Módulo: Expresión diferencial

Bioinformática y Estadística 2

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Día 2

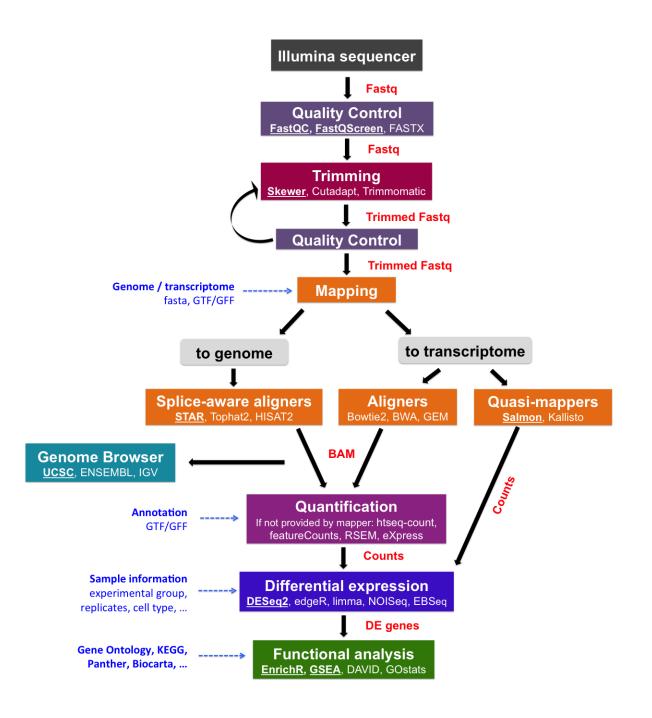
- Gene Set Enrichment Analysis (GSEA) -Análisis funcional
- Información general
- Ejemplos de graficas



Pipeline bioinformática

Dónde estamos...

mRNA-Seq data analysis workflow "https://biocorecrg.github.io/RNAseq_course_2 019/workflow.html"

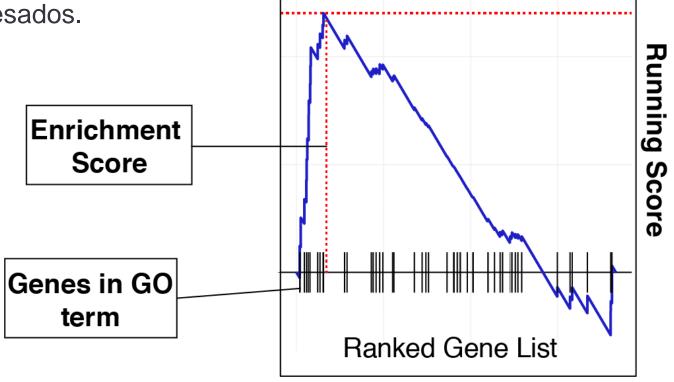


Gene Set Enrichment Analysis (GSEA)

Análisis de funcional de genes expresados.

- Comparación con una distribución hipergeométrica.
- Ejemplo con m&ms

https://www.youtube.com/watch?v=udyAvvaMjfM



GSE function workflow

Input

- gene_list = Ranked gene list (numeric vector, names of vector should be gene names)
- GO_file= Path to the "gmt" GO file on your system.
- pval = P-value threshold for returning results

