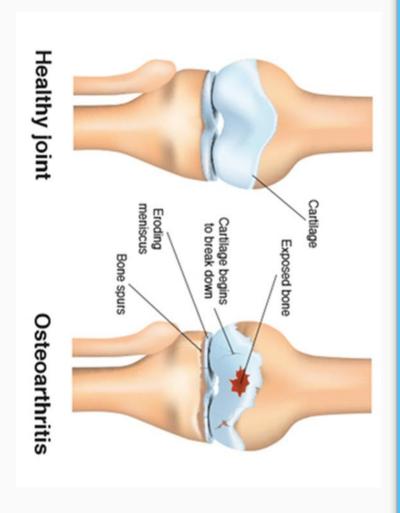
BIOINFORMATICS APPLICATIONS IN OSTEOARTHRITIS

Mackenzie Fleischer

Introduction

Osteoarthritis:

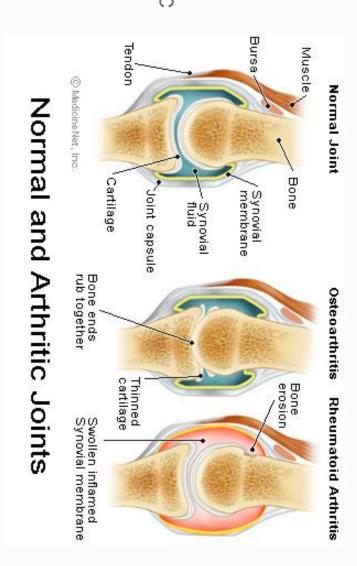
- Cartilage between joints breaks down leading to pain, stiffness, and swelling
- Most common chronic condition of the joints
- Affects 27 million Americans



Introduction

Osteoarthritis:

- Important to distinguish from other types of arthritis
- Typically a result of a traumatic joint injury
- Progression may lead to joint replacement surgery



Data Description



- 16 patients: 10 RA, 6 OA
- Synovial membrane samples
- RNA microarray analysis

Analysis Methods

Project:

- Log2 transformation
- eBayes function to identify differentially expressed genes
- Clustering (kmeans, cmeans)
- Heatmap clustering
- Gene Interaction

Paper:

- Classifier training & testing- rules assign a to rank a gene
- Rules applied to samples prior to clustering

```
203915_at
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217986_s_at 0.00176448 BAZ1A
                                   210031_at
                                                                                                                                                                              205159_at
                                                                                                                                                                                                              205890_s_at 0.00092858 UBD///GABB ubiquitin D///gamma-aminobutyric acid type B receptor subunit 1
                                                                                                                                                                                                                                                 209604_s_at 0.00020665 GATA3
                                                                                                                                                                                                                                                                                      221003_s_at 0.00020571 CAB39L
                                                                                                                                                                                                                                                                                                                        0.00020571 ADAMDEC1 ADAM like decysin 1
                                       0.00176448 CD247
                                                                                                                                                                                                                                                                                                                                                             adj.P.Val
                                                                                                                                                                               0.00103936 CSF2RB
                                                                       0.00176448 CXCL9
                                                                                                        0.00176448 PRELP
                                                                                                                                            0.00119727 PSMB9
                                                                                                                                                                                                                                                                                                                                                             Gene Symbol Gene.title
                                                                                                                                                                              colony stimulating factor 2 receptor beta common subunit
  bromodomain adjacent to zinc finger domain 1A
                                                                                                                                           proteasome subunit beta 9
                                                                                                                                                                                                                                                                                    calcium binding protein 39 like
                                      CD247 molecule
                                                                   C-X-C motif chemokine ligand 9
                                                                                                          proline and arginine rich end leucine rich repeat protein
                                                                                                                                                                                                                                                   GATA binding protein 3
```

K-means clustering:

```
2 1 1 2 2 2 1 2 6

GSM1339626 GSM1339627 GSM1339628 GSM1339629 GSM1339630 GSM1339631 GSM1339632 GSM1339633
2 1 2 1 1 1 1 1 1
                                                                                                                               GSM1339618 GSM1339619 GSM1339620 GSM1339621 GSM1339622 GSM1339623 GSM1339624 GSM1339625
                                                                                                                                                                                      Clustering vector:
```

