**Beam time: June 2015 (for example)**

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| **Name** | **Badge number**  | **affiliation** | **PI** |
| **Your name** | **XXXXX** | **Department, Lab , organization you work for** | **Your PI’s name or your name if you are the PI** |

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| --- | --- | --- | --- | --- | --- | --- |
| **Previous Projects****(**short but informative title/) | Last beam time used (month/year) | Status (complete/

|  |  |  |
| --- | --- | --- |
|  | abnegate) |  |

 | Outcome of the research(any papers/manuscript/abstract to which conference etc.) |
| **Structural Studies of xxx Complex** | **06/2014** | **complete** | **Publication list** |
| **Structural Studies of xxx**  | **03/2015** | **ongoing** | **Not yet** |
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| --- | --- | --- |
| **Current Project1** **for this beam time****(**short but informative title) | Status (new/ongoing) | Outcome of the research(any papers/manuscript/abstract etc.) |
| **Structural Studies of xxx** | **ongoing** | **Not yet** |

**Sample information (the sample name, MW, origin, amount, buffer composition)****and name of experimental file (***letter and number combination, no more than 6 characters, no “-“,”\_” or other symbol characters***). The file names you give will be used for data collection.**

Lysozyme (14.3 kDa), 7 mg/ml, commercial purchased, Non- hazardous, Non- Radioactive

*Sample note: Proteins*

*Name of experimental file: Lys*

Buffer: 40 mM Na ascorbate, 0.05% NaN3, 150 mM NaCl, 10 mM DTT, pH 4.3

12x167 nucleosome array + GH5 (2655.5 kDa), 4 mg/ml, recombinant expressed and purified

*Sample note:* Complex of DNA and histone proteins +linker

*Name of experimental file: GH5arr*

Buffer: 10 mM HEPES, 0.1 mM EDTA, 0.05% NaN3, 1mM DTT, pH 8.0

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| **Current Project2** **for this beam time** **(**short but informative title) | Status (new/ongoing) | Outcome of the research(any papers/manuscript/abstract etc.) |
|  |  |  |

**Sample information (the sample name, MW, origin, amount, buffer composition)****and name of experimental file (***letter and number combination, no more than 6 characters, no “-“,”\_” or other symbol characters***). The file names you give will be used for data collection.**

UXXXX (10kDa), 4 mg/ml, commercial purchased, Non- hazardous, Non- Radioactive

*Sample note: Protein-Protein complex*

*Name of experimental file: ub*

Buffer: 20 mM TRIS, 150mM NaCl, 0.05% NaN3, 10mM DTT, pH 5.5

EcERA+GNP+RNA Non- hazardous, Non- Radioactive

EcERA (33.68 kDa), 2.2 mg/ml, recombinant expressed and purified

GNP (521.18 kDa), 10mM, commercial purchased

RNA oligo (3676.2 kDa), commercial purchased

*Sample note: Protein-RNA complex*

*Name of experimental file: EGR*

Buffer: 50 mM Tris, pH 8.0, 150 mM NaCl, 3mM BME