

User Registration

Return to: med.facs@uni-ulm.de

Please provide separate registrations for each individual project

user			
name		position	
phone		email	
group leader		email	
institute		<input type="checkbox"/> Ulm University	<input type="checkbox"/> Ulm Hospital
adress			
cost account			

samples		
species	organ	celltype
celline:		
<input type="checkbox"/>	adherent	<input type="checkbox"/> in suspension
approx. expected cell size :		

According to "BioStoffV" samples are classified as "Risikogruppe 2" or higher?

<input type="checkbox"/>	no	<input type="checkbox"/>	yes: Please complete „samples according Risikogruppe 2“
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Samples according Risikogruppe 2

We are not allowed to sort samples according Risikogruppe 3 or higher !

material

	pathogen	specify:
	patient material	specify:
	untested human material * (e.g. buffy coat,...)	specify:
	other	specify:

*Should be tested for HIV; HBV/HCV or other infectious hazards

According to the German "Gentechnikgesetz" samples are classified as

<input type="checkbox"/>	no
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<input type="checkbox"/>	S1
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<input type="checkbox"/>	S1 (formerly S2)
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<input type="checkbox"/>	S2
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Samples according GenTG

We are not allowed to sort infectious samples or samples according S3 or higher !

project leader (GenTG)	<input type="text"/>
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genetic modification (method/vector/...)

S	<input type="text"/>
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Short project description (general aim,)

<input type="text"/>

Agreements

<input type="checkbox"/>	I accept the <i>user guidelines</i> from the Core Facility Cytometry Uni Ulm
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<input type="checkbox"/>	I accept the <i>service fees</i> from the Core Facility Cytometry Uni Ulm
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<input type="checkbox"/>	I will inform the Core Facility Cytometry Uni Ulm of any changes in classification („BioStoffV“ or „GenTG“) of sample material
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Signature **user**

Signature **Group Leader**

Signature of the **Project Leader** according to the “Gentechnikgesetz”
(*relevant for all samples with genetical modification*)

Core Facility Cytometry

date