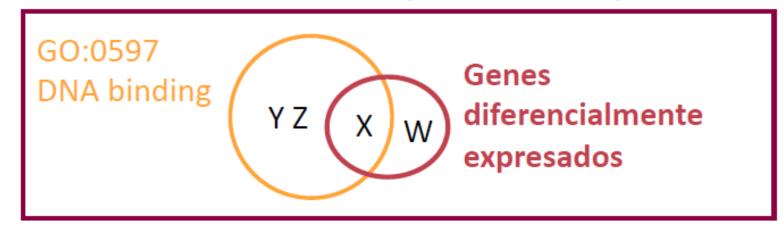
Paquetes

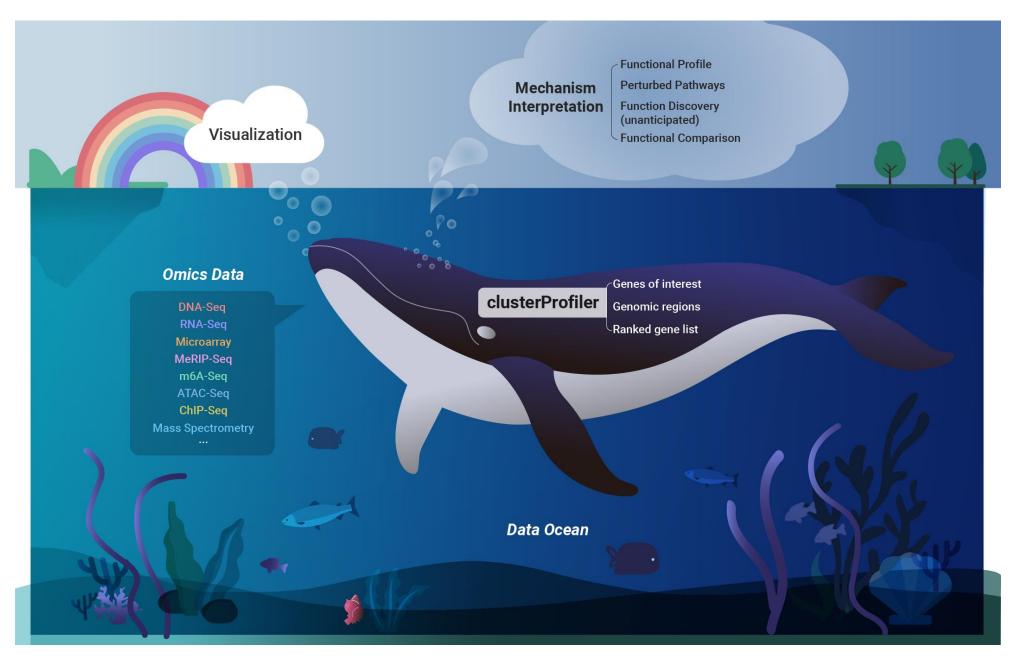
• fgsea, clusterProfiler, EnrichR, Gprofiler2, GORilla, topGO, AnnotationHub, goseq

Comparaciones

- Universo
- Genes expresados

Universo: Genes de ratón expresados en experimento





https://github.com/YuLab-SMU/clusterProfiler

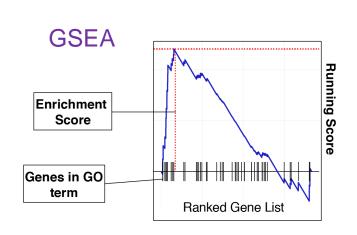
GOterm

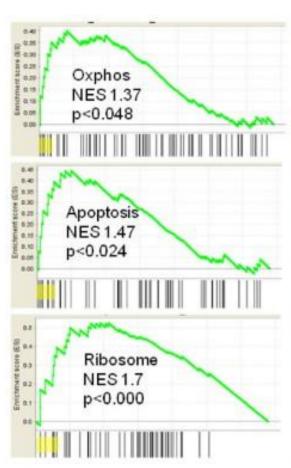
- BP: Biological process
 - · Oligopeptide transport, response to abiotic stimulus, cell cycle
- MF: Molecular function
 - Identical protein binding, endopeptidase activity
- CC : Celular component
 - Apoplast, cell wall, extracellular region, cell surface

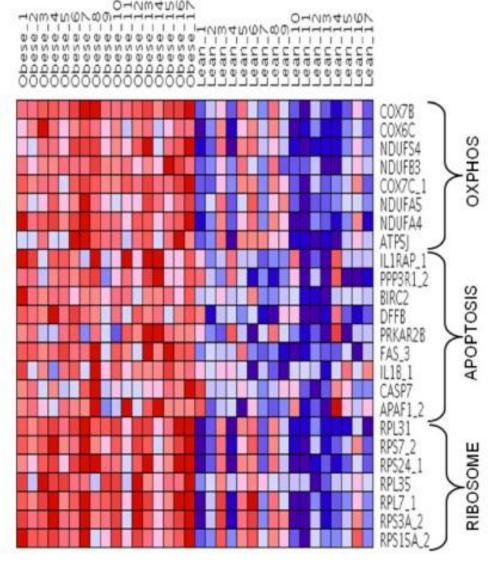
KEGG pathway

Rutas metabólicas / enzimáticas.

