Applied Property Testing For Drug Repurposing

Name: Omer Mualem Advisor: Dr. Sarel Cohen

Workshop Track: Research Project Number: 15006801

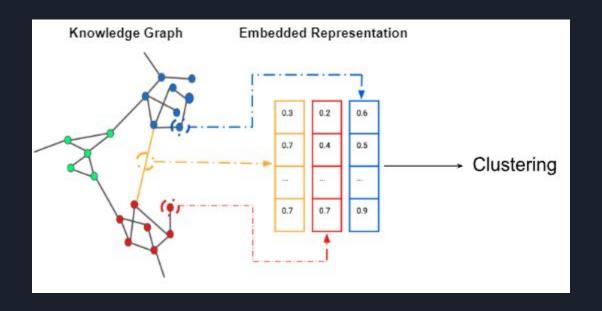
BackGround

- In the face of a global viral outbreak, the immediate need for effective treatments becomes paramount
- My project aims to circumvent these challenges by leveraging existing
 FDA-approved drugs through a process known as drug repurposing
- Test a new clustering algorithm

Solution

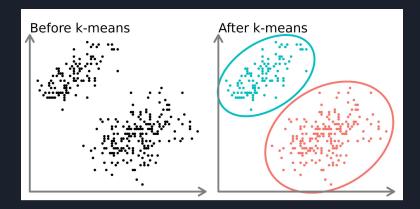
- Relying on recent work on drug repurposing by Cohen et al[PLOS ONE 2023]
- Change clustering algorithm used
- Test for improved result

Drug Repurposing Model Architecture



Current Clustering Algorithm

- K-means
- Minimize the variance within each cluster and maximize the variance between clusters



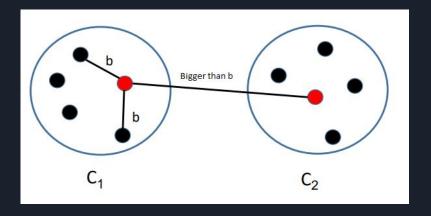
New Clustering Algorithm

- first suggested in a paper by Michal Pernas, Dar, Ron and Alon in 2003
- Property testing
- Utilize the power of randomness

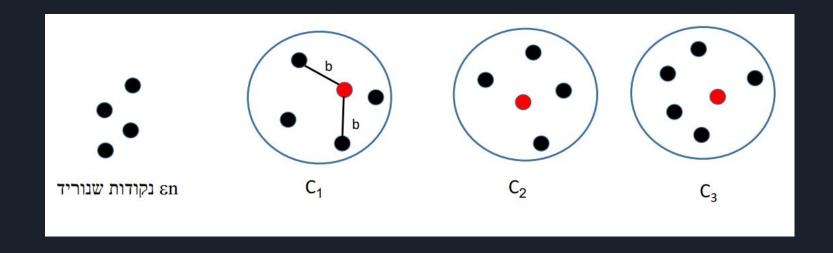
Parameters

- In the new algorithm we introduce new parameters
 - epsilon outlier percentage parameter
 - b cluster max radius
 - k max number of clusters

Parameters Meaning



Parameters Meaning



New Clustering Algorithm

Input: Set of points X, parameters k, b and a outlier parmeter epsilon

Output: Is X (k,b)-clusterable or epsilon-far from being (k,2b)-clusterable

Steps:

- 1. Let Rep_1 be a random points from X
- 2. i = 1, Find_New_Rep = True
- 3. For i < k + 1 and Find_New_Rep = True:
 - 3.1. Choose In(3k) / epsilon points from a uniform distribution on X
 - 3.2. If a point x in the sample has > b distance from all current representors

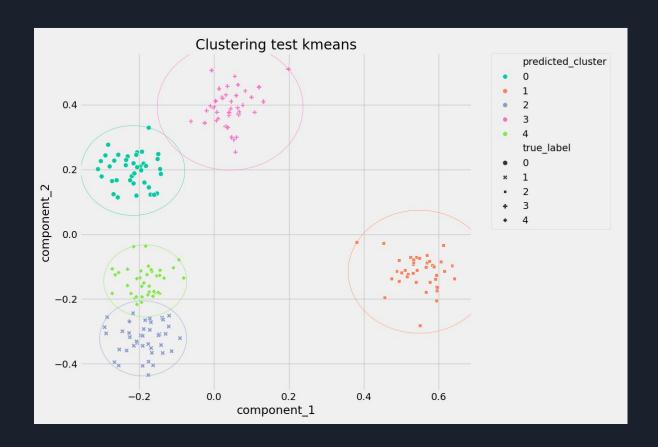
then
$$Rep_i+1 = x$$
 and $i = i + 1$

- 3.3. Else Find_New_Rep = False
- 4. If i <= k we accept the representors and parameters, else we decline

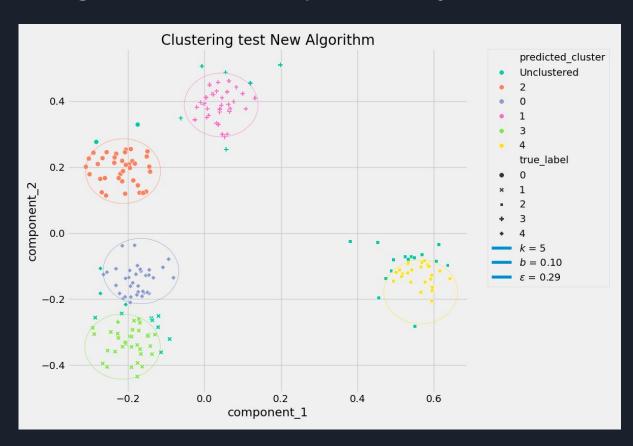
Current Progress

- Created a general pipeline for clustering algorithms using tensorboard to save outputs
- Tested the pipeline with the k-means algorithm on synthetic data
- Implemented the new clustering algorithm
- Tested the new clustering algorithm on synthetic data
- Tested the new clustering algorithm on SIFT 1M dataset training set including 100,000 samples with dimension 128

