1. Projections

a) 
$$\text{proj}_{\vec{a}}(\vec{b}) = \frac{\langle \vec{b}, \vec{a} \rangle}{\|\vec{a}\|^2} \vec{a} = \frac{3+0-1}{(1+0+1)^2} \vec{a} = \frac{2}{2} \vec{a} = \vec{a}$$

$$\operatorname{proj}_{\tilde{a}}(\tilde{b}) = 0$$

$$\|\vec{e}\|^2 = \|\text{proj}\vec{a}(\vec{b}) - \vec{b}\|^2 = \|-2\|^2 = \sqrt{4 + 4 + 4} = [12]$$

1

b) 
$$proj_{[3]}(\vec{b}) + proj_{[3]}(\vec{b}) = \frac{-4}{2} \begin{bmatrix} 1 \\ 0 \\ 1 \end{bmatrix} = \frac{4$$

$$\begin{vmatrix} -2 \\ = 4 \\ -2 \end{vmatrix}$$

$$\begin{vmatrix} 2 & -3 & 2 \\ \|\vec{e}\| = \| & 0 & \| = 18 \\ 3 & 3 & \end{vmatrix}$$

2. Least Squares ax = B 2 | = | 0.233  $\|\vec{e}\| = \|\vec{a} \times - \vec{b}\|^2 = \|2\|$ 1.117 -1 = 3.367 6 -1.533 6 0.3 8 0.933 ax1+x2 = 6 7 8 81 x = 120 0.05 -0.25 134 134 0.95 20 1.5 23 -0.25 20  $\vec{a}(0.95) + 1 = \vec{b}$ 0.9  $\|\vec{e}\|^2 = \|\vec{a}x_1 + \vec{x}_2 - \vec{b}\| = \|0.3\| =$ 2.7 0.6 Yes, it is better for the data. This is because error in part (b) is lower than error in post (a) which means model in part (b) fits better

3. Trilateration with Noise  $\|(x,y)-(x,y,y)\|=d_1$ a 1 6) let  $\vec{p} = (x,y)$ Pi=(x1,y1), P=(x2,y2), P==(x3,y3), Pi=(x4,y4) 1  $\|\vec{p}\| - 2 < \vec{p}, \vec{p}, > + \|\vec{p}_i\|^2 = d_i^2$  $= \frac{||\vec{p}||^2 - 2 < \vec{p}_1 \vec{p}_2) + ||\vec{p}_2||^2 = d^2}{||\vec{p}||^2 - 2 < \vec{p}_1 \vec{p}_2) + ||\vec{p}_3||^2 = d^2}$   $||\vec{p}||^2 - 2 < \vec{p}_1 \vec{p}_3) + ||\vec{p}_3||^2 = d^2$   $||\vec{p}||^2 - 2 < \vec{p}_1 \vec{p}_3) + ||\vec{p}_3||^2 = d^2$  $2 < \vec{p}, \vec{p}_1 > -2\vec{p}, \vec{p}_2 > + \|\vec{p}_2\|^2 - \|\vec{p}_1\|^2 = d_2^2 - d_1^2$   $2 < \vec{p}, \vec{p}_1 > -2 < \vec{p}, \vec{p}_3 > + \|\vec{p}_2\|^2 - \|\vec{p}_1\|^2 = d_3^2 - d_1^2$   $2 < \vec{p}, \vec{p}_1 > -2 < \vec{p}, \vec{p}_4 > + \|\vec{p}_2\|^2 - \|\vec{p}_1\|^2 = d_2^2 - d_1^2$  $\begin{aligned} 2\vec{p} & (\vec{p}_1 - \vec{p}_2) = d_2^2 - d_1^2 - ||\vec{p}_2||^2 + ||\vec{p}_1||^2 \\ 2\vec{p} & (\vec{p}_1 - \vec{p}_3)^T = d_3^2 - d_1^2 - ||\vec{p}_3||^2 + ||\vec{p}_1||^2 \\ 2\vec{p} & (\vec{p}_1 - \vec{p}_4)^T = d_4^2 - d_1^2 - ||\vec{p}_4||^2 + ||\vec{p}_1||^2 \end{aligned}$  $A \times = \vec{b}$ -(x,y,)-(x,y,)) X 2((x,y,)-(x,y,y)) X 2x,-0 dz - d, 2 - ||(x2, y2)|| + ||(x1, y1)||  $= |d_3^2 - d_1^2 - ||(x_3, y_3)||^2 + ||(x_1, y_1)||^2$   $|d_4^2 - d_1^2 - ||(x_4, y_4)||^2 + ||(x_1, y_1)||^2$ 