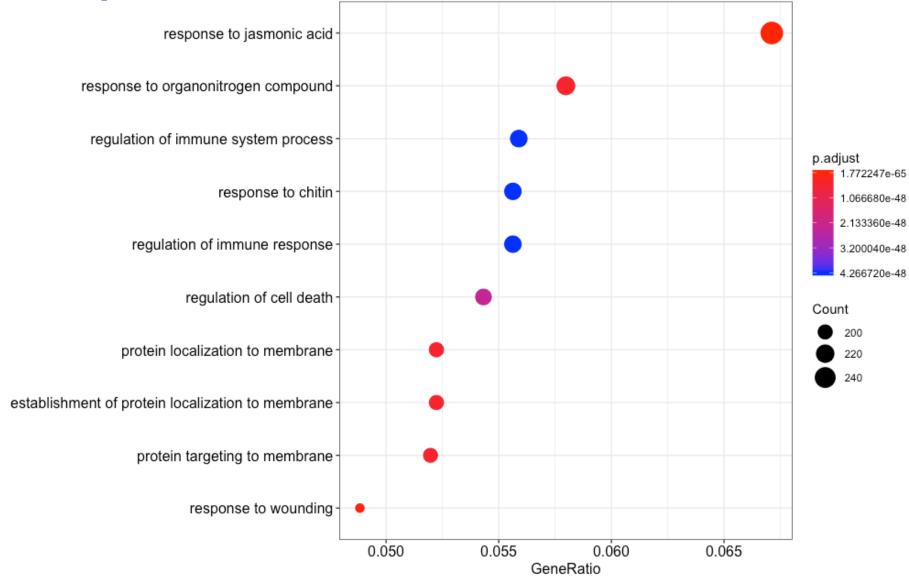
Gene expression profiling in whole blood identifies distinct biological pathways associated with obesity