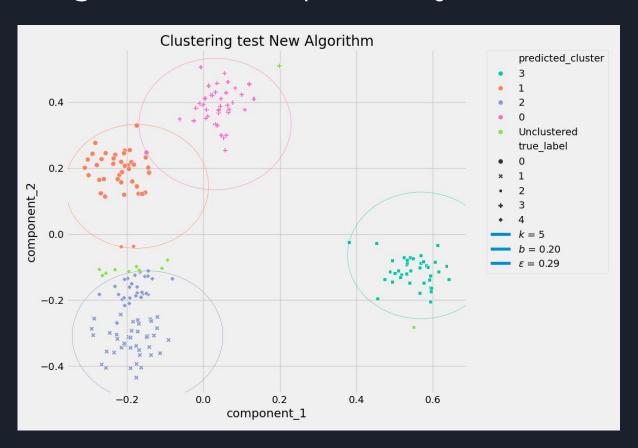
K-means Example On Synthetic Data



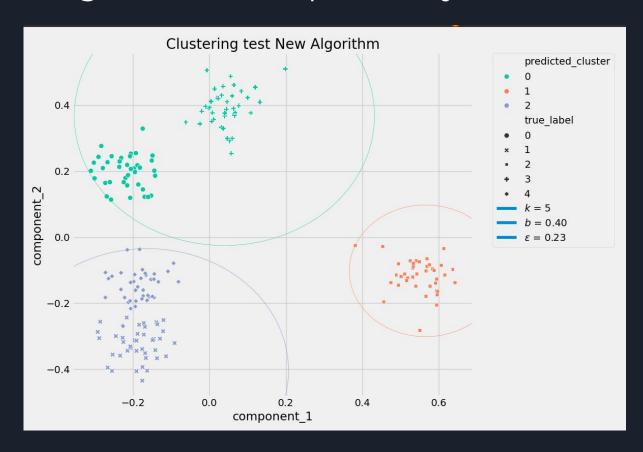
New Algorithm Example On Synthetic Data

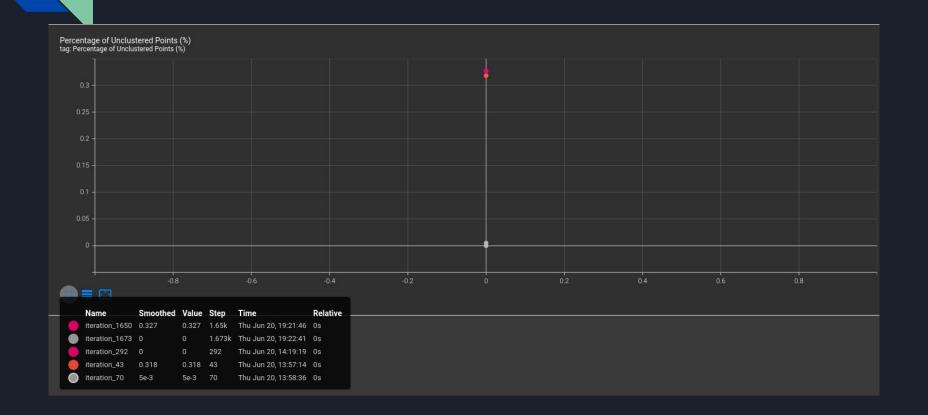


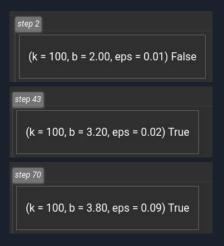
New Algorithm Example On Synthetic Data

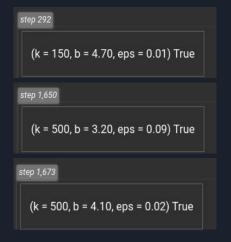


New Algorithm Example On Synthetic Data









```
        Name
        Smoothed
        Value
        Step
        Time
        Relative

        iteration_1650
        0.327
        0.327
        1.65k
        Thu Jun 20,19:21:46
        0s

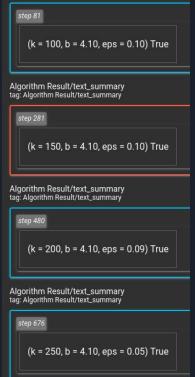
        iteration_1673
        0
        0
        1.673k
        Thu Jun 20,19:22:41
        0s

        iteration_292
        0
        0
        292
        Thu Jun 20,14:19:19
        0s

        iteration_43
        0.318
        0.318
        43
        Thu Jun 20,13:57:14
        0s

        iteration_70
        5e-3
        5e-3
        70
        Thu Jun 20,13:58:36
        0s
```





```
(k = 300, b = 3.80, eps = 0.09) True
Algorithm Result/text summary
tag: Algorithm Result/text_summary
     (k = 350, b = 4.40, eps = 0.01) True
Algorithm Result/text_summary
tag: Algorithm Result/text_summary
     (k = 400, b = 4.40, eps = 0.01) True
Algorithm Result/text_summary tag: Algorithm Result/text_summary
     (k = 450, b = 4.40, eps = 0.01) True
Algorithm Result/text_summary
tag: Algorithm Result/text. summary
     (k = 500, b = 4.10, eps = 0.09) True
```



Conclusions And Moving Forward

- Testing the new algorithm provides us with deeper insights into the data structure
- If the drug repurposing dataset is structured as clusters of circles we can possibly improve current clustering by using a different technique
- Next we need to test our clustering on SIFT 1M test dataset
- Finally use the new algorithm on our drug repurposing problem