots high = 15)  
    knots = range(6,16)  
    #polynomial orders to grid search over (order low = 1, order high = 5)  
    orders = range(1,6)  
    #smoothing parameters to grid search over (spar low = 0, spar high = 1)  
    spars = np.linspace(0,1,100)  
    #lambda parameters to grid search over (lambda low = 1, lambda high = 4)  
    lambdas_ = np.arange(0.1,4,0.1)  
  
    #initialize empty array to hold all calculated MSEs for each hyperparameter  
    ↪grid searched over  
    cs_mse = np.zeros(len(knots))  
    bs_mse = np.zeros(len(knots) * len(orders))  
    ss_mse = np.zeros(len(spars))  
    gk_mse = np.zeros(len(lambdas_))  
  
    #create a multiprocessing Queue object for each process to write results to  
    cs_q = Queue()  
    bs_q = Queue()  
    ss_q = Queue()  
    gk_q = Queue()  
  
    #list to maintain PID for all created processes  
    cs_process = []  
    bs_process = []  
    ss_process = []
```



```

gk_process = []

#cubic mse multiprocessing
for i in knots:
    p = Process(target=cubic_mse, args=(x, y, i, cs_q))
    p.start()
    cs_process.append(p)

#join results from each cubic process
for pos, p in enumerate(cs_process):
    p.join()
    cs_mse[pos] = cs_q.get()

#b-spline mse multiprocessing
for i in knots:
    for j in orders:
        p = Process(target=bspline_mse, args=(x, y, j, i, bs_q))
        p.start()
        bs_process.append(p)

#join results from each b-spline process
for pos, p in enumerate(bs_process):
    p.join()
    bs_mse[pos] = bs_q.get()

#smoothing splines mse multiprocessing
for i in spars:
    p = Process(target=smoothing_mse, args=(x,y,i,ss_q))
    p.start()
    ss_process.append(p)

#join results from each smoothing spline process
for pos, p in enumerate(ss_process):
    p.join()
    ss_mse[pos] = ss_q.get()

#gaussian mse multiprocessing
for i in lambdas_:
    p = Process(target=gaussian_mse, args=(x,y,i, gk_q))
    p.start()
    gk_process.append(p)

#join results from each gaussian process
for pos, p in enumerate(gk_process):

```

```

p.join()
gk_mse[pos] = gk_q.get()

#plot results
fig, axes = plt.subplots(nrows=2, ncols=2, figsize=(18,10))

ax1 = plt.subplot(221)
ax1.plot(knots, cs_mse)
ax1.scatter(knots[np.argmin(cs_mse)], cs_mse[np.argmin(cs_mse)],
↪color='red', s=50)
ax1.set_xlabel('knots')
ax1.set_ylabel('mse')
ax1.set_title('cubic splines')

ax2 = plt.subplot(222)
points = [(k,o) for k in knots for o in orders]
xp = [x[0] for x in points]
yp = [x[1] for x in points]
ax2.scatter(xp,yp, s=bs_mse * 300)
ax2.scatter(xp[np.argmin(bs_mse)], yp[np.argmin(bs_mse)], color='r',
↪s=bs_mse[np.argmin(bs_mse)]*300)
plt.text(xp[np.argmin(bs_mse)], yp[np.argmin(bs_mse)] + 0.5, 'Min MSE = {}'.
↪format(round(bs_mse[np.argmin(bs_mse)],2)), color='green')
ax2.set_xlabel('knots')
ax2.set_ylabel('polynomial order')
ax2.set_title('b-splines')

ax3 = plt.subplot(223)
ax3.plot(spars, ss_mse)
ax3.scatter(spars[np.argmin(ss_mse)], ss_mse[np.argmin(ss_mse)], color='r',
↪s=50)
ax3.set_xlabel('spars')
ax3.set_ylabel('mse')
ax3.set_title('Smoothing Splines')

ax4 = plt.subplot(224)
ax4.plot(lambdas_, gk_mse)
ax4.scatter(lambdas_[np.argmin(gk_mse)], gk_mse[np.argmin(gk_mse)],
↪color='r', s=50)
ax4.set_xlabel('lambda')
ax4.set_ylabel('mse')
ax4.set_title('Gaussian Kernel')

plt.subplots_adjust(left=None, bottom=None, right=None, top=None,
↪wspace=None, hspace=0.3)

```

