

# Analysis of Parasitic Protozoa at the Single-cell Level using Microfluidic Impedance Cytometry

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The dataset contains the experimental data and Matlab codes needed to generate the figures of the article. In particular:

- “Figure4.mat” and “Figure4.m”, corresponding to the experimental data and code, respectively, to plot figure 4;
- “Figures\_5\_S2.mat”, “Figure5.m” and “FigureS2.m”, corresponding to the experimental data and code, respectively, to fit the relaxation curves and plot both figure 5 and S2;
- “Figure6.mat” and “Figure6.m”, corresponding to the experimental data and code, respectively, to plot figure 6;
- “Figure7.mat” and “Figure7.m”, corresponding to the experimental data and code, respectively, to plot figure 7;
- “Figure8.mat” and “Figure8.m”, corresponding to the experimental data and code, respectively, to plot figure 8;
- “FigureS1.mat” and “FigureS1.m”, corresponding to the experimental data and code, respectively, to plot figure S1;
- “FigureS3.mat” and “FigureS3.m”, corresponding to the experimental data and code, respectively, to plot figure S3;
- “FigureS4.mat” and “FigureS4.m”, corresponding to the experimental data and code, respectively, to plot figure S4;

Extract the “.mat” (data) and “.m” (code) to the same directory. Run the script files in Matlab 2016 (or later) to generate the plots corresponding to each figure.