| Experiment/BL parameters (Most BL parameters are the same as MX) |
| --- |
| Experiment_type (chip/injector) |
| Sample support |
| Jet_material |
| Mono_stripe |
| Beam_Size |
| Temperature |
| See_gleb_data_model for extensions |

| Data collection |
| --- |
| Exp.time |
| Transmission |
| Flux |
| Repetition rate |
| Energy_bandwidth |
| Beam_center_x |
| Beam_center_y |
| Det_distance |
| **Sample/crystal slurry** |
| Protein |
| Buffer |
| Avg_xtal_size |
| Xtal_concentration |

| Crystfel_merging |
| --- |
| Processing_ids |
| Cc* |
| Rsplit |
| Number of indexed images |

| Crystfel_processing |
| --- |
| Hits (integer) |
| Indexed(Integer) |
| Lattice type / point group |
| Unit_cell_params (a,b,c, alpha,beta, gamma) |
| Min-max-avg-std_dev |
| Lattice_type |
| Resolution_estimate |
| **PyFAI_parameters** |
| Polarization_value |
| Sigma_value |
| Mask_file_name |
| Error_model |
| Number of hits |
|  |