

Building monocyte derived macrophage behavior

Experimental steps

This tab includes data related to wet-lab experiments, such as laboratory protocols and FASTQ files, done on monocyte derived macrophages to calibrate and validate intracellular models of monocyte derived macrophages.

Bioinformatics steps

This tab includes data related to bioinformatics analysis, such as scripts and pipelines, used to analyze data gathered in the wet-lab part.

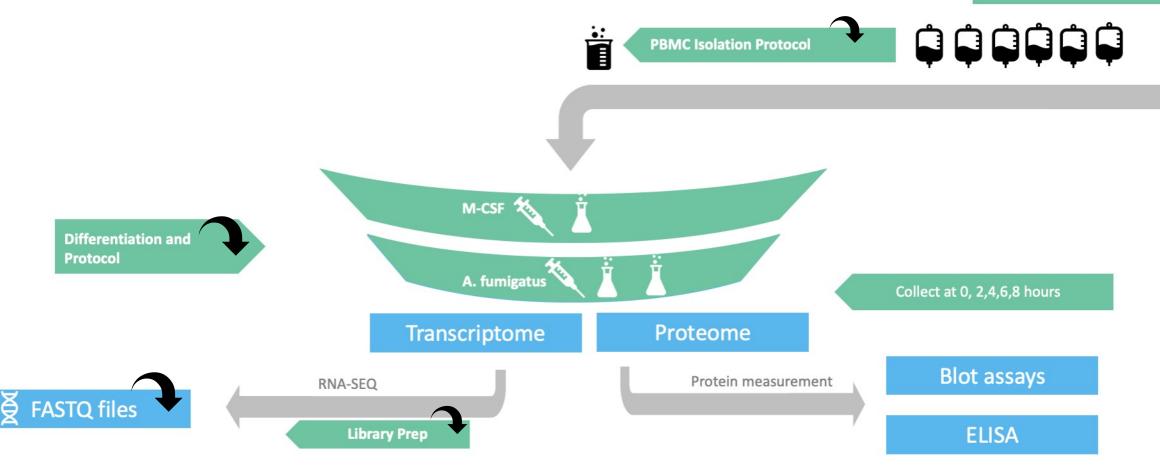
Modeling steps

This tab includes data related to the intracellular mathematical model of macrophages, including literature used to build the models, links to scripts, results, etc.

Experimental flow for coculture experiment

Sample metadata

Monocyte derived macrophages



Bioinformatics pipeline

FASTQ files for macrophages



Adapters FASTA file

Get reference genome

MultiQC posttrim

Analysis scripts





Cut adapters (Trimmomatic)

STAR aligner (index and align)

Qualimap, SAM tools MultiQC

DESEQ2, CHEA3, enrichment

Aligned reads