	4 Labelia Dataste visina
	4. Labeling Patients using Gene Expression Data
	Gene Expression Doctor
a)	Unknowns: X, , Xz, Xs, Xu, X5
	001,200,000
	5. Image Analysis
a)	$\alpha(x^2+y^2)+dx+ey=1$
	$A\vec{x} = \vec{b}$
	$\left(a\left(0.3^{2}+0.69^{2}\right)+0.3d-0.69e=1\right)$
	$a(0.5^2+0.87^2)+0.5d+0.87e=1$
	$a(0.9^2+0.86^2)+0.9d-0.86e=1$
	$\alpha(1^2 + 0.88^2) + d + 0.88e = 1$
	$a(1.2^{2}+0.82^{2})+1.24-0.82e=1$
	a(1.52+0.642)+1.5d+0.64e=1
	$(a(1.8^2) + 1.8d = 1$
	0.566 0.3 -0.69 0
	1.007 0.5 0.87 d = 1
	1.55 0.9 -0.86 e 1
	1.774 1 0.88
	2.112 1.2 -0.82
	2.66 1.5 0.64
	3.24 1.8 0
***	$\overrightarrow{X} = \overrightarrow{b}$
V	1 T 1 OT C
,	₹= (ATA)TAT 6

	Similar Ax=6	to p	part	(a), we	uil	find	Α	mat	ri×.	
	$0.3^{2}$	(0.3)(-0	2.69)	0.69		0.3	-0.65	Ma		1
	0.52	(0.5)(0.		0.87		0.5	0.87			١
	0.92	o-Xe.o)		0.862		0.9	-0.86	C	=	١
	1	0.88		0.582		1	0.88	9		1
	1.22	(1.2)(-0	0.82)	0.82		1.2	-0.82	le		1
	11.5	(1.5)(0.	64)	0.64	-	1.5	0.64	_		1
	1.82	0		0		1-8	0			
	₹ = (A	TATIATE	»		•					
10										
			\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	~						
(d)	Error	in po	urt (	d) (s	lowe	r th	an er	ror i	^ F	oar
0)	Becau	use of	2 tu	at, t	he a	ellipse	tecl	ror i	ne	ar
d)	Becau	use of	2 tu	d) is	he a	ellipse	tecl	ror i	n f	oar
d)	Becau	use of	2 tu	at, t	he a	ellipse	tecl	ror i	n F	ar
d)	Becau	the of	2 + tv	at, the ci	rcle	ellipse meth	tecl	nnic	n f	Jar
d)	Becau	the of	2 + tv	at, t	rcle	ellipse meth	tecl	nnic	jue jue	Jar
d)	Becaubetter	the of	>. Hc	the ci	Proce	ellipse meth	tech	nnic	juè	
	Becau better I wo	the of	on.	the cir	Proce	ellipse meth ess	tech sol.	nnic	jue - r	rea
	Because Better	the of the	on res	the cir	Proce	ellipse meth ess ork ed t	tech sol.	nnic	jue - r	rea
	Because Better	the of	on res	the cir	Proce	ellipse meth ess ork ed t	tech sol.	nnic	jue - r	rea
	Because Better	the of the	on res	the cir	Proce	ellipse meth ess ork ed t	tech sol.	nnic	jue - r	rea
	Because Better	the of the	on res	the cir	Proce	ellipse meth ess ork ed t	tech sol.	nnic	jue - r	rea
	Because Better	the of the	on res	the cir	Proce	ellipse meth ess ork ed t	tech sol.	nnic	jue - r	rea

### **EECS16A: Homework 13**

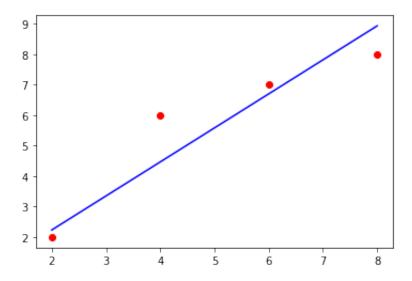
```
In [1]: from __future__ import division
%pylab inline
import numpy as np
import matplotlib.pyplot as plt
import scipy.io
import sys
```