C-means fuzzy clustering: data point can belong to one or more cluster

```
Closest hard clustering:
GSM1339618 GSM1339619 GSM1339620 GSM1339621 GSM1339622 GSM1339623 GSM1339624 GSM1339625
1 1 2 1
```

> table(cluster\$clust)

70% ACCURATE

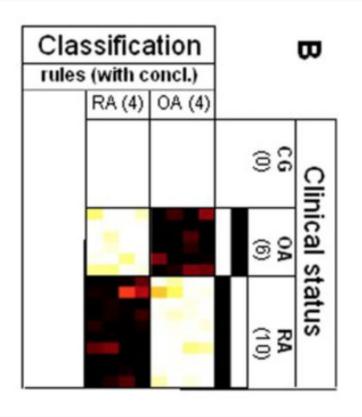
9 7

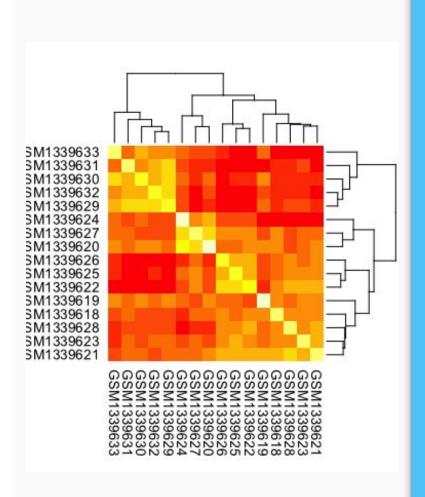
> table(ccluster\$clust)

2 68% ACCURATE

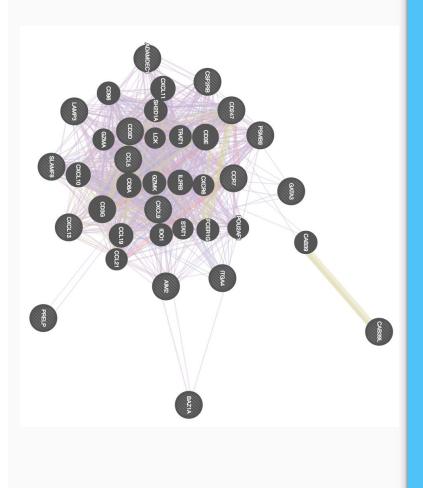
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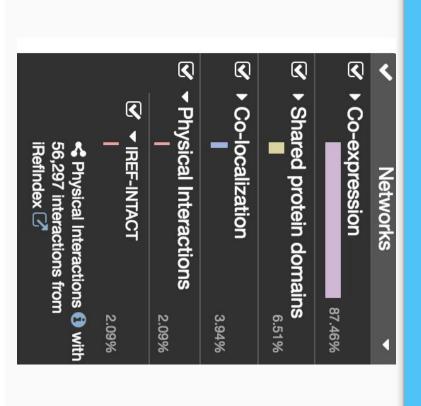
0	0	0	0	0	0	70	Z	70	70	70	70	70	70	RA	70	G	
OA	OA	OA	OA	OA	OA	RA	۶	RA	Group								
GSM1339633	GSM1339632	GSM1339631	GSM1339630	GSM1339629	GSM1339628	GSM1339627	GSM1339626	GSM1339625	GSM1339624	GSM1339623	GSM1339622	GSM1339621	GSM1339620	GSM1339619	GSM1339618	Accession	
																4	





Gene Relationships





Conclusion

- Identifying patients with OA purely on the basis of differential gene expression and clustering methods is not very accurate
- appropriate method (91% accuracy) Outcomes indicate application of rule-based clustering is a more
- More accurate rule-based clustering methods may allow for identification of pathogenetically or therapeutically relevant genetic targets

Bibliography

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https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2694558/

http://genemania.org/