

```
In [166]: AECheLon = forwardElimination(np.array([[1.0,2.0,3.0,-14.0],
[5.0,4.0,6.0,-37.0],[10.0,9.0,8.0,-41.0]]))
```

```
In [167]: inconsistentSystem(AECheLon)
```

```
Out[167]: False
```

```
In [168]: backsubstitution(AECheLon)
```

```
Out[168]:
```

```
array([[ 1.,  0.,  0., -3.],
       [ 0.,  1., -0.,  5.],
       [ 0.,  0.,  1., -7.]])
```

```
In [170]: A = np.loadtxt('h2m1.txt')
```

```
In [171]: AECheLonA = forwardElimination(A)
```

```
In [172]: inconsistentSystem(AECheLonA)
```

```
Out[172]: False
```

```
In [173]: backsubstitution(AECheLonA)
```

```
Out[173]:
```

```
array([[ 1.,  0.,  0.,  0.,  0.,  0.,  1.],
       [ 0.,  1., -0., -0., -0., -0.,  1.],
       [ 0.,  0.,  1.,  0.,  0.,  0.,  2.],
       [ 0.,  0.,  0.,  1.,  0.,  0.,  3.],
       [ 0.,  0.,  0.,  0.,  1., -0.,  5.],
       [ 0.,  0.,  0.,  0.,  0.,  1.,  8.]])
```

```
In [175]: B = np.loadtxt('h2m2.txt')
```

```
In [176]: AECheLonB = forwardElimination(B)
```

```
In [177]: inconsistentSystem(AECheLonB)
```

```
Out[177]: True
```

```
In [179]: C = np.loadtxt('h2m3.txt')
```

```
In [180]: AECheLonC = forwardElimination(C)
```

```
In [181]: inconsistentSystem(AECheLonC)
```

```
Out[181]: False
```

```
In [182]: backsubstitution(AECheLonC)
```

```
Out[182]:  
array([[ 1.00000000e+00,  0.00000000e+00,  0.00000000e+00,  
         0.00000000e+00,  0.00000000e+00,  0.00000000e+00],  
       [ 0.00000000e+00,  1.00000000e+00, -0.00000000e+00,  
       -0.00000000e+00, -0.00000000e+00, -0.00000000e+00],  
       [ 0.00000000e+00,  0.00000000e+00,  1.00000000e+00,  
         0.00000000e+00,  0.00000000e+00,  0.00000000e+00],  
       [ 0.00000000e+00,  0.00000000e+00,  0.00000000e+00,  
         1.00000000e+00, -0.00000000e+00, -0.00000000e+00],  
       [ 0.00000000e+00,  0.00000000e+00,  0.00000000e+00,  
         0.00000000e+00,  1.00000000e+00,  0.00000000e+00],  
       [ 0.00000000e+00,  0.00000000e+00,  0.00000000e+00,  
         0.00000000e+00,  0.00000000e+00,  3.55271368e-15]])
```

```
In [184]: D = np.loadtxt('h2m4.txt')
```

```
In [185]: AECheLonD = forwardElimination(D)
```

```
In [186]: inconsistentSystem(AECheLonD)
```

```
Out[186]: False
```

```
In [187]: backsubstitution(AECheLonD)
```

```
Out[187]:  
array([[ 1.,  0.,  0., -1.04878049, -2.41463415,  
         2.],  
       [ 0.,  1.,  0.,  2.07317073,  4.12195122,  
       -2.],  
       [ 0.,  0.,  1., -1.90243902, -4.17073171,  
         4.]])
```