Populating the interactive namespace from numpy and matplotlib

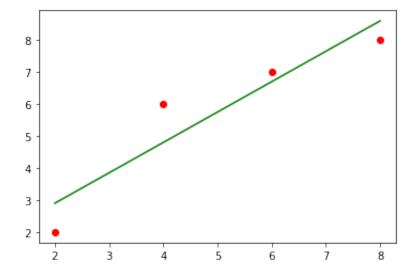
## Mechanical Linear Least Squares (optional)

## In [15]: # use for plotting print("part A") a = np.array([2,4,6,8]) b = np.array([2,6,7,8]) for i in range(len(a)): plt.plot(a[i], b[i], color="red", marker = 'o') plot(a, a\*1.1166666667, 'r', color="blue") plt.show() print("part B") for i in range(len(a)): plt.plot(a[i], b[i], color="red", marker = 'o') plot(a, a\*0.95+1, 'r', color="green") plt.show()

### part A



part B



### **Demonstration: Trilateration With Noise!**

In lecture, we learned how the GPS receiver determines its location once it knows the distance of the various signaling beacons from itself. This method is called *trilateration*.

In this demonstration, we're going to further explore the connection between trilateration and least squares through a toy problem with four beacons and one GPS receiver.

We are given *three* possible sets of measurements for the distances of each of the beacons from the receiver:

- 1. First, the ideal set of measurements.  $d_1 = d_2 = d_3 = d_4 = 5$ .
- 2. Next, a set of imperfect measurements.  $d_1 = 5$ ,  $d_2 = 4.5$ ,  $d_3 = 5$ ,  $d_4 = 5.5$ .
- 3. Finally, a set of mostly perfect measurements, but  $d_1$  is a very bad measurement. We have  $d_1=6.5$  and  $d_2=d_3=d_4=5$ .

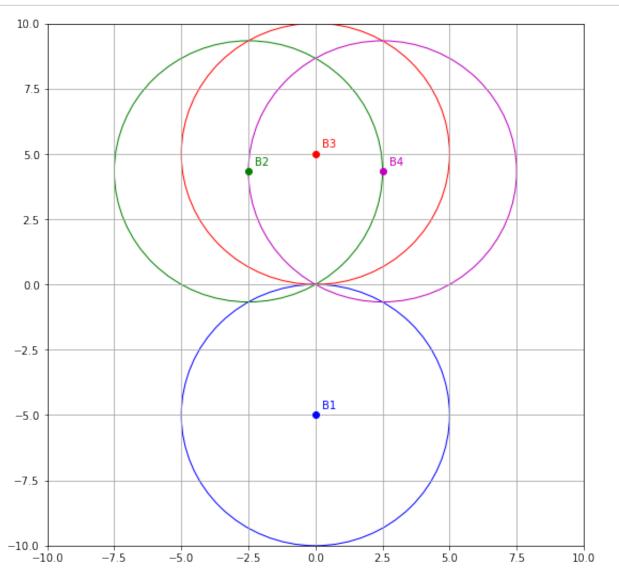
First, we set up some notation for the positions of the beacons, as well as their respective distances from the receivers.

```
In [16]: from utils import *
         ideal\_distances = [5, 5, 5, 5]
         imperfect_distances = [5.5, 4.5, 5, 5]
         one_bad_distances = [6.5, 5, 5, 5]
         #these are the coordinates of the beacons
         positions = np.array([
              [0, -5],
              [-5 / 2, 5 * np.sqrt(3) / 2],
              [5 / 2, 5 * np.sqrt(3) / 2],
         ])
         xpositions = positions[:, 0]
         ypositions = positions[:, 1]
         # setup to make the helper functions work
         register(positions=positions)
         register(xpositions=xpositions)
         register(ypositions=ypositions)
```

