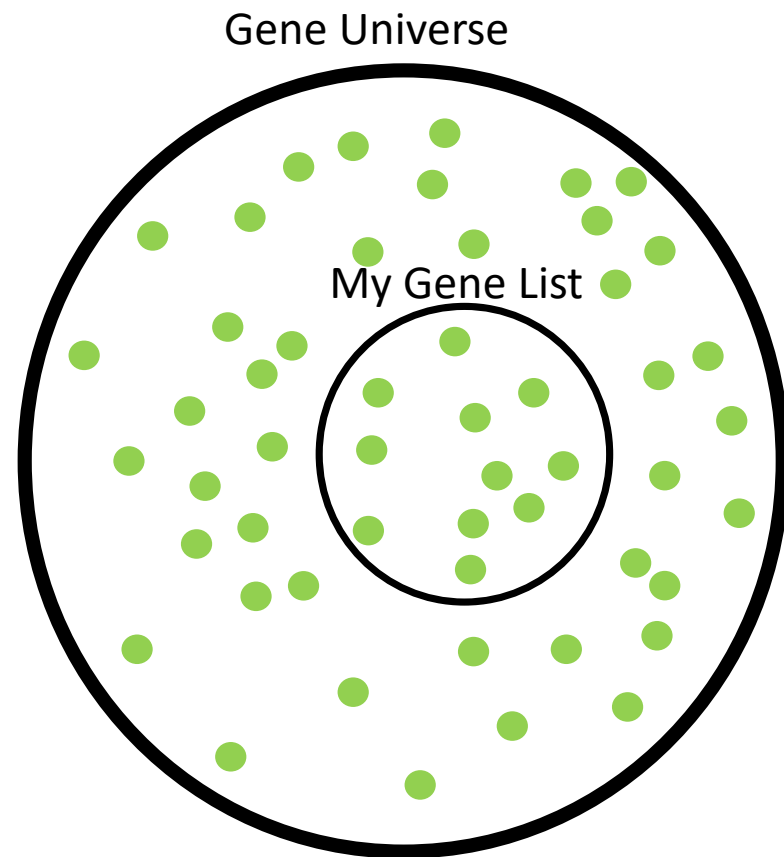
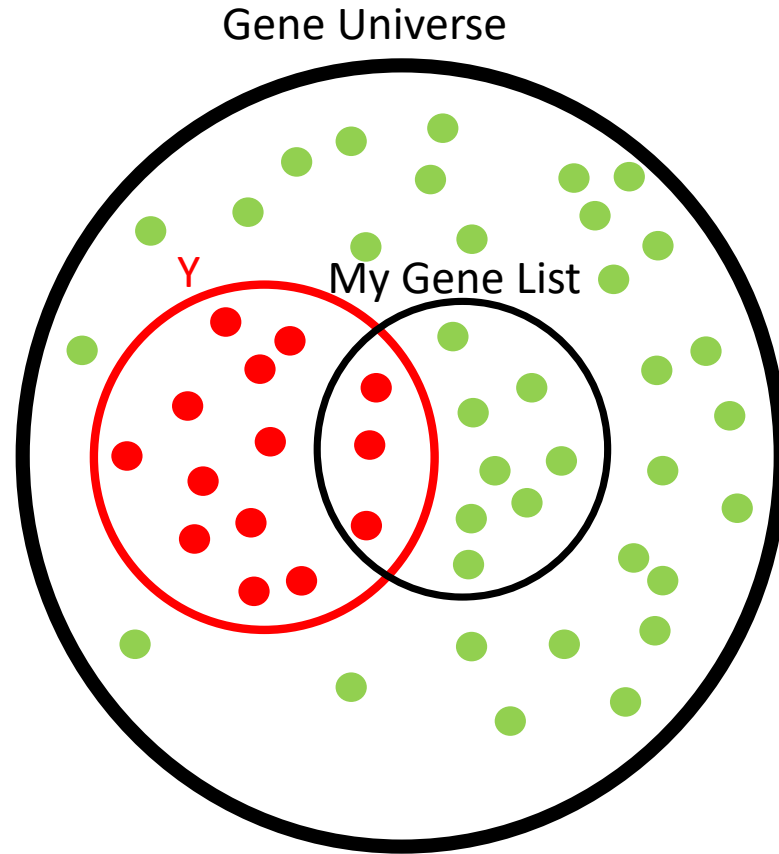


