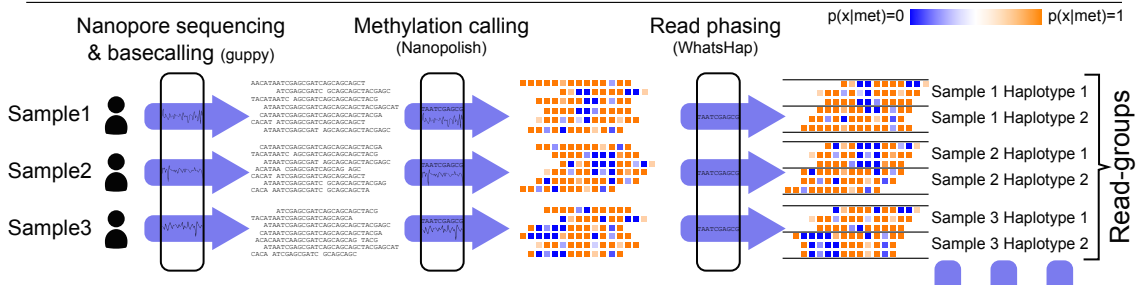


# Data preparation



## MetH5 and pycoMeth

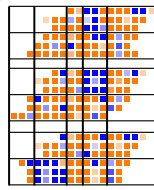
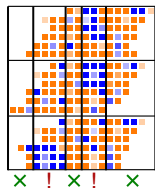
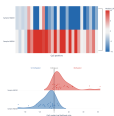
**pycoMeth**

Reporting

Differential Methylation

Methylome Segmentation

HTML report



**MetH5 CLI**  
store as MetH5 files  
incl. read-groups

Sample 1.m5

Sample 2.m5

Sample 3.m5

**MetH5**