

# 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



## 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 samples with the project ID

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

Sample ID	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