Overrepresentation Test



Overrepresentation Test

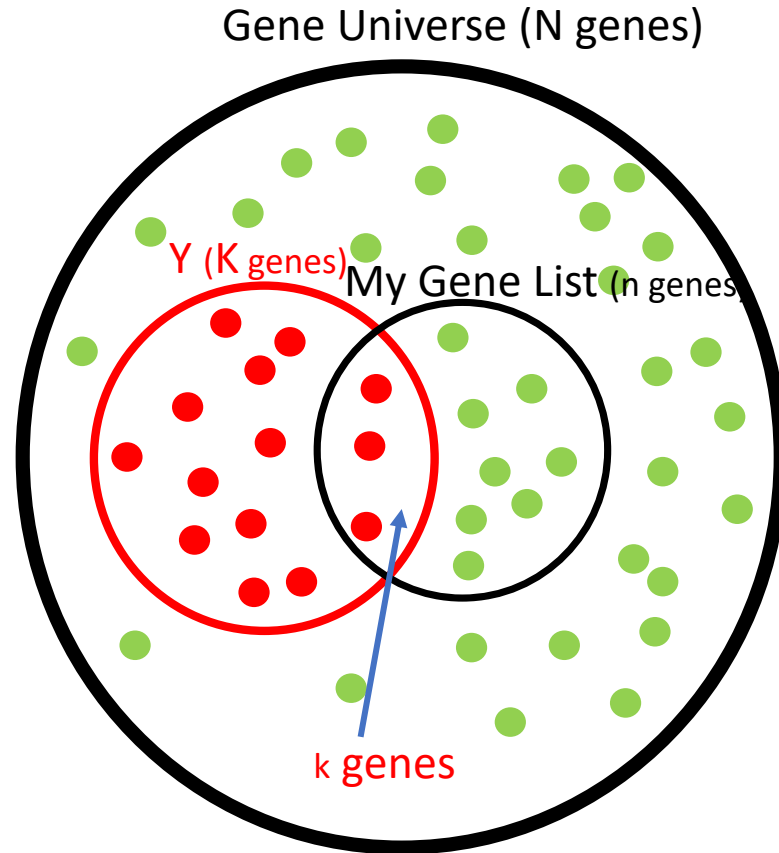


A subset of genes are annotated with **category Y**

My Gene List has 11 genes and **3 are annotated with category Y.**

Are **category Y genes** overrepresented in My Gene List?

Overrepresentation Test



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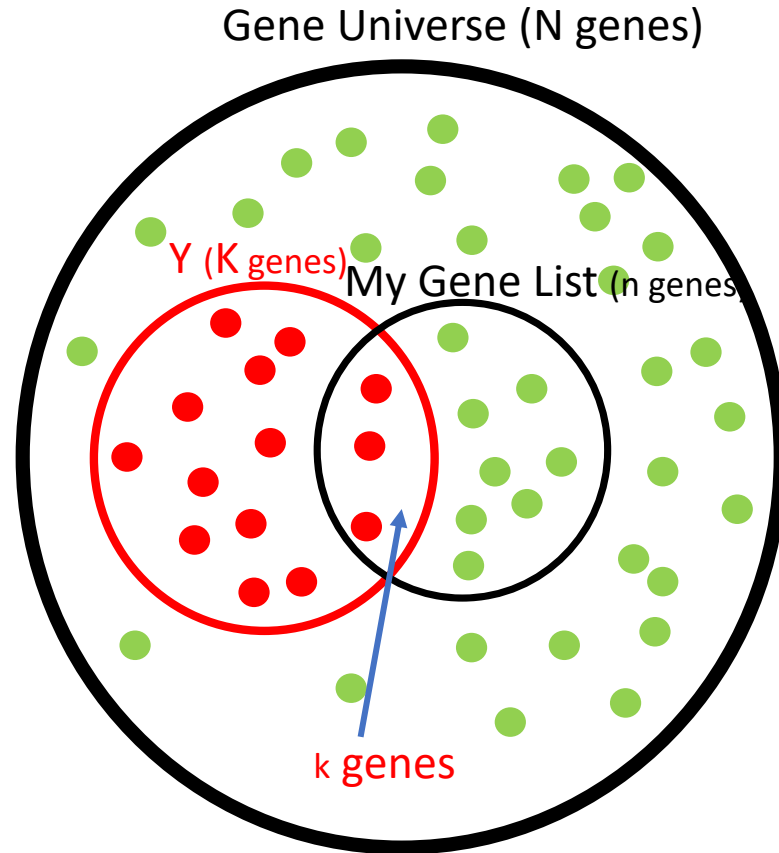
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Hypergeometric Distribution