```

plt.show()

results = {
    'cubic_splines': {'knot': knots[np.argmin(cs_mse)], 'mse':
→cs_mse[np.argmin(cs_mse)]},
    'bsplines': {'knot': xp[np.argmin(bs_mse)], 'order': yp[np.
→argmin(bs_mse)], 'mse': bs_mse[np.argmin(bs_mse)]},
    'smooth_splines': {'spar': spars[np.argmin(ss_mse)], 'mse':
→ss_mse[np.argmin(ss_mse)]},
    'gaussian_kernel': {'lambda': lambdas_[np.argmin(gk_mse)], 'mse':
→gk_mse[np.argmin(gk_mse)]}
}

results_string = '''
Cubic Splines:      Optimal Knots {}
                   MSE {}

B-Splines:          Optimal Knots {}
                   Optimal Degree {}
                   MSE {}

Smoothing Splines: Optimal Smoothing Parameter {}
                   MSE {}

Gaussian Kernel:    Optimal Lambda {}
                   MSE {}

''' .format(results['cubic_splines']['knot'],
            results['cubic_splines']['mse'],
            results['bsplines']['knot'],
            results['bsplines']['order'],
            results['bsplines']['mse'],
            results['smooth_splines']['spar'],
            results['smooth_splines']['mse'],
            results['gaussian_kernel']['lambda'],
            results['gaussian_kernel']['mse'])

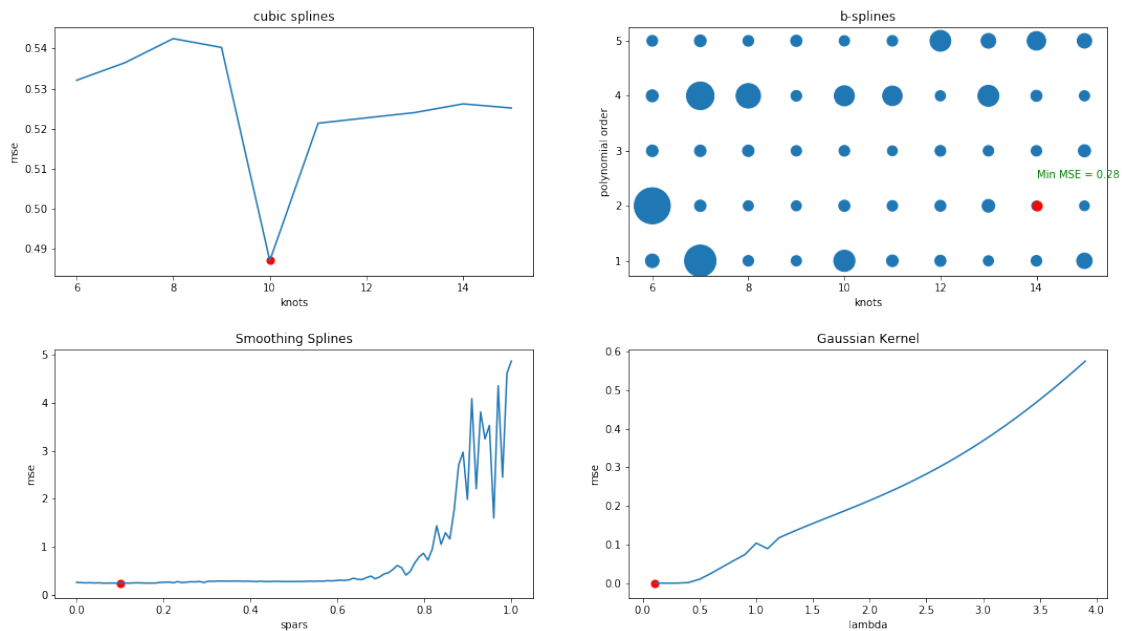
print(results_string)

