

In [29]:

```
import networkx as nx
import numpy as np
from sklearn.cluster import KMeans, SpectralClustering
from sklearn.datasets import make_circles
import matplotlib.pyplot as plt
```

In [8]:

```
# SSBM Graph
n = 30
k = 3
A = 0.7
B = 0.1
W = [[A, B, B], [B, A, B], [B, B, A]]

G = nx.Graph()
labels = [0, 1, 2]

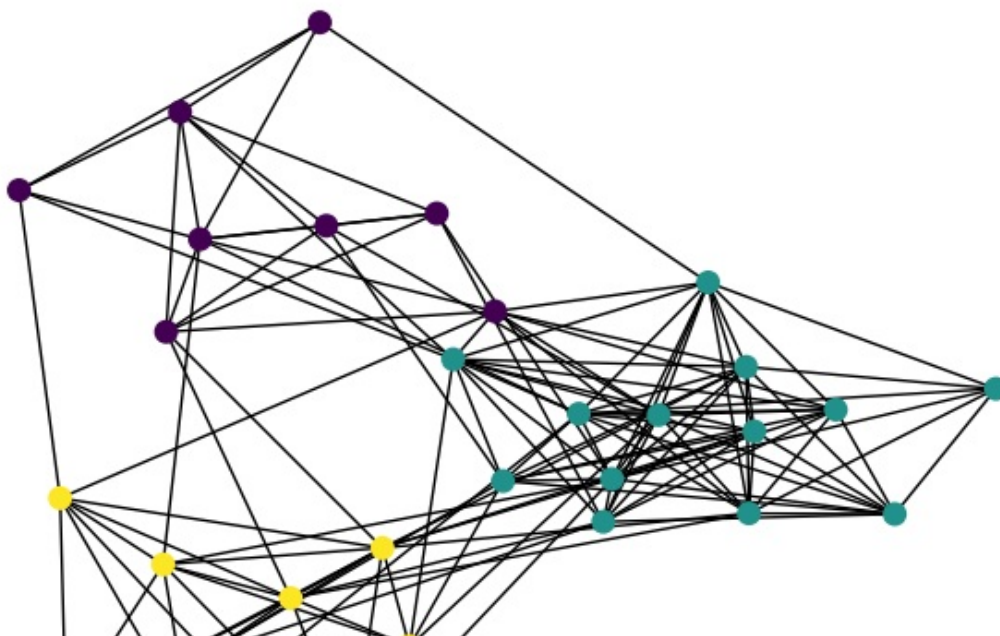
for i in range(n):
    label = np.random.choice(labels)
    G.add_node(str(i) + " " + str(label))

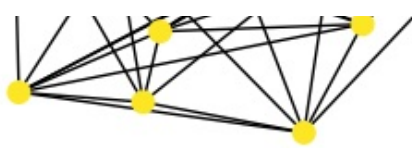
nodes = list(G.nodes)
for i in range(n):
    for j in range(i + 1, n, 1):
        node1 = nodes[i]
        node2 = nodes[j]
        label1 = node1.split(" ")[1]
        label2 = node2.split(" ")[1]
        p = W[int(label1)][int(label2)]
        if np.random.random() < p:
            G.add_edge(node1, node2)
```

In [15]:

```
spectral = SpectralClustering(k)
clusters = spectral.fit_predict(nx.to_numpy_array(G))
nx.draw(G, node_color=clusters, node_size=70)
```

```
/opt/homebrew/lib/python3.10/site-packages/sklearn/cluster/_spectral.py:717: UserWarning:
The spectral clustering API has changed. ``fit`` now constructs an affinity matrix from data. To use a custom affinity matrix, set ``affinity=precomputed``.
warnings.warn(
```

