

Proteomics Core Facility - Checklist



1. Initial Discussion with PCF Staff

Shortly explain your project to us and get Info about feasible Proteomic workflows



2. Optional: Request an Aliquot of Sample Lysis Buffer in PPMS

We kindly provide you with Lysis buffer. Use this Link to request an Aliquot:
<https://ppms.dkfz-heidelberg.de/PACF/req/?pf=2&project=true&form=18>



3. Request a Sample Block Randomisation in PPMS

Especially for large cohorts of samples it is necessary to randomize the sample order. Use this link to request Block Radomization: <https://ppms.dkfz-heidelberg.de/PACF/req/?pf=2&project=true&form=16>



4. Wait until you receive a randomized sample list

We will send you the block randomized sample list. Please submit your samples in the given order.



5. Complete your sample submission list

Enter valid concentration and volume values and make sure all samples have the same concentration

Sample [N]	Concentration [µg/µl]	Volume [µl]
1	1.5	20
2	1.5	20
3	1.5	20
4	1.5	20
5	1.5	20



6. Create a Project Request in PPMS

Use this link to request a Project in PPMS:
<https://ppms.dkfz-heidelberg.de/PACF/req/?pf=2&project=true&form=14>



7. Wait for a confirmation e-Mail containing your project ID

We are crosschecking your request and will contact you in case anything is unclear.



8. Submit your samles with the project ID

Once you received your Project ID you may submit your samples any weekday between 10 - 11am

DH1234