return results

```

0.1.1 Perform LOOCV on Unscaled X data First

```
[23]: results = LOOCV(X,y)
```



Cubic Splines: Optimal Knots 10
 MSE 0.4869035180512312

B-Splines: Optimal Knots 14
 Optimal Degree 2
 MSE 0.2757741596820151

Smoothing Splines: Optimal Smoothing Parameter 0.101010101010102
 MSE 0.24488285318203684

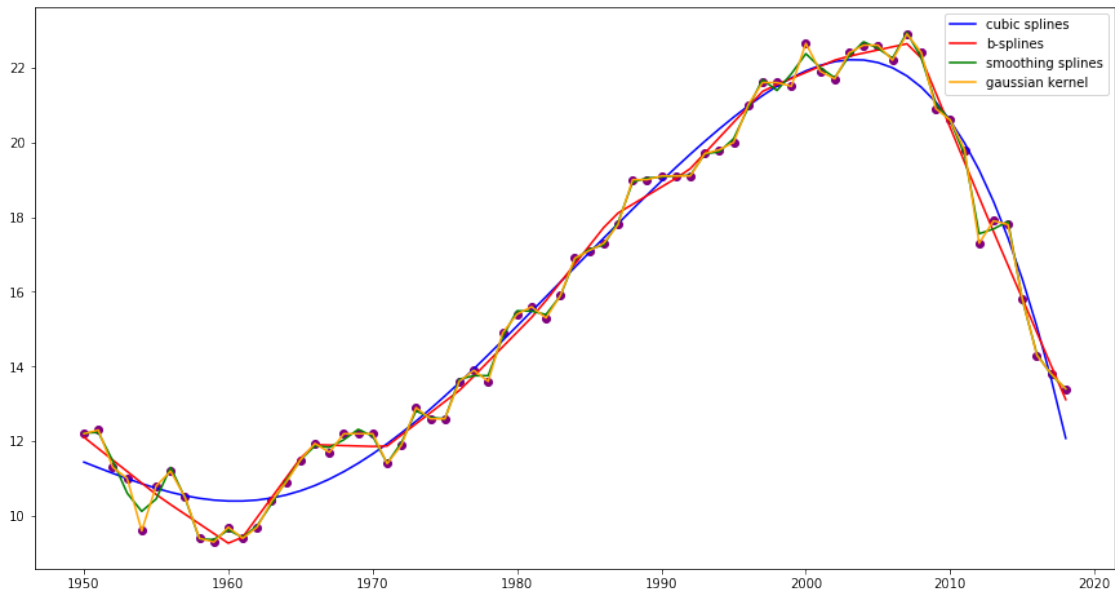
Gaussian Kernel: Optimal Lambda 0.1
 MSE 1.0060834733253483e-30

0.1.2 Utilizing best parameters for each algorithm

```
[24]: cs_knots = results['cubic_splines']['knot']  
bs_knots = results['bsplines']['knot']  
bs_degree = results['bsplines']['order']  
ss_spar = results['smooth_splines']['spar']  
gk_lambda = results['gaussian_kernel']['lambda']
```

```
[25]: b, yhatcs, transformfunc = cubic_splines(X,y,num_knots=cs_knots)  
weights, yhatbs, transform_object = bspline(X, y, bs_degree, num_knots=bs_knots)  
yhatss, predict_obj = smoothing_splines(x=X, y=y, spar=ss_spar)  
gk = gaussian_kernel(X,y,gk_lambda)
```

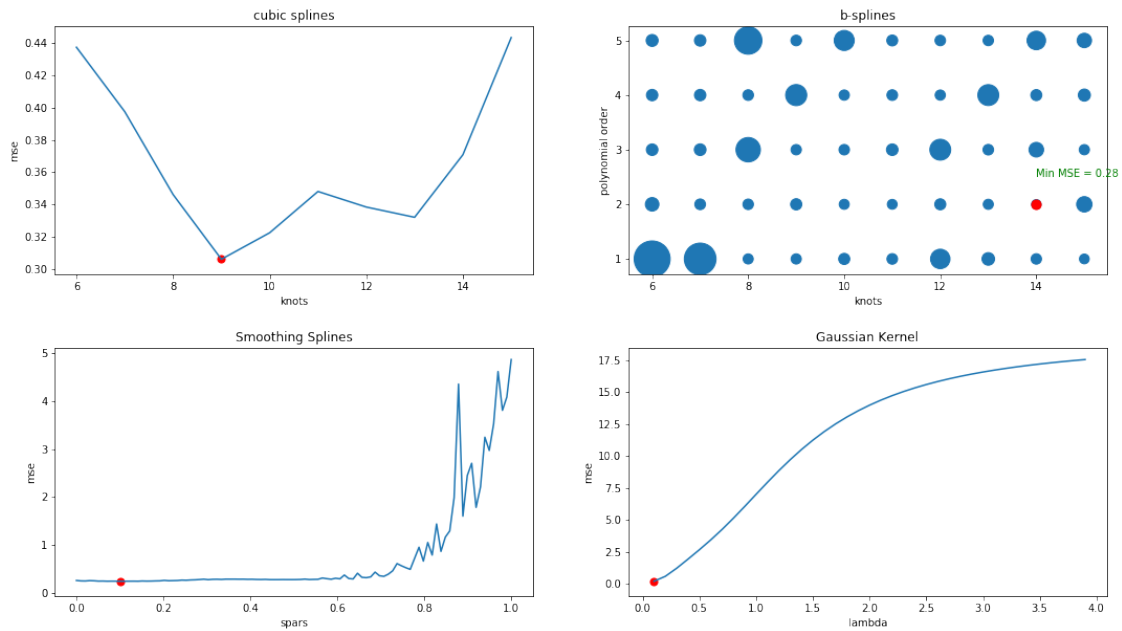
```
[26]: plt.figure(figsize=(15,8))  
plt.scatter(X,y, color='purple')  
plt.plot(X,yhatcs, color='blue', label='cubic splines')  
plt.plot(X,yhatbs, color='red', label='b-splines')  
plt.plot(X,yhatss, color='green', label='smoothing splines')  
plt.plot(X,gk(X), color='orange', label='gaussian kernel')  
plt.legend()  
plt.show()
```



0.1.3 Perform LOOCV on Scaled Data

```
[27]: scaler = StandardScaler()
scaler.fit(X.reshape(-1, 1))
X_transformed = scaler.transform(X.reshape(-1, 1))
X_transformed = np.array([i[0] for i in X_transformed.tolist()])
```

```
[28]: results = LOOCV(X_transformed,y)
```



Cubic Splines: Optimal Knots 9
MSE 0.30610506353725286

B-Splines: Optimal Knots 14
Optimal Degree 2
MSE 0.27577415968201885

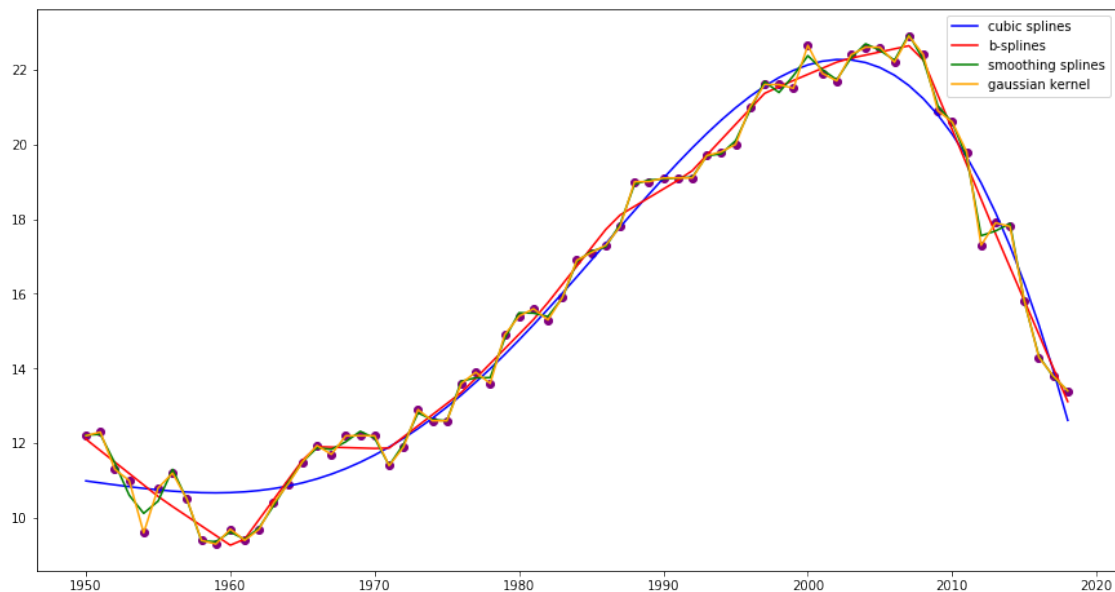
Smoothing Splines: Optimal Smoothing Parameter 0.10101010101010102
MSE 0.24488285318203865

Gaussian Kernel: Optimal Lambda 0.1
MSE 0.21302888604096862

```
[29]: cs_knots = results['cubic_splines']['knot']
bs_knots = results['bsplines']['knot']
bs_degree = results['bsplines']['order']
ss_spar = results['smooth_splines']['spar']
gk_lambda = results['gaussian_kernel']['lambda']
```

```
[30]: b, yhatcs, transformfunc = cubic_splines(X,y,num_knots=cs_knots)
weights, yhatbs, transform_object = bspline(X, y, bs_degree, num_knots=bs_knots)
yhatss, predict_obj = smoothing_splines(x=X, y=y, spar=ss_spar)
gk = gaussian_kernel(X,y,gk_lambda)
```

```
[31]: plt.figure(figsize=(15,8))
plt.scatter(X,y, color='purple')
plt.plot(X,yhatcs, color='blue', label='cubic splines')
plt.plot(X,yhatbs, color='red', label='b-splines')
plt.plot(X,yhatss, color='green', label='smoothing splines')
plt.plot(X,gk(X), color='orange', label='gaussian kernel')
plt.legend()
plt.show()
```



```
[32]: fig, axes = plt.subplots(nrows=2, ncols=2, figsize=(20,15))

ax1 = plt.subplot(221)
ax1.scatter(X,y, color='red', edgecolor='k')
ax1.plot(X,yhatcs, color='blue', label='cubic splines')
ax1.set_xlabel('years')
ax1.set_ylabel('energy')
```