Probability of My Gene List having **k red genes** if it is a random sample of size **n** from the Gene Universe (size **N**)

$$P(X = k) = \frac{\binom{K}{k} \binom{N-K}{n-k}}{\binom{N}{n}}$$

Overrepresentation Test



P-value: Probability of having **k or more red genes** by chance:

If this is small, our result is significant

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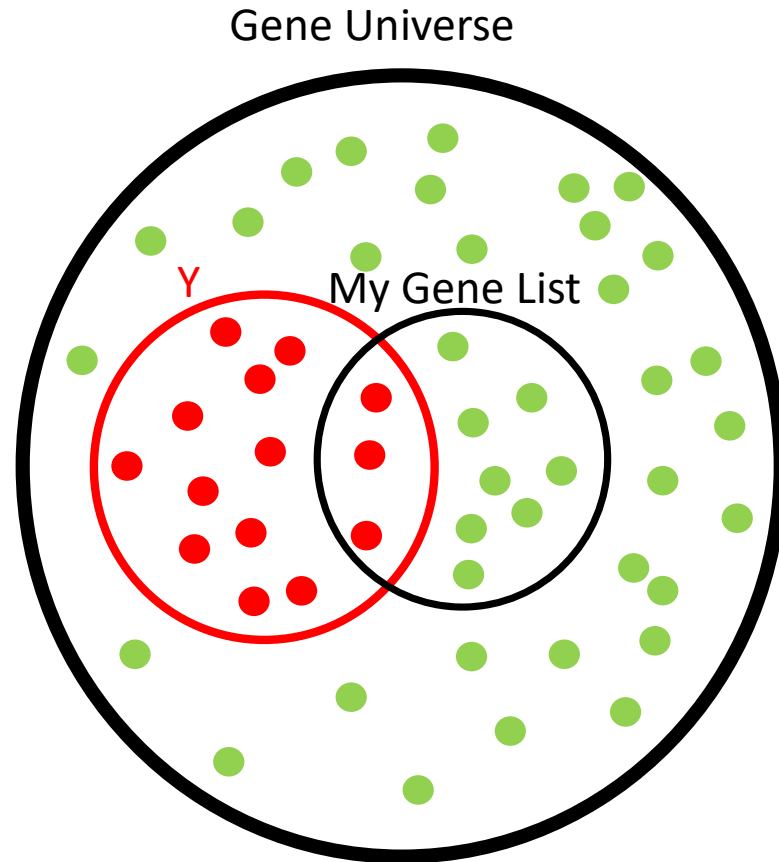
Hypergeometric Distribution

Probability of My Gene List having **k red genes** if it is a random sample of size n from the Gene Universe (size N)

$$P(X = k) = \frac{\binom{K}{k} \binom{N-K}{n-k}}{\binom{N}{n}}$$

$$P(X \geq k | n; N; K) = \sum_{i=k}^n \frac{\binom{K}{i} \binom{N-K}{n-i}}{\binom{N}{n}}$$

