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# What is wrong with my flow cytometry data?

Hints, tricks and pitfalls

**Webinar 3**  
**04.04.2023**

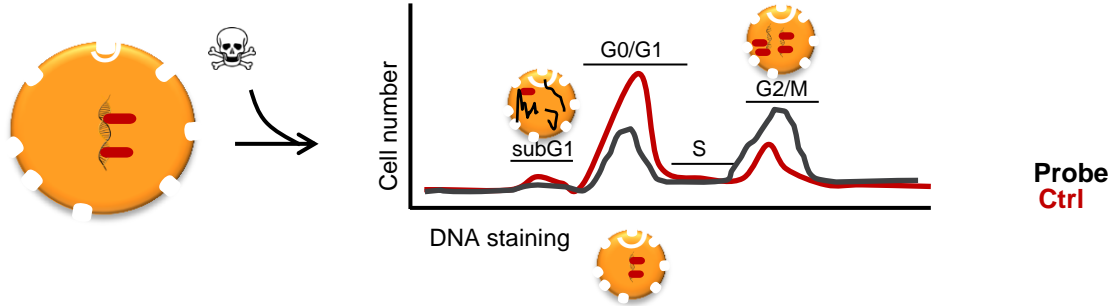


# Cell cycle analysis – simple but complicated?

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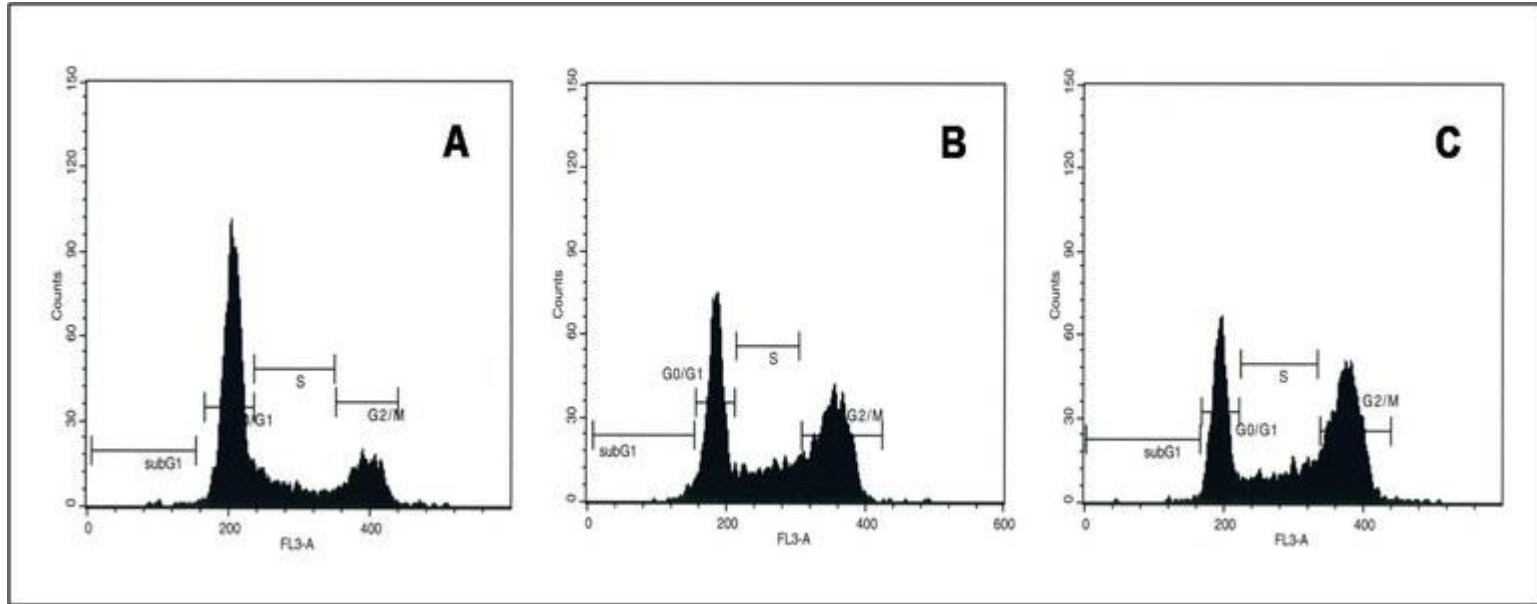
- simple:
  - use of intercalating DNA-Dyes
  - only one colour needed → can be done on nearly any flow cytometer
  - one of first applications of flow cytometry in history
- complicated:
  - many different protocols available
  - small differences in treatment can have big effects
  - RNA, doublets and clumps disturb cell cycle analysis

