

## Sorting Request

Please read the general terms of the FACS facility prior to requesting an appointment for sorting:  
<https://www.ccrc-hauner.de/technology-platforms/flow-cytometry/8e97a79744e9af8a>

### 1. User Information:

Name:	Phone:
Department:	E-mail:
Research Group:	

dd/mm/yy

hh:mm

Estimated duration

### 2. Preferred date and time for sorting:

### 3. Sample details:

Cell type:

Number of samples:  Total cell no./sample:  Estimated % of target cells:

### 4. Biological Safety Questionnaire

Please note that sorting of infectious samples or genetically modified organisms of biosafety level S2 or above is not possible.

Do the samples contain primary human cells?

If yes, have donors been screened for infectious pathogens (e.g. HIV, HCV)?

Do the samples contain any known human or murine pathogens?

If yes, specify:

Are the cells genetically engineered?

If yes, provide safety level:

### 5. Sorting details

Fluorochromes: please specify

other: specify

405 nm

488 nm

561 nm

633 nm

Nozzle Size

Target populations sorted into

date previous sorting

Is there a similar set-up available from a previous sorting?

Data export format:

FCS 3.0

Printscreen PDF

other

**Please send the filled report to:**

**[Raffaele.Conca@med.uni-muenchen.de](mailto:Raffaele.Conca@med.uni-muenchen.de)/[Susanne.Wullinger@med.uni-muenchen.de](mailto:Susanne.Wullinger@med.uni-muenchen.de)**