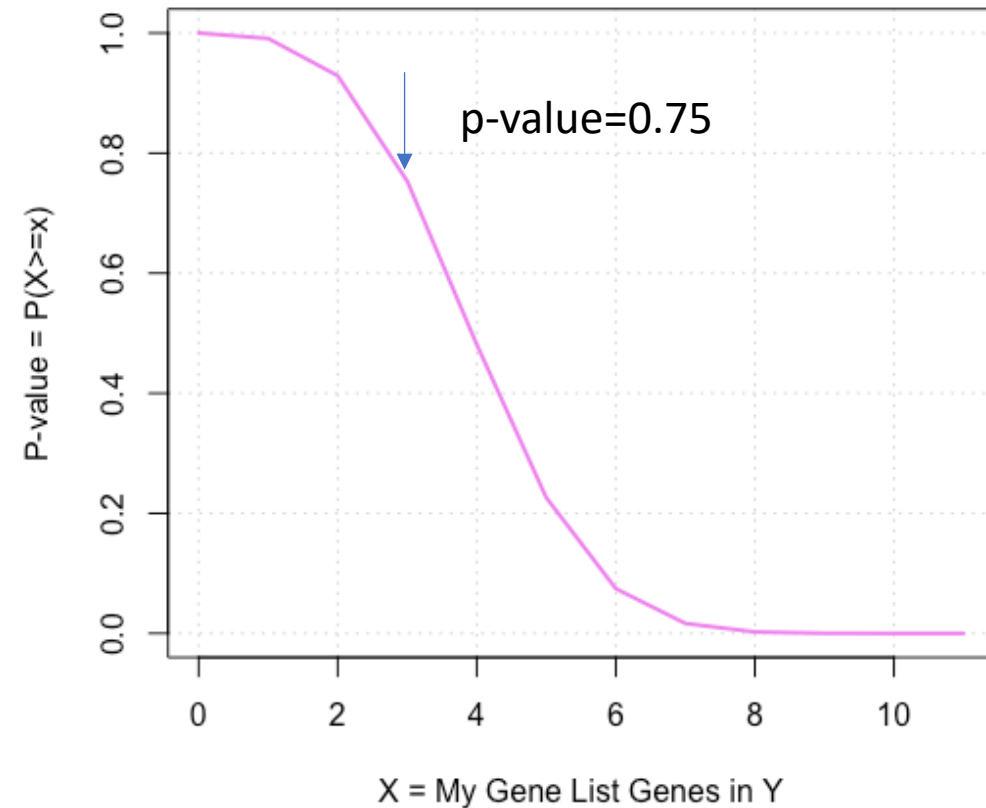
Overrepresentation Test



P-value: Probability of having **k or more red genes** by chance:

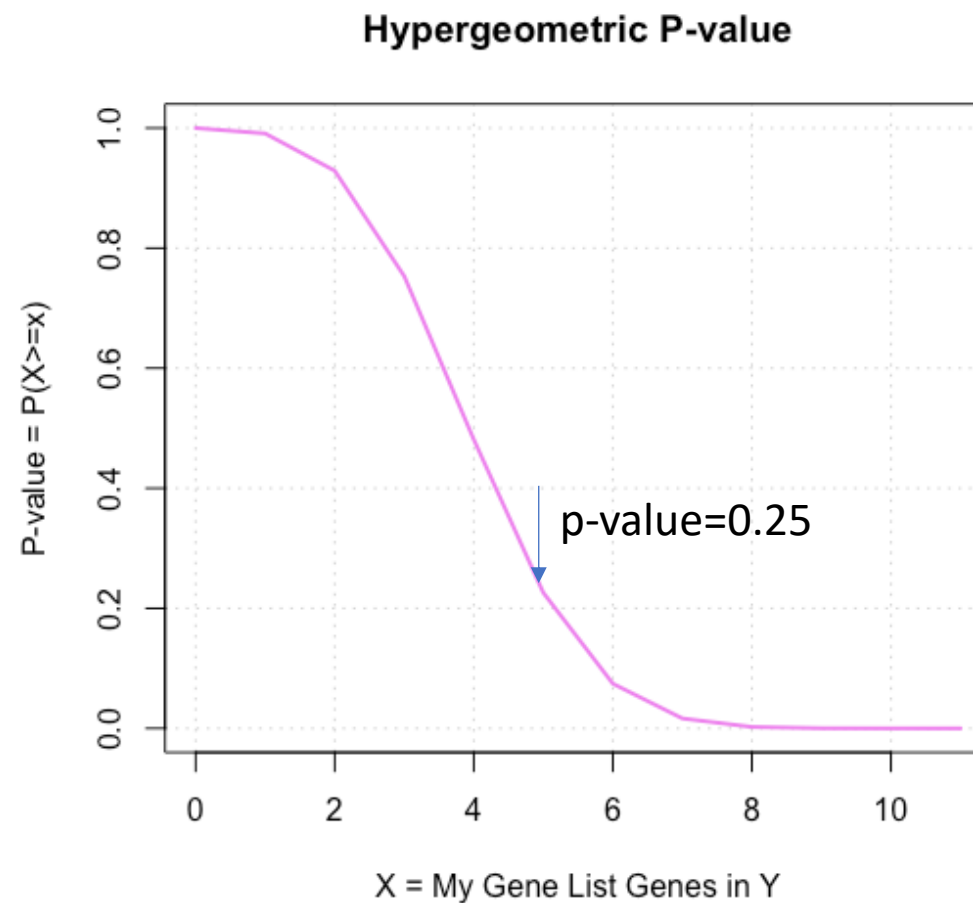
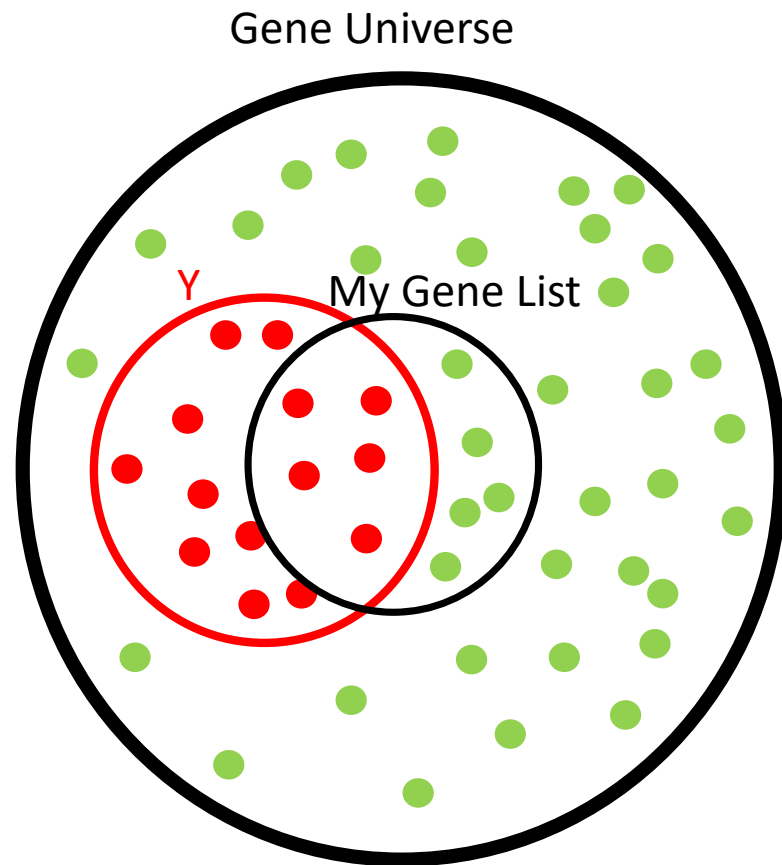
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Hypergeometric P-value