## principle and general steps

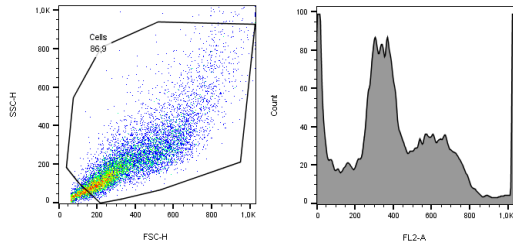


1. Fix cells that have been harvested and washed in PBS in 70% v/v ethanol. Adding the ethanol dropwise to the cell pellet while vortexing will ensure that all cells are fixed and will minimize clumping.
2. Fix cells for 30 min at 4°C, after which wash cells twice in PBS (850 × g). Be careful to avoid cell loss when discarding the supernatants.
3. Treat cells with RNase (50 µL, 100 µg/mL) in order to ensure that only DNA is stained.
4. Add PI (200 µL PI, 50 µg/mL stock solution) immediately before analyzing

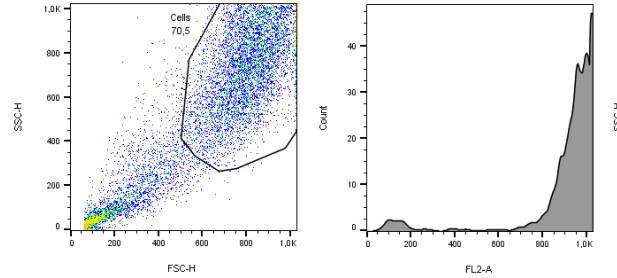
# Example of cell cycle arrest



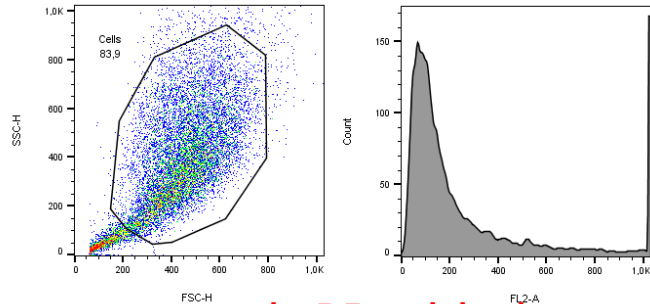
# Cell cycle analysis – What had happened here?



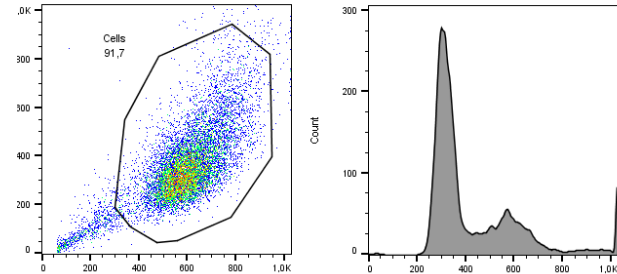
too many dead cells,  
low number of living cell



PMTV for scatter and dye  
not set properly



too much PI added



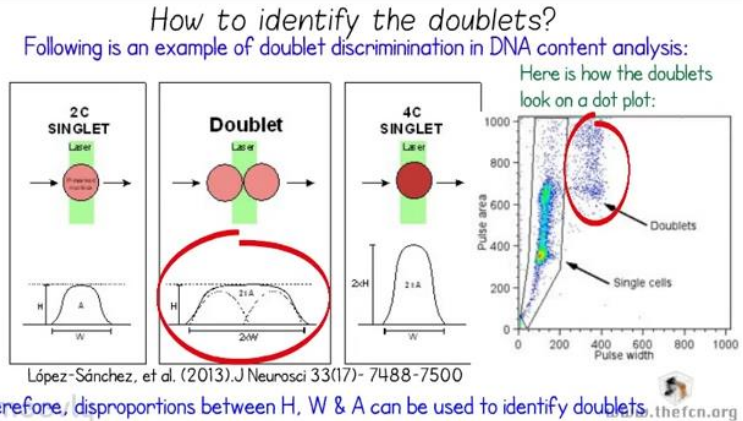
no doublet exclusion

## What to consider?

- Many DNA Dyes available:
  - PI, Hoechst stains, TO-PRO-3, SYTOX, acridine orange, pyronin Y, 7-AAD, DAPI, DRAQ5™, and DRAQ7™
  - bind RNA (PI), others less (7-AAD, DAPI), staining of living cells (DRAQ Dyes)
- PFA-fixation reduces resolution, because of chromatin crosslinking
- Always count cells before DNA staining → DNA-dye – cell – ratio should be stable