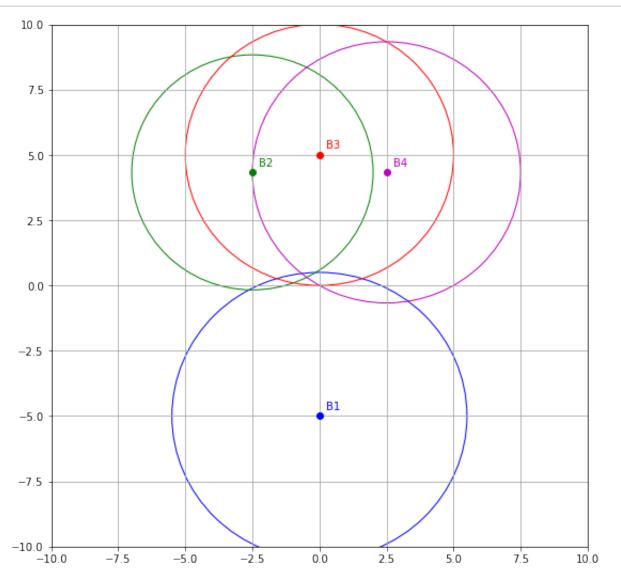
### Part (d)

Now, for each of the three above cases, let's plot a circle for each beacon whose radius corresponds to the reported distance of the receiver. The intersection of the circles will tell us, intuitively, where the receiver is located! (Note that the circles do not necessarily intersect at one point for all three of the cases. Think about what this means).

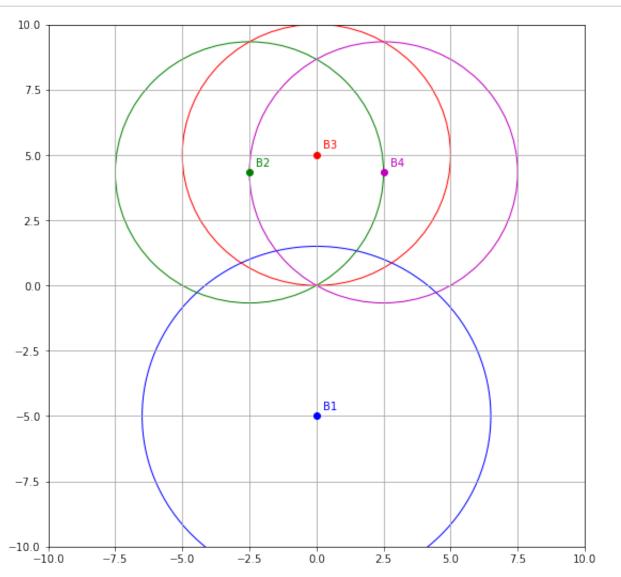
In [17]: plot(ideal\_distances)



In [18]: plot(imperfect\_distances)



In [19]: plot(one\_bad\_distances)



### Part (e)

Now, let's solve for the location of the receiver,  $\begin{bmatrix} x \\ y \end{bmatrix}$ , using least squares.

Recall that we made the system of equations linear by subtracting the equation for the first beacon from each of the equations for the other beacons.

This will result in the system of equations:

$$A\begin{bmatrix} x \\ y \end{bmatrix} = \vec{b}$$

You will define A and  $\vec{b}$  in the code blocks below.

### Part (f)

Fill in the entries of b in the below function to correspond to the entries of  $\hat{b}$  from the problem.

```
In [61]: def make_b(distances):
    """
    Since `b` depends on `distances`, we implement it using a function different `b` vectors depending of which set of distances we are : (i.e. ideal_distances, imperfect_distances, or one_bad_distances)

Examples of how to call the function:
    make_values(ideal_distances) OR
    make_values(imperfect_distances) OR
    make_values(one_bad_distances)

b = np.zeros(3)

b[0] = -(xpositions[1]**2 + ypositions[1]**2) + (xpositions[0]**2
    b[1] = -(xpositions[2]**2 + ypositions[2]**2) + (xpositions[0]**2
    b[2] = -(xpositions[3]**2 + ypositions[3]**2) + (xpositions[0]**2
    return b

make_b(ideal_distances)
```

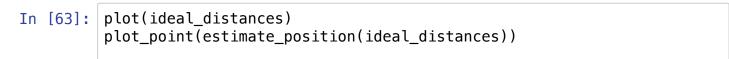
Out[61]: array([3.55271368e-15, 0.00000000e+00, 3.55271368e-15])