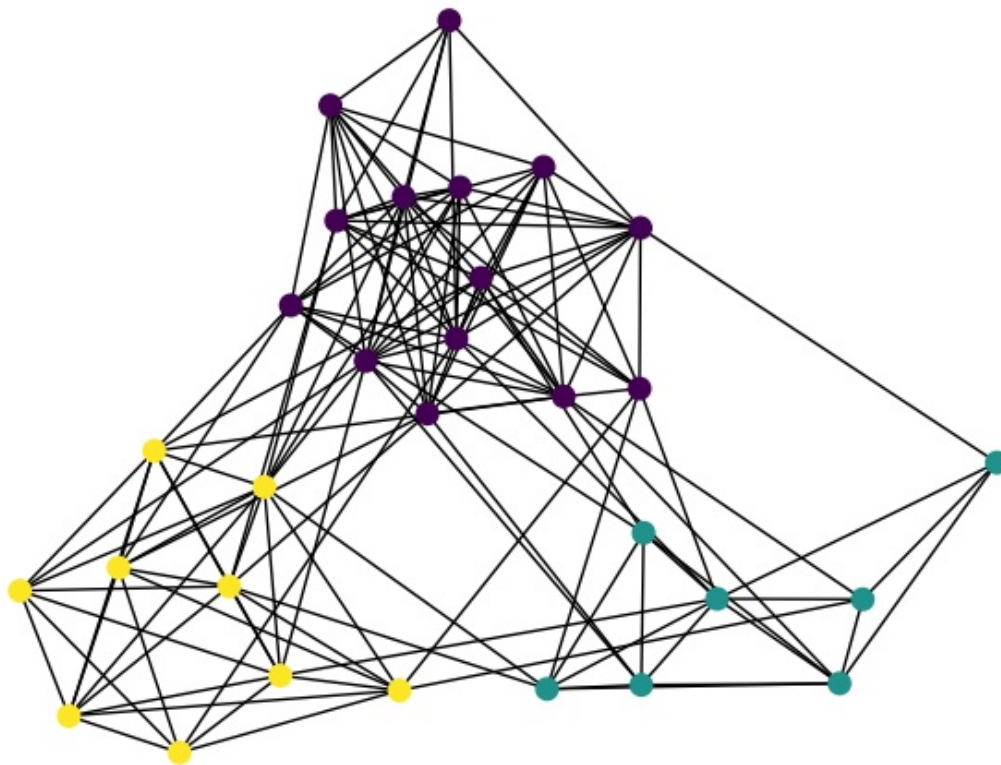




In [16]:

```
kmeans = KMeans(k)
clusters = kmeans.fit_predict(nx.to_numpy_array(G))
nx.draw(G, node_color=clusters, node_size=70)
```

/opt/homebrew/lib/python3.10/site-packages/sklearn/cluster/_kmeans.py:870: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning
warnings.warn(



There is a slight difference with 2 of the clusters containing differing numbers of nodes, but overall both algorithms capture very similar clusters. Both algorithms seem to capture the SSBM clusters pretty well as well.

In [19]:

```
circles = make_circles(500, noise=0.01)
```

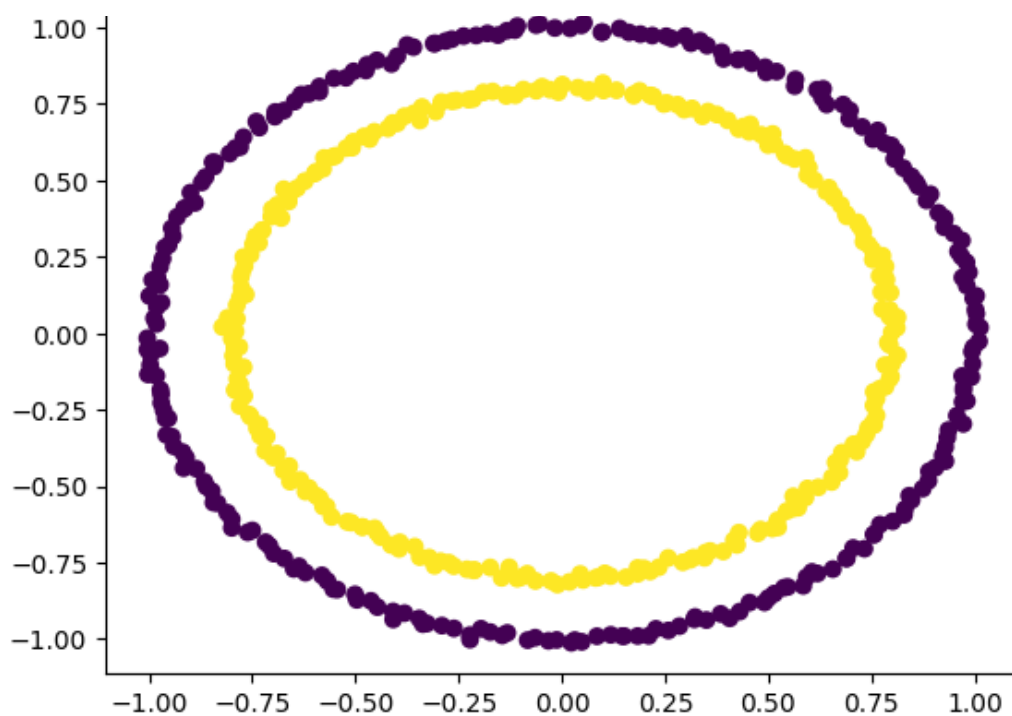
In [34]:

```
spectral = SpectralClustering(2, affinity="nearest_neighbors")
clusters = spectral.fit_predict(circles[0])
plt.scatter(circles[0][:, 0], circles[0][:, 1], c=clusters)
```

/opt/homebrew/lib/python3.10/site-packages/sklearn/manifold/_spectral_embedding.py:274: UserWarning: Graph is not fully connected, spectral embedding may not work as expected.
warnings.warn(

Out[34]:

<matplotlib.collections.PathCollection at 0x17fe87190>



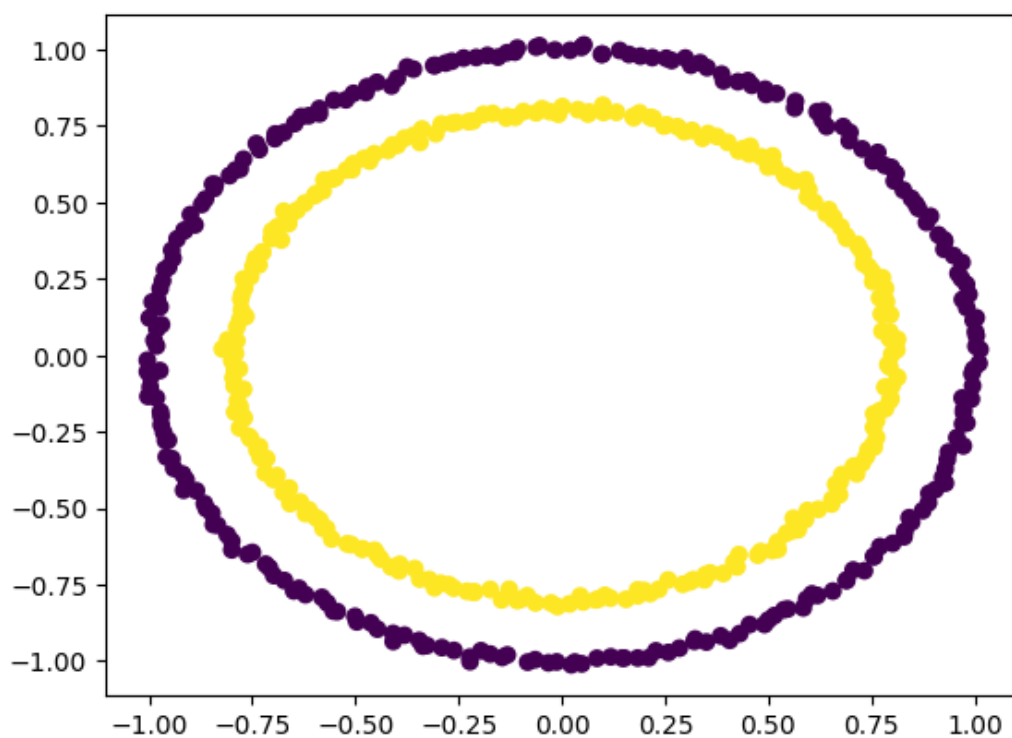
In [37]:

```
kmeans = KMeans(2)
clusters = spectral.fit_predict(circles[0])
plt.scatter(circles[0][:, 0], circles[0][:, 1], c=clusters)
```

/opt/homebrew/lib/python3.10/site-packages/sklearn/manifold/_spectral_embedding.py:274: UserWarning: Graph is not fully connected, spectral embedding may not work as expected.
warnings.warn(

Out[37]:

<matplotlib.collections.PathCollection at 0x2a794c280>



Both algorithms perform essentially the same, separating the inside vs the outside circles well.

3.

A.

		Player 2				
		a	b	c	d	e
Player 1	w	(3,1)	(1,1)	(6,1)	(5,2)	(2,2)
	x	(10,4)	(1,5)	(7,3)	(3,3)	(0,0)
	y	(0,2)	(2,5)	(2,4)	(3,3)	(1,0)
	z	(0,0)	(0,0)	(8,0)	(0,1)	(3,3)
	t	(4,3)	(1,2)	(1,4)	(4,6)	(1,2)

b d e
 w (1,1) (5,2) (2,2)
 y (2,5) (3,3) (1,0)
 z (0,0) (0,1) (3,3)

player 2 has no incentive to play A or C,

Thus player 1 then has no incentive to play X.

player 1 has no incentive to play t either.

Thus, bde survive for p2, w y z for p1.

B.

$$p(w) = p$$

$$p(b) = x$$

$$p(y) = q$$

$$p(d) = y$$

$$p(z) = 1 - p - q$$

$$p(e) = 1 - x - y$$

$$\text{indiff. w, y: } p + 5q + 2(1 - p - q) = 2p + 3q + 1 - p - q$$

$$\text{indiff. b, d: } x + 5y = 2x + 3y + 1 - x - y$$

$$d, e: 2x + 3y + 1 - x - y = 2x + 3(1 - x - y)$$

$$y, z: 2p + 3q + 1 - p - q = 3(1 - p - q)$$

$$x = \frac{1}{6} \quad y = \frac{1}{3}, \quad 1 - x - y = \frac{1}{2}$$

$$p = \frac{1}{2} \quad q = 0 \quad 1 - p - q = \frac{1}{2}$$

$$\text{So: } p1: \left(\frac{1}{2}, 0, 0, \frac{1}{2}, 0\right) \quad p2: \left(0, \frac{1}{6}, 0, \frac{1}{3}, \frac{1}{2}\right)$$