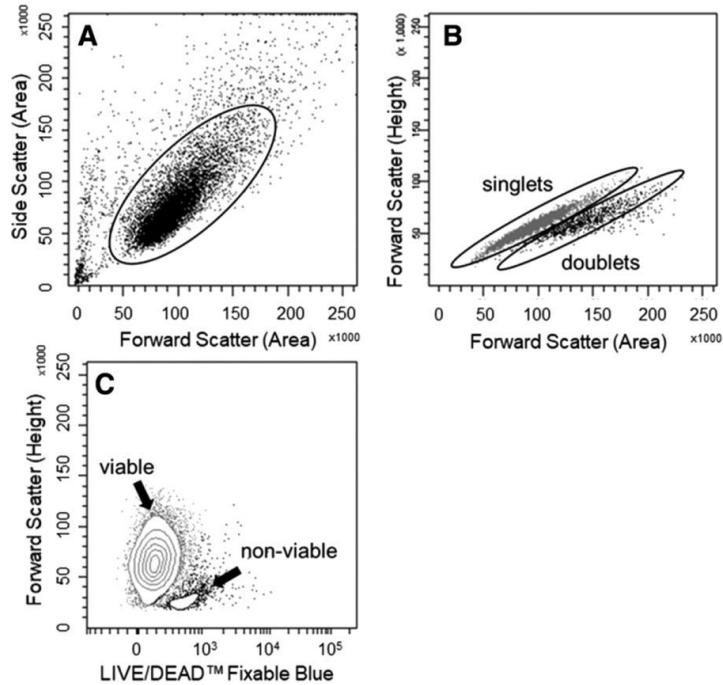
## What to consider?

- exclude doublets
- if possible exclude dead cells  
(scatter or better fixable live-dead-stain)
- lower flow rate results in more accurate acquisition → sharp peaks
- measure & analyze with linear scaling
- record 10.000 – 20.000 cells
- adjust G1 peak to a definite value, e.g. ~ 200, and check for the G2/M to appear at the doubled value , e.g. 400 (doubling of DNA content)



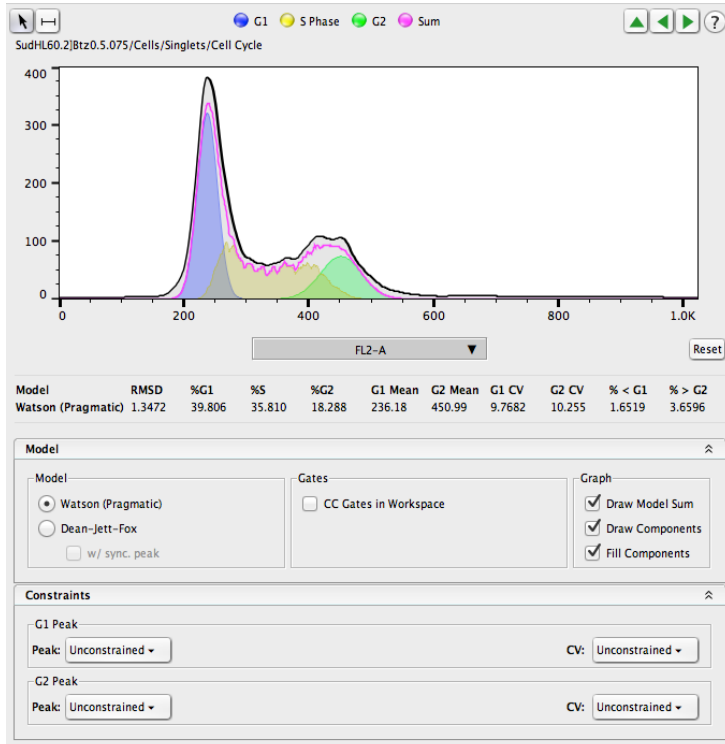


# experimental improvements



- adjust concentration of treatment
  - (too many apoptotic cells)
- include live-dead-staining
- include untreated control (always same confluency)
- maybe starve and synchronize cells before assays start

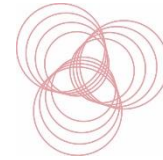
# experimental improvements



- For this purpose specialized programs have been designed such as:
  - ModFit LT™ from Verity Software House (<http://www.vsh.com/products/mflt/mfFeatures.asp>)
  - Multicycle AV™ from Phoenix Flow Systems (<http://www.phnxflow.com/MultiCycle.standalone.html>)

## cell cycle analysis –TAKE HOME MESSAGE

- improve data quality:
  - dye selection and protocol optimization
  - doublet exclusion and low flow rate
  - because of linear scale small differences have big effect
  - stable cell number and dye concentration needed
- FlowJo provides easy analysis tool
  - needs clear G1-peak
  - good untreated control necessary
- ‚easy‘ cell cycle analysis can be more complicated than thought



Thank you for your attention.

See you next month **2th Mai**.

Next topic:

**Aggravating compensation –  
discover errors and correct them**