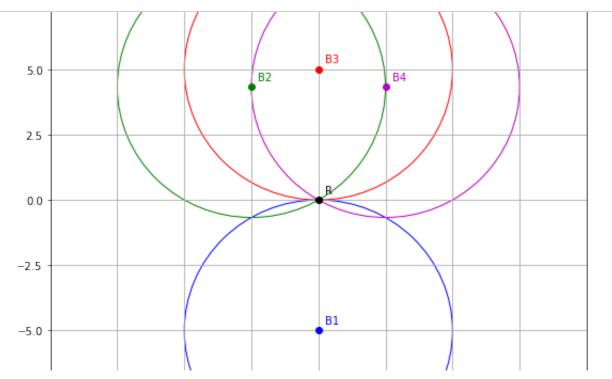
### Part (g)

Now, calculate the linear least squares estimate for the ideal\_distances data and plot the results. We have given you code for the implementation for this part.

```
In [62]: def estimate_position(distances):
    U1 = np.dot(A.T, A)
    U2 = np.dot(A.T, make_b(distances))
    least_squares_sol = np.dot(np.linalg.inv(U1),U2)
    return least_squares_sol
    estimate_position(ideal_distances)
```

Out[62]: array([ 8.59260618e-33, -1.20930181e-16])

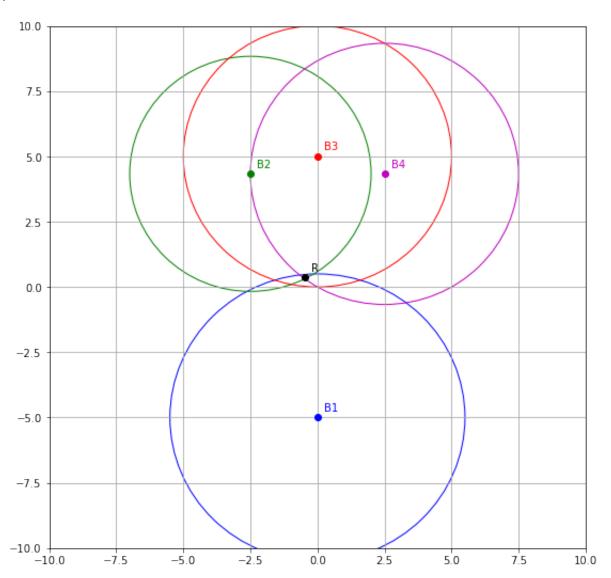




What about for the imperfect\_distances and one\_bad\_distances? Copy and modify the above code to compute and plot the least-squares trilateration solutions for those two cases, and comment on the quality of the solution in each case.

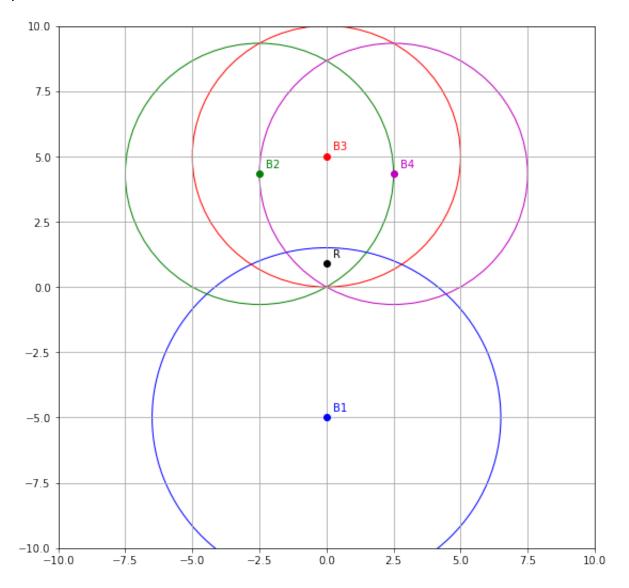
In particular, for one\_bad\_distances, is the solution the best possible? In other words, if you were trying to identify your position from the graph by hand, would you have chosen the same point that our trilateration solution did?

### point: [-0.475 0.35531308]



```
In [65]: ### one_bad_distances case
           plot(one_bad_distances)
           p = estimate_position(one_bad_distances)
print("point:", p)
           plot_point(p)
```

point: [-6.40790822e-17 9.01832908e-01]



### Part (h)

You should see that linearizing and solving least squares did not always do well in the above cases. Now, let's try something else. For any candidate location of the receiver, we can define the distance from  $i_{th}$  beacon as  $\hat{d}_i$  and define  $\hat{d}$  as:

$$\vec{\hat{d}} = \begin{bmatrix} \hat{d}_1 \\ \hat{d}_2 \\ \hat{d}_3 \\ \hat{d}_4 \end{bmatrix}$$