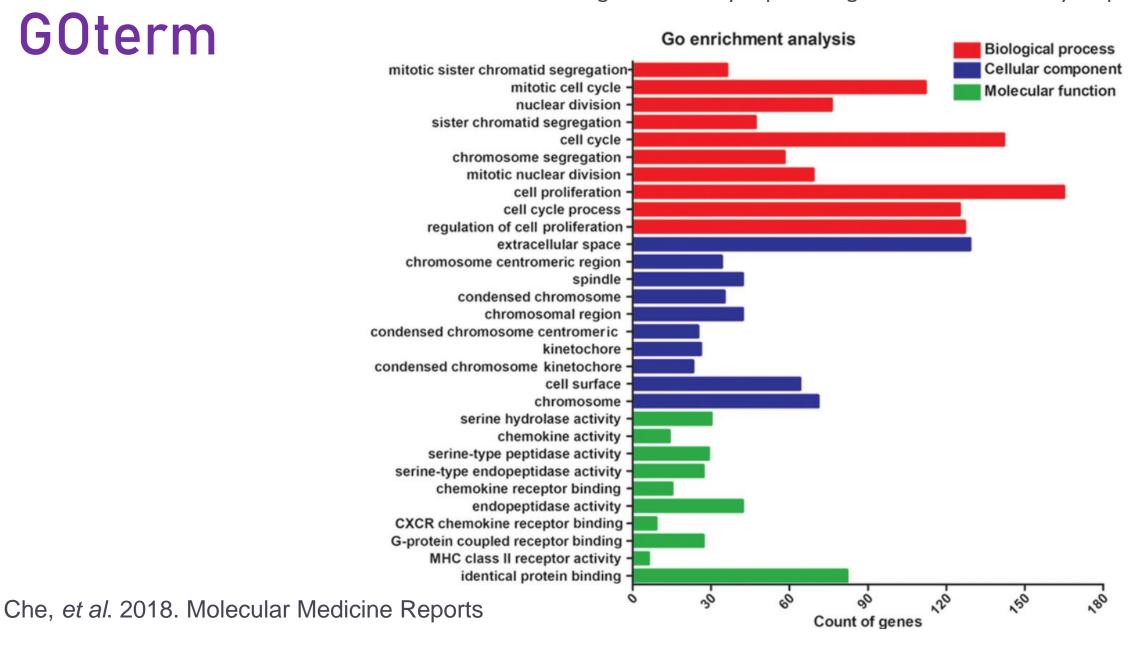


Tus primeros pasos

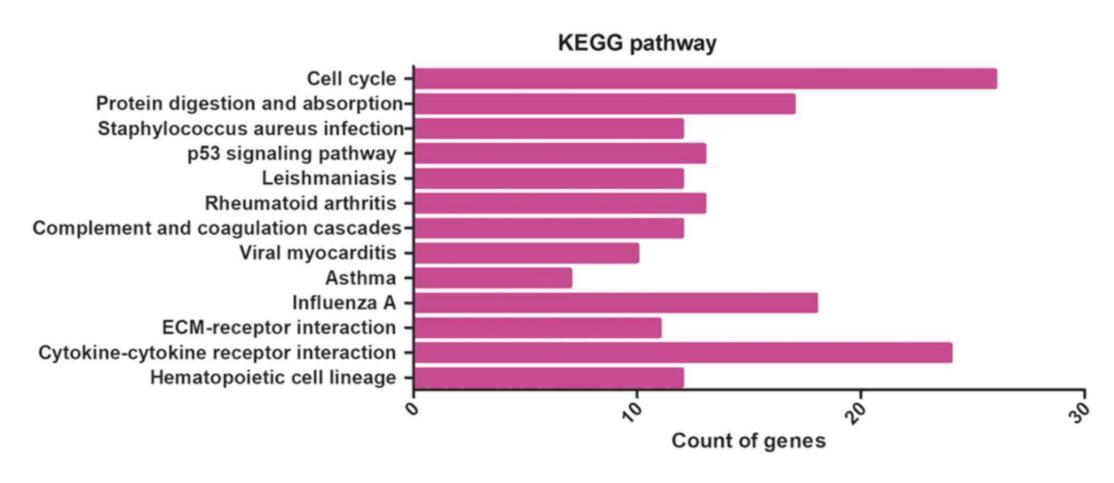


Identification of putative drugs for gastric adenocarcinoma utilizing differentially expressed genes and connectivity map

GOterm



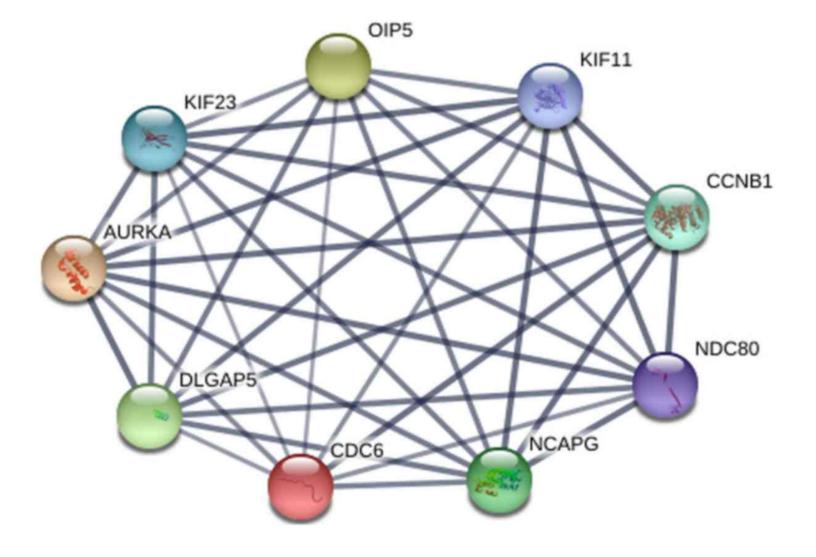
KEGG pathway



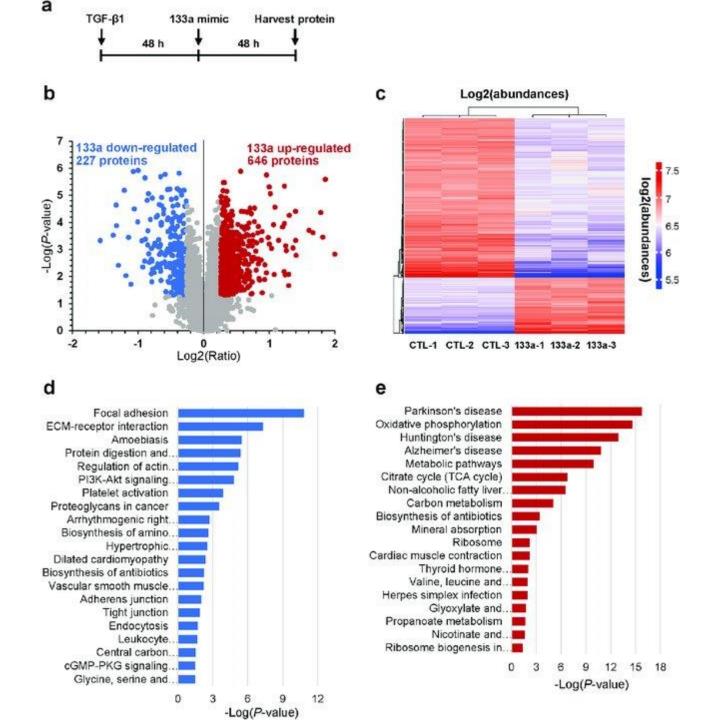
Redes

Identification of putative drugs for gastric adenocarcinoma utilizing differentially expressed genes and connectivity map





Transforming growth factor (TGF)- β 1-induced miR-133a inhibits myofibroblast differentiation and pulmonary fibrosis



Práctica

https://github.com/EveliaCoss/RNAseq_classFEB2023/blob/main/RNA_seq/README.md#practica4