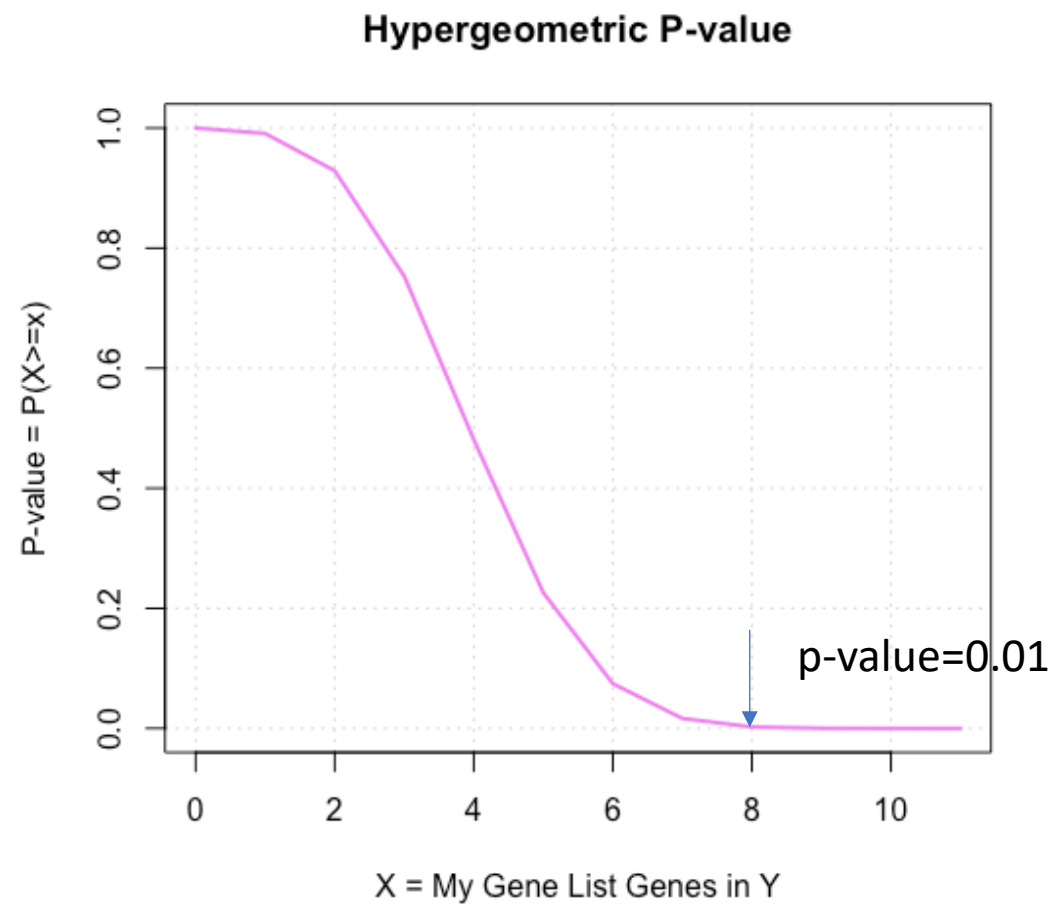
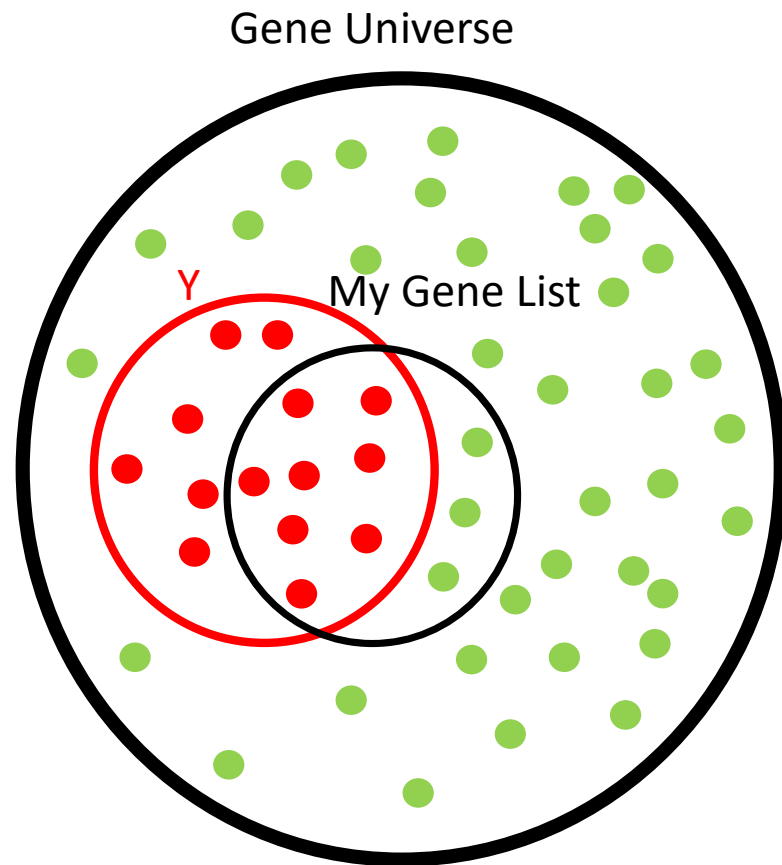


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Overrepresentation Test



Overrepresentation Test



GO Enrichment with ClusterProfiler (R)

```
ego = enrichGO(geneList,  
  OrgDb,  
  ont = 'BP',  
  pvalueCutoff = 0.05)
```

Fraction of genes in
My Gene List with the
Annotation

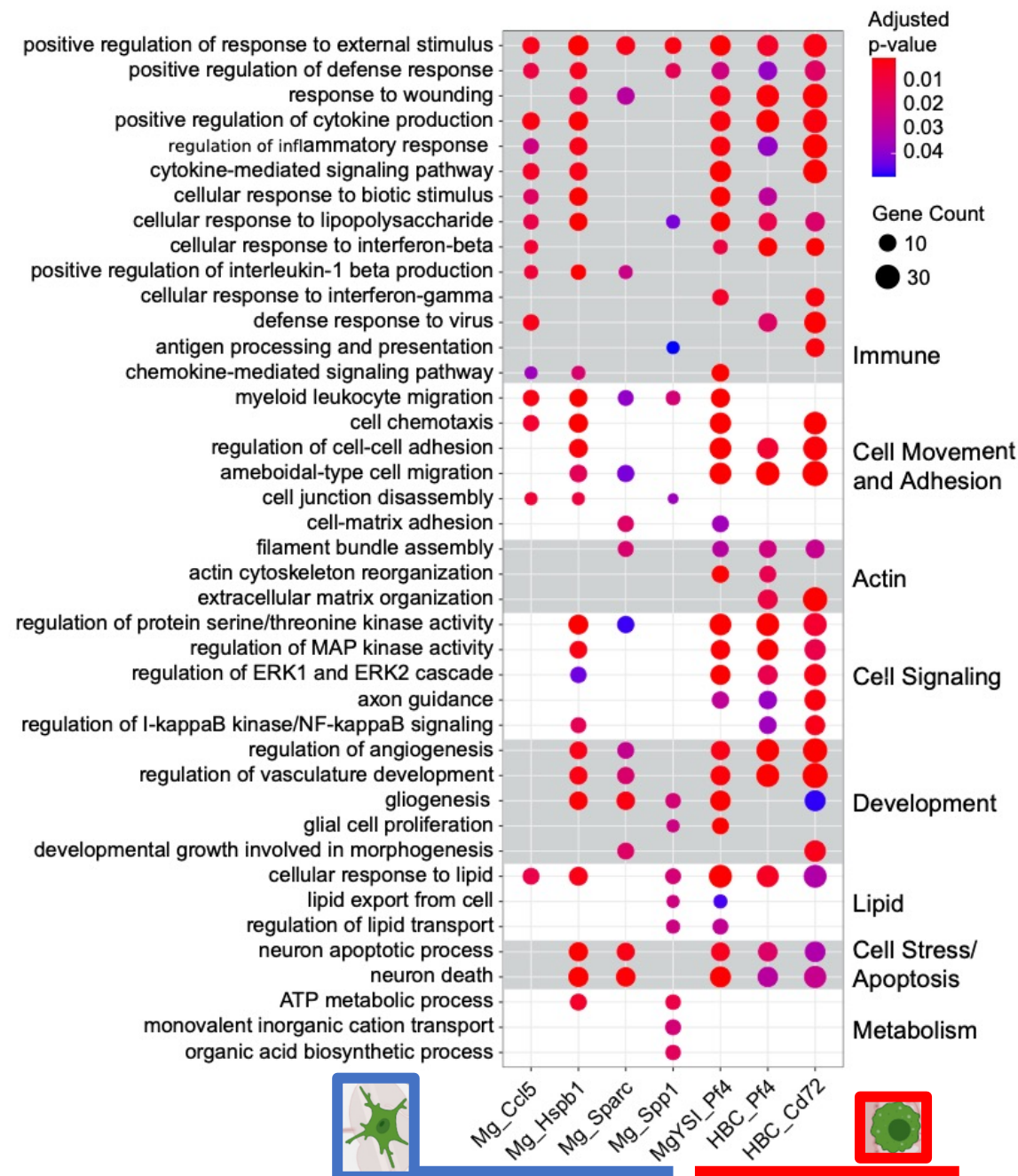
Fraction of genes in
the database with the
Annotation

ID	Description	GeneRatio	BgRatio	pvalue	p.adjust	geneID
GO:0007059	chromosome segregation	53/467	309/16976	3.60E-27	1.36E-23	Birc5/Ube2c/ Top2a/Mki67/ ...
GO:0000070	mitotic sister chromatid segregation	34/467	147/16976	4.02E-22	7.61E-19	Birc5/Ube2c/ Cdca8/Nusap 1/...

...

Next, visualize many 100's of results

GO Enrichment for Macrophage subset Marker Genes



We find mostly overlapping functions for Brain Microglia and Placental Hofbauer cells

Do these cells respond similarly to Maternal Obesity?