So we might want to minimize the following:

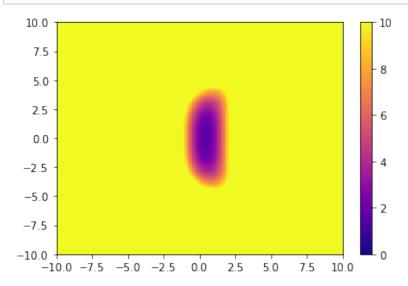
$$\sum_{i=1}^{4} (d_i - \hat{d}_i)^2$$

Let's make a plot that will help us visualize this. (x, y) in the plot correspond to candidate positions of the receivers and the color corresponds to the cost.

Let's first plot the costs when the one\_bad\_distances data is used. You should see that this cost is minimized at (0,0).

In [66]: def cost\_function(point, actual\_distances):
 cost = 0
 for pos, actual\_distance in zip(positions, actual\_distances):
 cost += (distance\_between\_points(point, pos) - actual\_distance
 return cost

plot\_cost(one\_bad\_distances, cost\_function)



Let's verify numerically that the above approach does indeed do better than the least-squares trilateration approach in the one\_bad\_distances case. Use cost\_function to compare the cost of (0, 0) with the cost of your estimated position obtained from the least-squares solution in all three cases. When does least squares do worst, compared to the new approach?

In [70]: with ideal distances:", cost\_function([0,0], ideal\_distances))
 ted position with ideal distances:", cost\_function(estimate\_position(i
 with imperfect distances:", cost\_function([0,0], imperfect\_distances))
 ted position with imperfect distances:", cost\_function(estimate\_positi
 with one bad distance:", cost\_function([0,0], one\_bad\_distances))
 ted position with one bad distance:", cost\_function(estimate\_position()

Cost of (0,0) with ideal distances: 0.0
Cost of estimated position with ideal distances: 0.0
Cost of (0,0) with imperfect distances: 0.5
Cost of estimated position with imperfect distances: 0.12767439181123
577
Cost of (0,0) with one bad distance: 2.25
Cost of estimated position with one bad distance: 2.31715040201415

## **Labeling Patients**

```
In [37]: import numpy as np
In [47]: # Part B
         A = np.load('gene_data_train.npy')
         b = np.load('diabetes_train.npy')
         model = list(np.linalg.lstsq(A, [i[0] for i in b], rcond=None)[0])
         print(model)
         [-0.15646168983619893, 0.09239417683325737, 0.4805397398225526, -0.58]
         47017953319975, -0.3535073412970997]
In [51]: # Part C
         # You may find np.sign useful to generate your +/- 1 prediction vector
         A_test = np.load('gene_data_test.npy')
         b_test = np.load('diabetes_test.npy')
         correct = 0
         for i in range(len(A_test)):
             data = A_test[i]
             pred = data[0] * model[0] + data[1] * model[1] + data[2] * model[2]
             print("Prediction:", int(np.sign(pred)), "Actual:", b_test[i][0])
             if int(np.sign(pred)) == b_test[i][0]:
                 correct += 1
         print("Accuracy:", correct / len(b_test) * 100, "%")
         Prediction: 1 Actual: 1
         Prediction: -1 Actual: -1
         Prediction: -1 Actual: -1
         Prediction: 1 Actual: 1
         Accuracy: 100.0 %
```