```

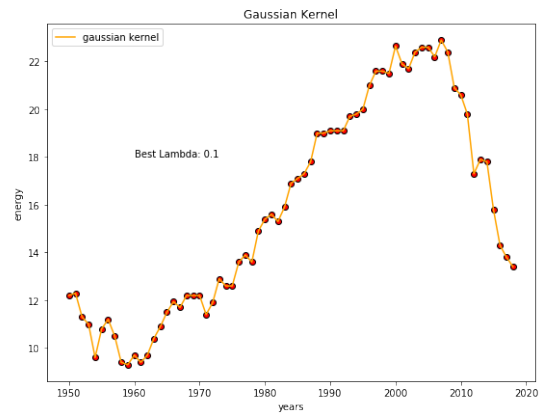
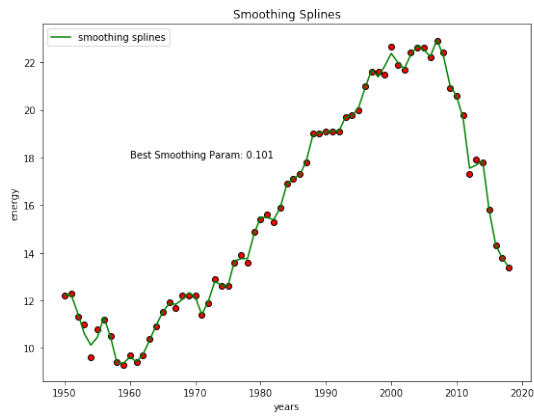
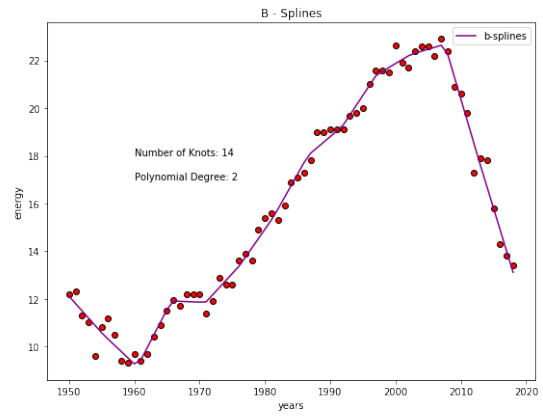
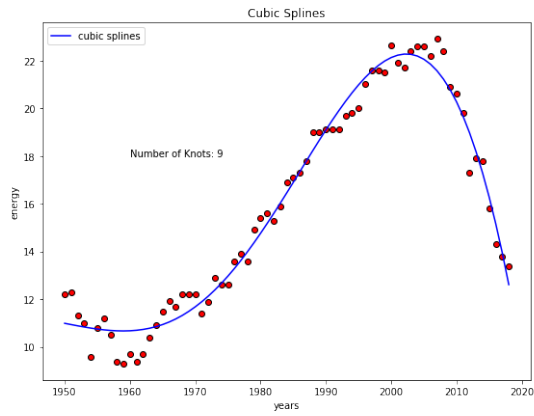
ax1.set_title('Cubic Splines')
plt.text(1960, 18, 'Number of Knots: {}'.format(cs_knots))
plt.legend()

ax2 = plt.subplot(222)
ax2.scatter(X,y, color='red', edgecolor='k')
ax2.plot(X,yhatbs, color='purple', label='b-splines')
ax2.set_xlabel('years')
ax2.set_ylabel('energy')
ax2.set_title('B - Splines')
plt.text(1960, 18, 'Number of Knots: {}'.format(bs_knots))
plt.text(1960, 17, 'Polynomial Degree: {}'.format(bs_degree))
plt.legend()

ax3 = plt.subplot(223)
ax3.scatter(X,y,color='red', edgecolor='k')
ax3.plot(X,yhatss, color='green', label='smoothing splines')
ax3.set_xlabel('years')
ax3.set_ylabel('energy')
ax3.set_title('Smoothing Splines')
plt.text(1960, 18, 'Best Smoothing Param: {}'.format(round(ss_spar,3)))
plt.legend()

ax4 = plt.subplot(224)
ax4.scatter(X,y, color='red', edgecolor='k')
ax4.plot(X,gk(X), color='orange', label='gaussian kernel')
ax4.set_xlabel('years')
ax4.set_ylabel('energy')
ax4.set_title('Gaussian Kernel')
plt.text(1960, 18, 'Best Lambda: {}'.format(gk_lambda))
plt.legend()
plt.show()

```

[]: