

COMP0114 Inverse Problems in Imaging. Coursework 1

1. Solving Underdetermined Problems

a.)

I define a function `phi` to calculate Φ .

```
def phi(x, p):  
    return np.sum(np.power(np.abs(x), p))
```

Just follow the definition of Φ .

b.)

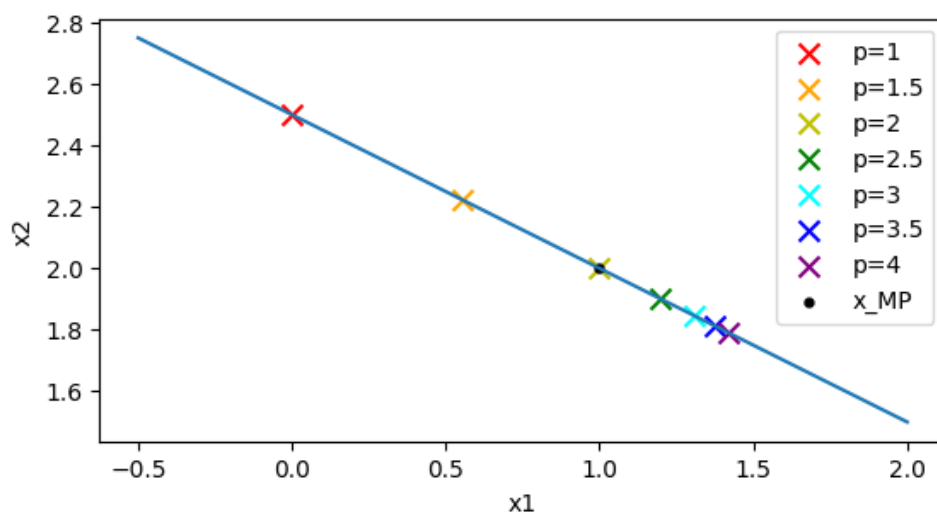
I use `scipy.optimize` to solve the problems.

```
def solve(p):  
    A = np.array([[1, 2]])  
    b = 5  
    cons = (  
        {  
            'type': 'eq',  
            'fun': lambda x: A @ x - b  
        })  
    res = optimize.minimize(phi, np.zeros((A.shape[1],)), args=(p),  
                           constraints=cons, method='SLSQP', options={'disp': False})  
    return res.x, phi(res.x, p)
```

The function takes `p` as the parameter, which represents the p in Φ .

c.)

The resulting plot is as below.



The blue line represents the constraint equation. The meanings of the crosses are as the legend.

d.)

Moore-Penrose generalised inverse corresponds $\bar{p} = 2$. As shown in the plot above, the dot represents x_{MP} is at the same location as $\bar{p} = 2$.

When $\bar{p} = 2$, we can rewrite the problem as:

$$\begin{aligned} \text{minimize } \Phi = x^T x, \quad \text{subject to } Ax = b \\ \text{where } A = (1 \ 2), \ b = 5 \end{aligned}$$

, which is equivalent to the problem:

$$\begin{aligned} \text{minimize } \Phi = x^T x - \lambda^T (Ax - b) \\ \text{where } A = (1 \ 2), \ b = 5 \end{aligned}$$

We can solve it:

$$\begin{aligned} & \begin{cases} \frac{d\Phi}{dx} = 0 \\ \frac{d\Phi}{d\lambda} = 0 \end{cases} \\ & \begin{cases} 2x - A^T \lambda = 0 \\ Ax - b = 0 \end{cases} \\ & \begin{cases} \lambda = 2(AA^T)^{-1}b \\ x = A^T(AA^T)^{-1}b = x_{\text{MP}} \end{cases} \end{aligned}$$

and $A^T(AA^T)^{-1}$ is the A^\dagger .

2. Singular Value Decomposition

a.)

I use `numpy.linspace` to create the spatial grid.

```
def spatial_grid(n):  
    return np.linspace(-1., 1., num=n)
```

b.)

I create a function to calculate the value of Gaussian function in a numpy vector.

```
def gaussian_func(x, n, mu=0., sig=0.2):  
    delta_n = 2. / (n - 1)  
    return delta_n / np.sqrt(2 * np.pi) / sig * np.exp(-(x - mu) * (x - mu) / (2. * sig  
* sig))
```

c.)