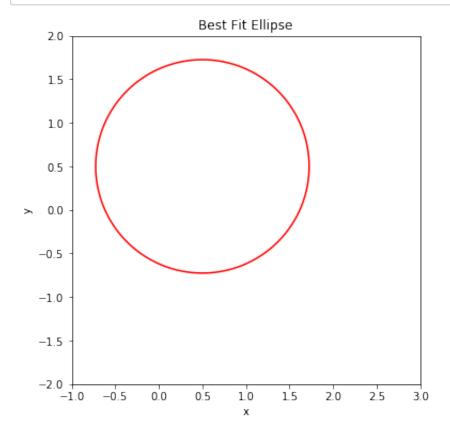
## **Image Analysis**

```
In [71]:
         def plot_circle(a, d, e):
             You can use this function to plot circles with parameters a,d,e.
             The parameters are described in the homework pdf.
             is\_circle = d**2 + e**2 - 4*a > 0
             assert is_circle, "Not a circle"
             XLIM_L0 = -1
             XLIM\ HI = 3
             YLIM L0 = -2
             YLIM HI = 2
             X_COUNT = 400
             Y_COUNT = 400
             x = np.linspace(XLIM_LO, XLIM_HI, X_COUNT)
             y = np.linspace(YLIM_LO, YLIM_HI, Y_COUNT)
             x, y = np.meshgrid(x, y)
             f = lambda x, y: a*(x**2 + y**2) + d*x + e*y
             c1 = plt.contour(x, y, f(x,y), [1], colors='r')
             plt.axis('scaled')
             plt.xlabel('x')
             plt.ylabel('y')
             plt.title("Best Fit Circle")
```

```
In [72]:
         def plot_ellipse(a, b, c, d, e):
             You can use this function to plot ellipses with parameters a-e.
             The parameters are described in the homework pdf.
             is_ellipse = b**2 - 4*a*c < 0
             assert is_ellipse, "Not an ellipse"
             XLIM_L0 = -1
             XLIM_HI = 3
             YLIM LO = -2
             YLIM_HI = 2
             X_COUNT = 400
             Y_COUNT = 400
             x = np.linspace(XLIM_LO, XLIM_HI, X_COUNT)
             y = np.linspace(YLIM_LO, YLIM_HI, Y_COUNT)
             x, y = np.meshgrid(x, y)
             f = lambda x,y: a*x**2 + b*x*y + c*y**2 + d*x + e*y
             c1 = plt.contour(x, y, f(x,y), [1], colors='r')
             plt.axis('scaled')
             plt.xlabel('x')
             plt.ylabel('y')
             plt.title("Best Fit Ellipse")
```

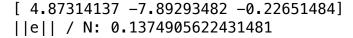
```
In [73]: # Here is an example of plot_ellipse.
# This plots (x-1)**2 + (y-1)**2 = 1,
# which is a circle centered at (1,1).

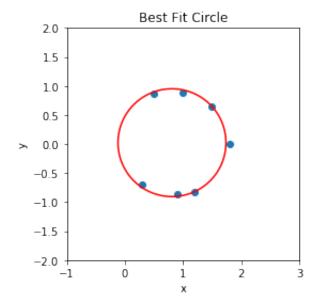
plt.figure(figsize=(6,6))
plot_ellipse(1, 0, 1, -1, -1)
```



You may find <u>plt.scatter (http://matplotlib.org/api/pyplot\_api.html)</u> useful for plotting the points.

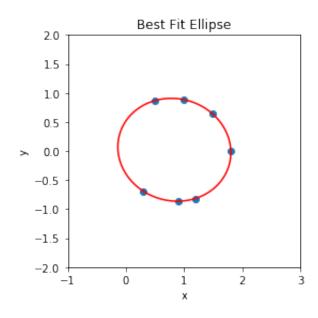
```
In [85]:
         # PART C
         xy = np.array([[0.3, -0.69],
                         [0.5, 0.87],
                         [0.9, -0.86],
                         [1, 0.88],
                         [1.2, -0.82],
                         [1.5, .64],
                         [1.8, 0]])
         x = xy[:,0]
         y = xy[:,1]
         # plot the data points
         plt.scatter(x,y)
         A = np.array([x**2+y**2,x,y]).T # Hint: this generates the A matrix
         b = [1] * 7
         circle_params = np.linalg.lstsq(A, b, rcond=None)[0]
         print(circle_params)
         plot_circle(circle_params[0],circle_params[1],circle_params[2])
         \# ||e|| / N
         print("||e|| / N:", np.linalg.norm(np.dot(A, circle_params) - b) / 7)
```





# In [86]: # PART D # plot the data points plt.scatter(x,y) A = np.array([x\*\*2, x\*y, y\*\*2, x, y]).T b = [1] \* 7 ellipse\_params = np.linalg.lstsq(A, b, rcond=None)[0] print(ellipse\_params) plot\_ellipse(ellipse\_params[0],ellipse\_params[1],ellipse\_params[2],ell # ||e|| / N print("||e|| / N:",np.linalg.norm(np.dot(A, ellipse\_params) - b) / 7)

[ 4.10382951 0.48711384 4.93938449 -6.85032284 -0.62259259] ||e|| / N: 0.012853829087236146



In [ ]: