Question 1:

```
Z:\C$432\A10>python kmeasure.py
44.0
44.5
38.0
29.7
28.05
50.0
41.0
29.0
33.8
30.1
Z:\C$432\A10>
```

K-Estimate Computations:

44.0

44.5

38.0

29.7

28.05

50.0

41.0

29.0

33.8

30.1

Name	K = 1	K = 2	K = 5	K = 10	K = 20
	44	44.5	38.0	29.7	28.05
http://f-measure.blogspot.com/					
http://ws- dl.blogspot.com/	50.0	41.0	29.0	33.8	30.1

After obtaining the necessary functions from the Collective Intelligence Programming textbook, the computations were able to be done correctly.

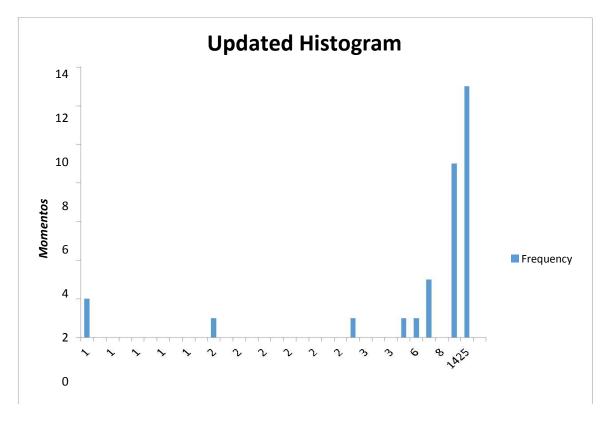
Question 2:

```
atria.cs.odu.edu - PuTTY
atria:~/C5432/A10> python libvsmCizer.py
       not-Zika
       not-Research
       not-Ebola
       not-HIV
       not-Outbreak
HIV:
        not-Research
        not-Cancer
       not-Ebola
       not-Outbreak
Research:
       not-Zika
       not-Cancer
       not-Ebola
       not-Outbreak
Ebola:
       not-Zika
       not-Research
       not-Cancer
       not-HIV
       not-Outbreak
Outbreak:
       not-Research
       not-Cancer
       not-Ebola
       not-HIV
Zika:
        not-Research
        not-Cancer
        not-Ebola
        not-HIV
        not-Outbreak
atria:~/CS432/A10>
```

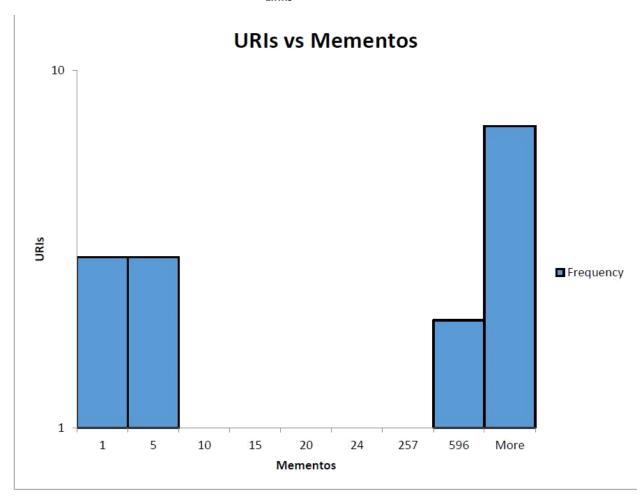
Category	Cross-Validation%	Correction%
Cancer	85.00%	66.00%
HIV	35.00%	42.00%
Research	68.00%	76.00%
Ebola	88.00%	66.00%
Outbreak	72.00%	46.00%
Zika	91.00%	95.00%

For question 2, I was able to figure out the LIBVSM scheme, however the 500 term vectors and 10-fold cross-validations was rather confusing. I researched on it for a while trying to figure out on how to implement it but no such luck up until Saturday when I decided to give it one last look.

Question 3 (Extra Credit):







For the Extra Credit question 3, the first graph is an updated histogram compared to the old one. As it can be seen the graphs differ when it was produced in Assignment 2.

1	2
1	0
1	0
1	0
1	0
1	0
1	0
1	0
1	0
1	0
2	1
2	0
2	0
2	0
2	0
2	0

2	0
2	0
2	0
2	0
2	0
3	1
3	0
3	0
3	0
5	1
6	1
8	3
8	0
43	9
1425	13