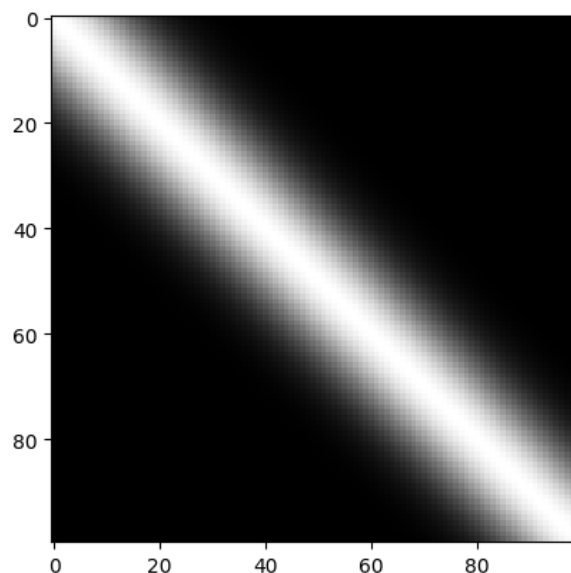
I create a function to generate a convolution matrix.

```
def convolution_matrix(n, mu=0., sig=0.2):  
    x = np.array([spatial_grid(n)])  
    x2d = np.tile(x, (n, 1)).T - x  
    return gaussian_func(x2d, n, mu, sig)
```

Just follow the definition of $A_{i,j}$.

d.)

The plot of the matrix A when $n = 100$ is as below.



e.)

I use `numpy.linalg.svd` to compute SVD.

```
def svd(A):  
    u, s, vt = np.linalg.svd(A, full_matrices=True)  
    return u, s, vt
```

I use two methods to verify $A = U W V^T$ and the results are in the comments.

```
u, s, vt = svd(A)  
print(((u * s) @ vt - A) < 1e-12).all() # True  
print(np.linalg.norm((u * s) @ vt - A)) # 3.865089147759237e-15
```

f.)

I create a function to calculate the pseudoinverse.

```
def pseudoinverse(A):  
    u, s, vt = svd(A)  
    W = np.diag(s)  
    Wi = np.linalg.pinv(W)  
    assert (Wi @ W - np.identity(W.shape[0]) < 1e-12).all()  
    assert (W @ Wi - np.identity(W.shape[0]) < 1e-12).all()  
    return vt.T @ Wi @ u.T
```

I use `numpy.linalg.pinv` to get the W^\dagger , and the assertion is used to check $W W^\dagger = W^\dagger W = I_{dn}$.

For $n = 10$, I write some code to calculate the SVD and verified the generated pseudoinverse. The results are in the comments.

```
n = 10  
A = convolution_matrix(n)  
Ai_np = np.linalg.pinv(A)  
Ai = pseudoinverse(A)  
print(np.linalg.norm(Ai - Ai_np)) # 3.5827277168872394e-15  
print((Ai @ A - np.identity(A.shape[0]) < 1e-12).all() # True  
print((A @ Ai - np.identity(A.shape[0]) < 1e-12).all() # True
```

The pseudoinverse is very similar to the true inverse.

g.)

For $n = 20$, the code and results are as below.

```

n = 20
A = convolution_matrix(n)
Ai_np = np.linalg.pinv(A)
Ai = pseudoinverse(A)
print(np.linalg.norm(Ai - Ai_np)) # 3.651758567089755e-10
print((Ai @ A - np.identity(A.shape[0]) < 1e-10).all()) # True
print((A @ Ai - np.identity(A.shape[0]) < 1e-10).all()) # True

```

Although the pseudoinverse is effective, the accuracy is not as good as the case of $n = 10$.

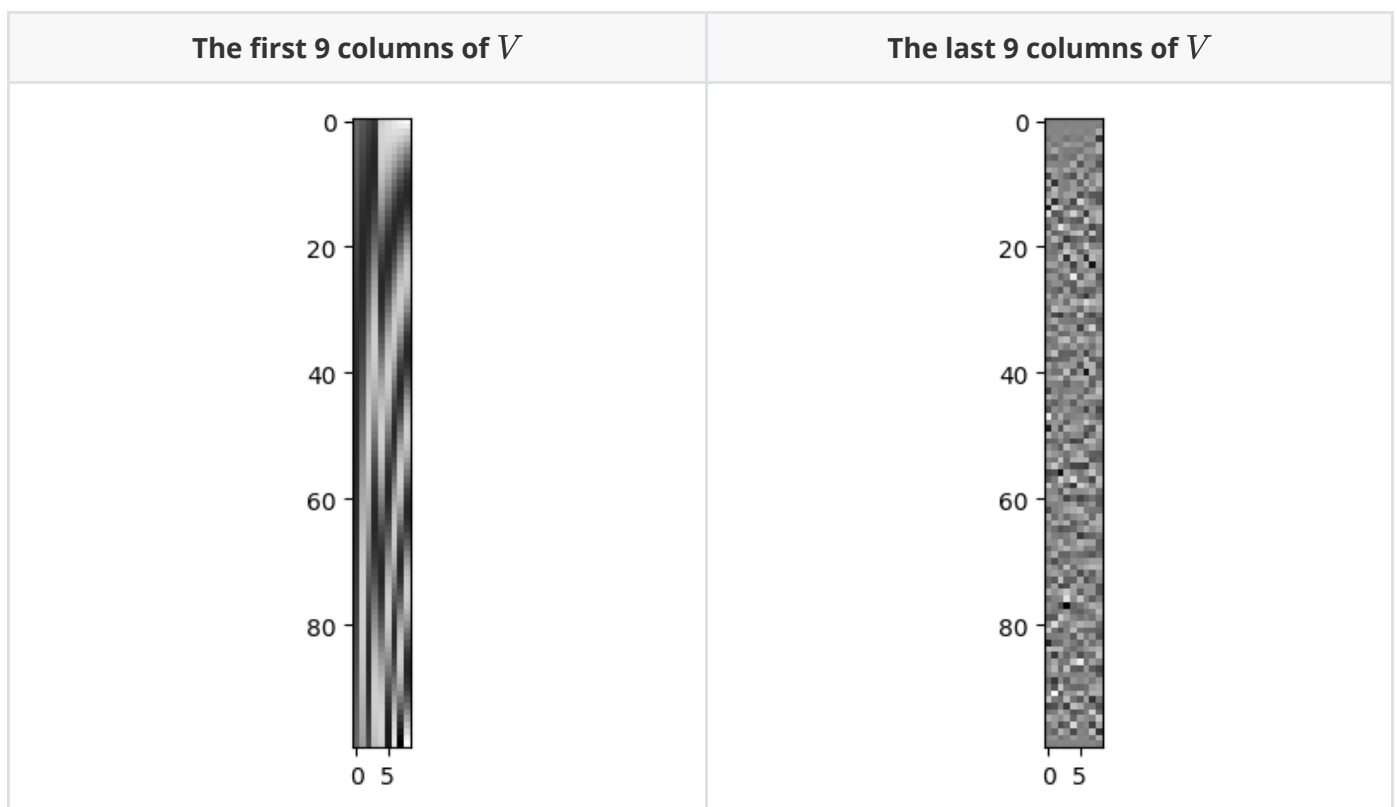
For $n = 100$, the code and results are as below.

```

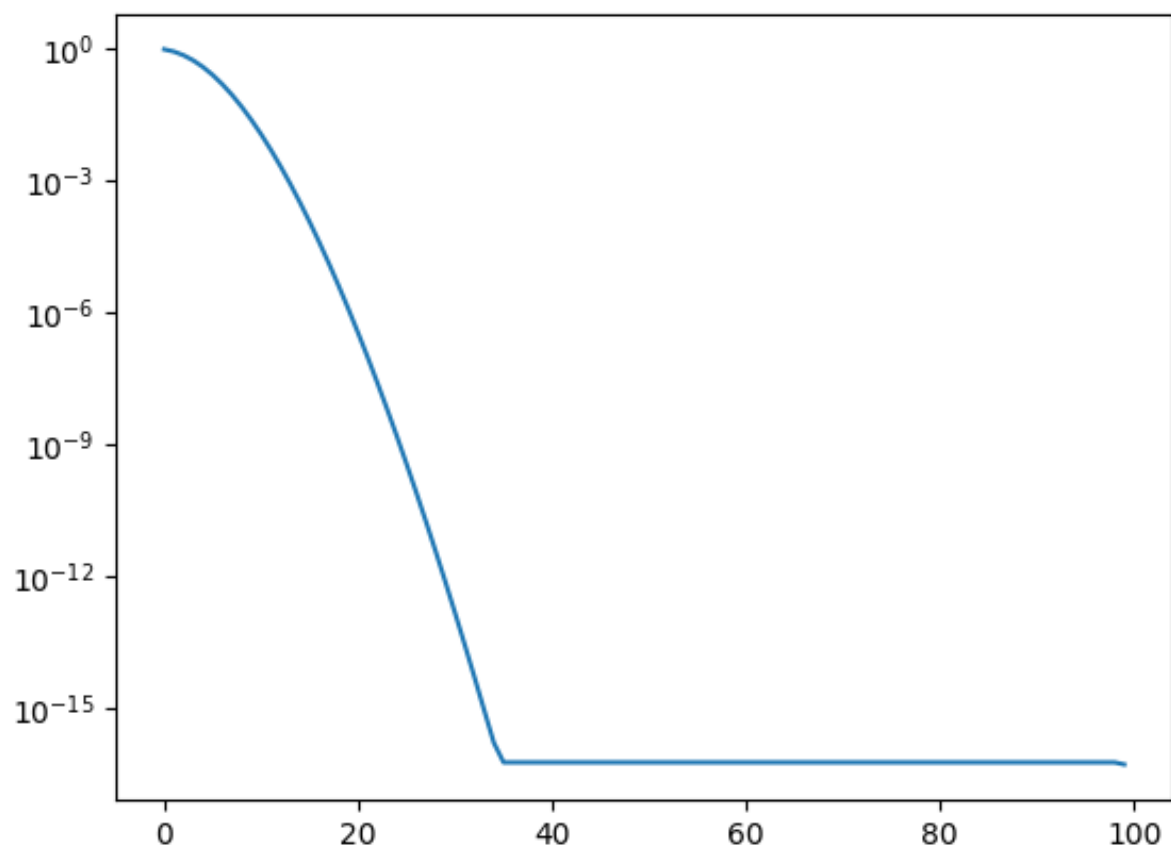
n = 100
A = convolution_matrix(n)
Ai_np = np.linalg.pinv(A)
Ai = pseudoinverse(A)
print(np.linalg.norm(Ai - Ai_np)) # 0.04085877562246467
print((Ai @ A - np.identity(A.shape[0]) < 1).all()) # True
print((A @ Ai - np.identity(A.shape[0]) < 1).all()) # True

```

The effect of pseudoinverse is not so close as the true inverse.



The first 9 columns is more meaningful than the last 9 columns. The corresponding singular values of last 9 columns is probably very small. From the plot of singular values below we can know it is true.



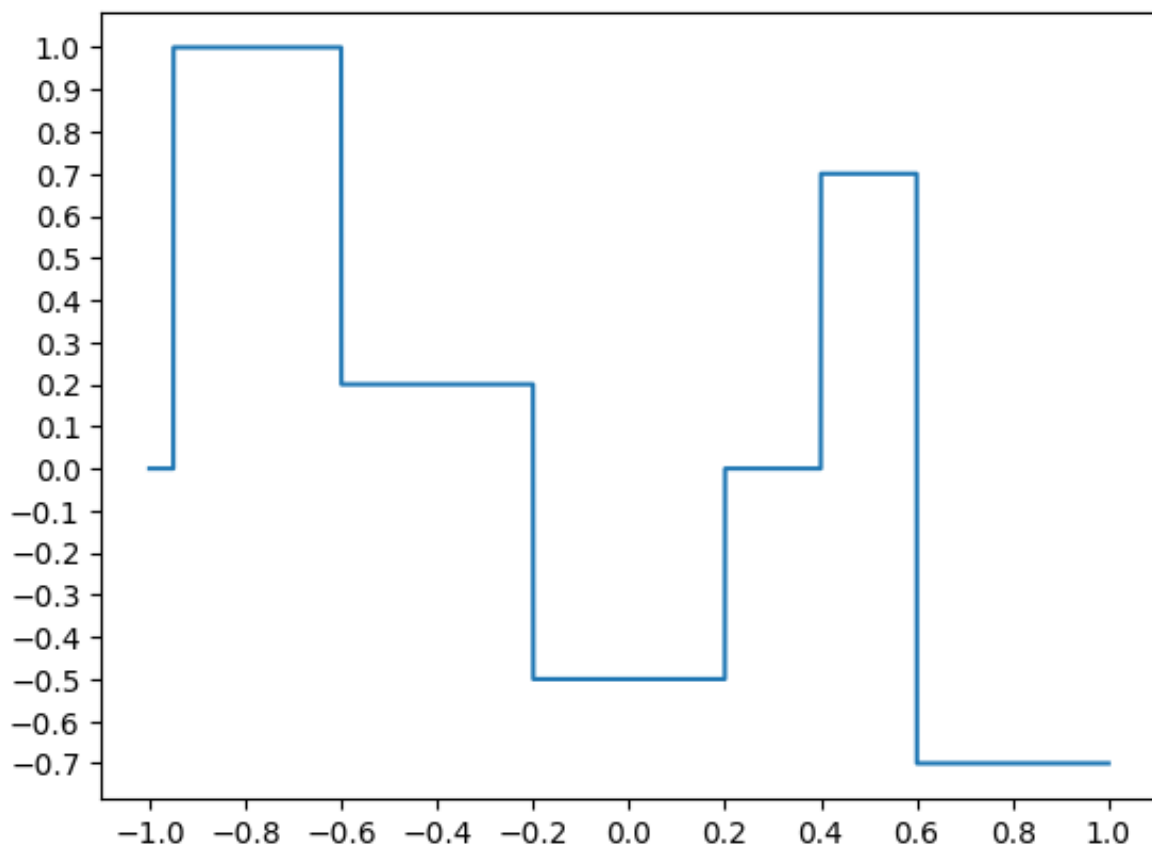
3. Convolutions and Fourier transform

a.)

The function calculating $f(x)$ is as below.

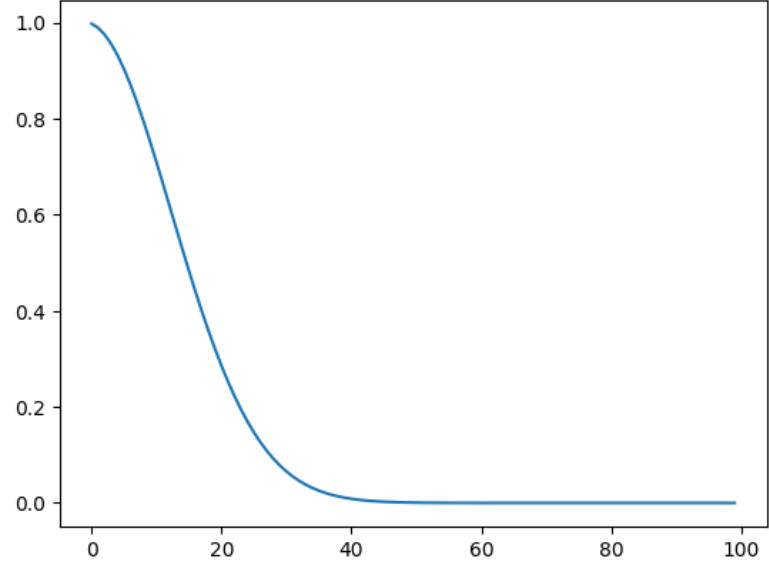
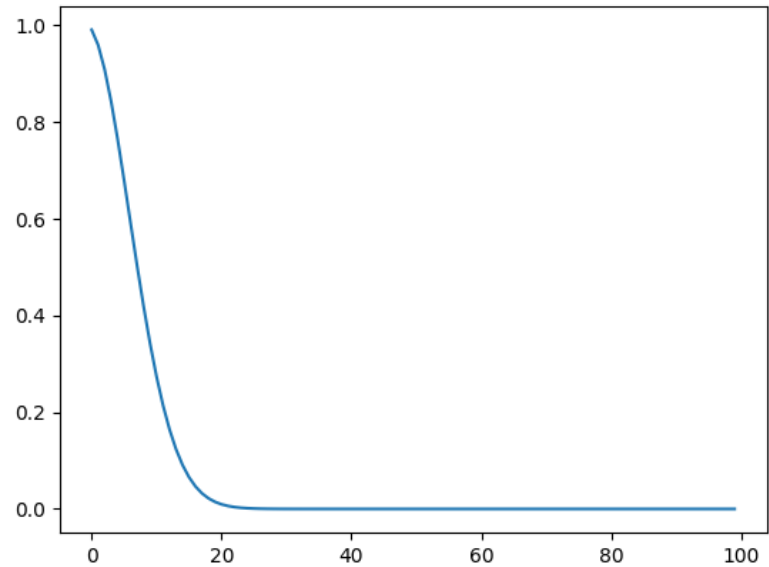
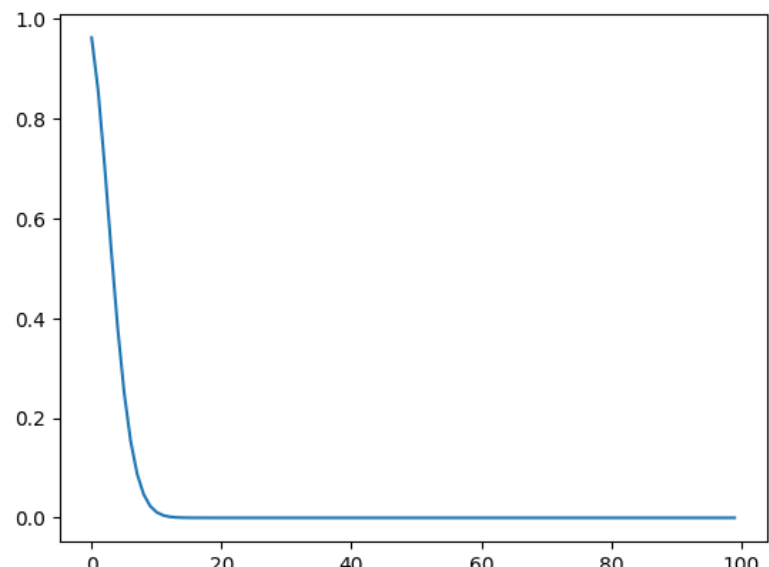
```
def X(x, a, b):  
    xc = x.copy()  
    idx = (a<x)&(x<=b)  
    xc[idx] = 1  
    xc[~idx] = 0  
    return xc  
  
def f(x):  
    return X(x, -0.95, -0.6) + 0.2 * X(x, -0.6, -0.2) - 0.5 * X(x, -0.2, 0.2) + 0.7 *  
    X(x, 0.4, 0.6) - 0.7 * X(x, 0.6, 1)
```

The plot is as below:



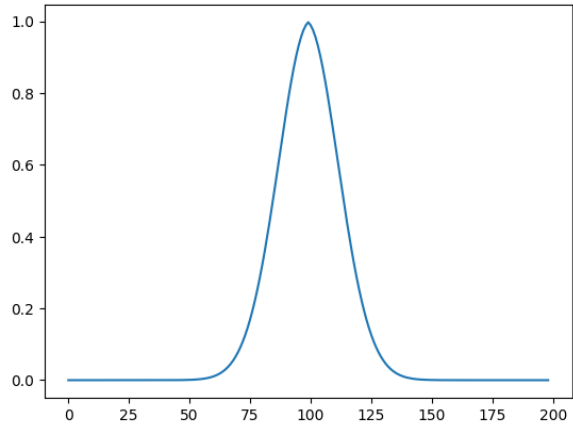
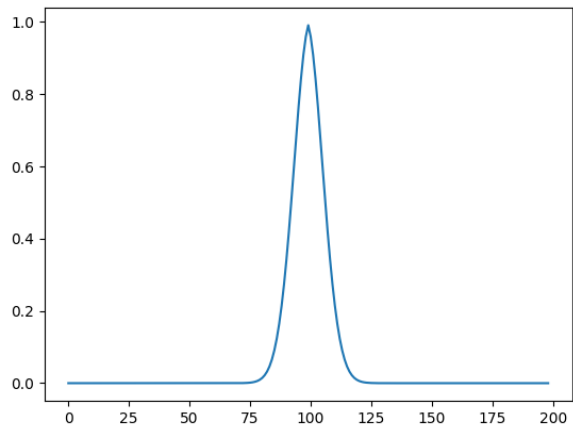
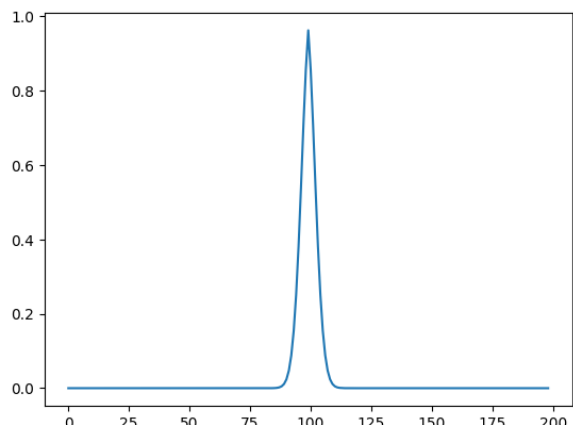
b.)

The plots of singular values:

σ	Plots of Singular Values																												
0.05	 <p>A line plot showing the singular values of a matrix for $\sigma = 0.05$. The x-axis represents the index of the singular value, ranging from 0 to 100 with major ticks every 20 units. The y-axis represents the magnitude of the singular value, ranging from 0.0 to 1.0 with major ticks every 0.2 units. The curve starts at (0, 1.0) and decreases rapidly, reaching approximately 0.4 at x=20, 0.1 at x=40, and then levels off to near zero for x > 60.</p> <table border="1"><thead><tr><th>Index</th><th>Singular Value</th></tr></thead><tbody><tr><td>0</td><td>1.00</td></tr><tr><td>10</td><td>0.75</td></tr><tr><td>20</td><td>0.40</td></tr><tr><td>30</td><td>0.15</td></tr><tr><td>40</td><td>0.05</td></tr><tr><td>50</td><td>0.02</td></tr><tr><td>60</td><td>0.01</td></tr><tr><td>70</td><td>0.01</td></tr><tr><td>80</td><td>0.01</td></tr><tr><td>90</td><td>0.01</td></tr><tr><td>100</td><td>0.01</td></tr></tbody></table>	Index	Singular Value	0	1.00	10	0.75	20	0.40	30	0.15	40	0.05	50	0.02	60	0.01	70	0.01	80	0.01	90	0.01	100	0.01				
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0.1	 <p>A line plot showing the singular values of a matrix for $\sigma = 0.1$. The x-axis represents the index of the singular value, ranging from 0 to 100 with major ticks every 20 units. The y-axis represents the magnitude of the singular value, ranging from 0.0 to 1.0 with major ticks every 0.2 units. The curve starts at (0, 1.0) and decreases rapidly, reaching approximately 0.4 at x=10, 0.1 at x=20, and then levels off to near zero for x > 30.</p> <table border="1"><thead><tr><th>Index</th><th>Singular Value</th></tr></thead><tbody><tr><td>0</td><td>1.00</td></tr><tr><td>10</td><td>0.40</td></tr><tr><td>20</td><td>0.10</td></tr><tr><td>30</td><td>0.02</td></tr><tr><td>40</td><td>0.01</td></tr><tr><td>50</td><td>0.01</td></tr><tr><td>60</td><td>0.01</td></tr><tr><td>70</td><td>0.01</td></tr><tr><td>80</td><td>0.01</td></tr><tr><td>90</td><td>0.01</td></tr><tr><td>100</td><td>0.01</td></tr></tbody></table>	Index	Singular Value	0	1.00	10	0.40	20	0.10	30	0.02	40	0.01	50	0.01	60	0.01	70	0.01	80	0.01	90	0.01	100	0.01				
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0.2	 <p>A line plot showing the singular values of a matrix for $\sigma = 0.2$. The x-axis represents the index of the singular value, ranging from 0 to 100 with major ticks every 20 units. The y-axis represents the magnitude of the singular value, ranging from 0.0 to 1.0 with major ticks every 0.2 units. The curve starts at (0, 1.0) and decreases very rapidly, reaching approximately 0.4 at x=5, 0.1 at x=10, and then levels off to near zero for x > 15.</p> <table border="1"><thead><tr><th>Index</th><th>Singular Value</th></tr></thead><tbody><tr><td>0</td><td>1.00</td></tr><tr><td>5</td><td>0.40</td></tr><tr><td>10</td><td>0.10</td></tr><tr><td>15</td><td>0.02</td></tr><tr><td>20</td><td>0.01</td></tr><tr><td>30</td><td>0.01</td></tr><tr><td>40</td><td>0.01</td></tr><tr><td>50</td><td>0.01</td></tr><tr><td>60</td><td>0.01</td></tr><tr><td>70</td><td>0.01</td></tr><tr><td>80</td><td>0.01</td></tr><tr><td>90</td><td>0.01</td></tr><tr><td>100</td><td>0.01</td></tr></tbody></table>	Index	Singular Value	0	1.00	5	0.40	10	0.10	15	0.02	20	0.01	30	0.01	40	0.01	50	0.01	60	0.01	70	0.01	80	0.01	90	0.01	100	0.01
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100	0.01																												

c.)

I flipped the original image of singular values horizontally and stitched it together with the original image to create a new image. The shapes look like Gaussian functions. The variances can be calculated by the largest singular values, which are the centre values of the Gaussian functions.

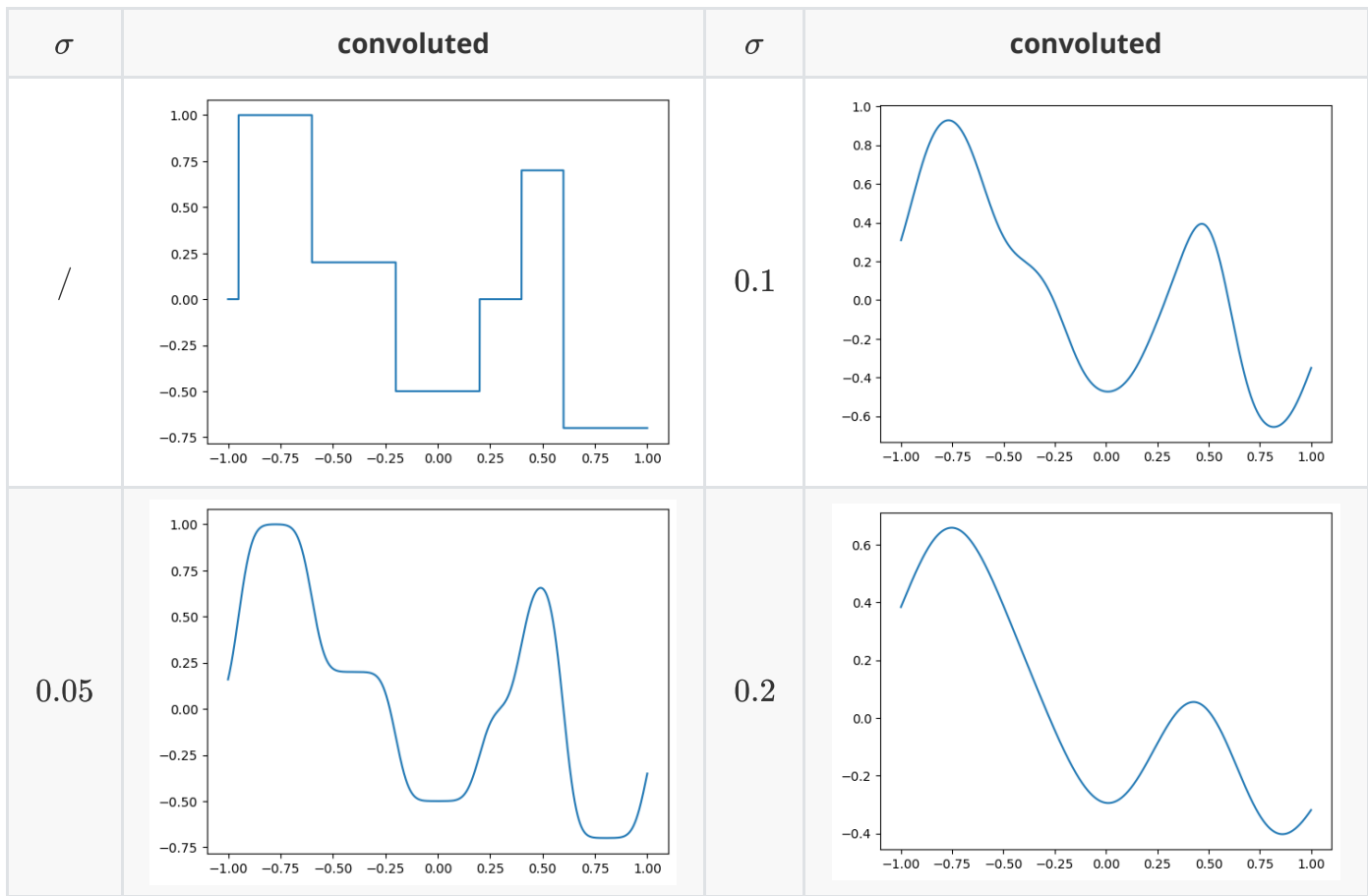
σ	Expanded Plots of Singular Values	Variance
0.05		0.400082
0.1		0.403275
0.2		0.414747

d.)

When calculating the convolution of f and A , we should make the height and width of A the same as the number of sampled points from f . The code example of $\sigma = 0.05$ is as below.

```
n = 10000
x = np.linspace(-1., 1., num=n)
fx = f(x)
A005 = convolution_matrix(n, 0, 0.05)
by = A005 @ fx
plt.plot(x, by)
plt.show()
```

The results are as below.



e.)

Actually, the matrix A is a Toeplitz matrix. The vector multiplication of the Toeplitz matrix can be accelerated by FFT. The Toeplitz matrix is characterised by the fact that all elements on the same diagonal are equal. Therefore, we can use the first row and the first column to represent the whole matrix, which can be written as a vector (assume it is z). Then we pad the data from f with 0 to align with the vector z to perform FFT convolution. The code example of $\sigma = 0.05$ is as below.

```
def conv1D(m, filter):
    m_feq = np.fft.fft(m)
    filter_feq = np.fft.fft(filter, m.shape[0])
    return np.fft.ifft(m_feq * filter_feq).real

n = 10000
x = np.linspace(-1., 1., num=n)
fx = f(x)
A005 = convolution_matrix(n, 0, 0.05)
z = np.hstack((A005[:, 0], np.zeros((1,)), A005[0, ::-1][:-1]))
by = conv1D(np.hstack((fx, np.zeros((n,)))), z)[:n]
plt.plot(x, by)
plt.show()
```

The results are the same as the normal matrix multiplication.

f.)

The matrix A , which is a Toeplitz matrix, is extremely regular and periodic, so we can use this property to accelerate the generation of the matrix A . The library function `scipy.linalg.toeplitz` can help us to do so. The code example of $\sigma = 0.05$ is as below.

```
def convolution_matrix_toeplitz(n, mu=0., sig=0.2):
    x = spatial_grid(n)
    xc = x - x[0]
    xr = -xc.copy()
    return toeplitz(gaussian_func(xc, n, mu, sig), gaussian_func(xr, n, mu, sig))
```

The effect is the same as the formal version, but it is faster. We can see the effect in the following case: using the new function to generate A makes it more than 8 times faster (4.2s to 0.5s).

$\sigma = 0.05$

```
n = 10000
x = np.linspace(-1., 1., num=n)
fx = f(x)
A005 = convolution_matrix(n, 0, 0.05)
z = np.hstack((A005[:, 0], np.zeros((1,)), A005[0, ::-1][:-1]))
by = conv1D(np.hstack((fx, np.zeros((n,)))), z)[:n]
plt.plot(x, by)
plt.show()
```

[186] ✓ 4.2s

Python

$\sigma = 0.05$

```
n = 10000
x = np.linspace(-1., 1., num=n)
fx = f(x)
A005 = convolution_matrix_toeplitz(n, 0, 0.05)
z = np.hstack((A005[:, 0], np.zeros((1,)), A005[0, ::-1][:-1]))
by = conv1D(np.hstack((fx, np.zeros((n,)))), z)[:n]
plt.plot(x, by)
plt.show()
```

[190]

✓ 0.5s

Python