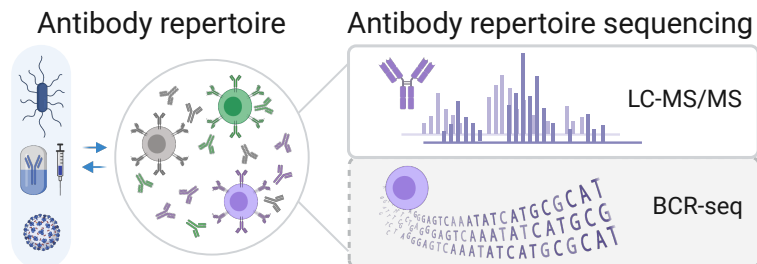
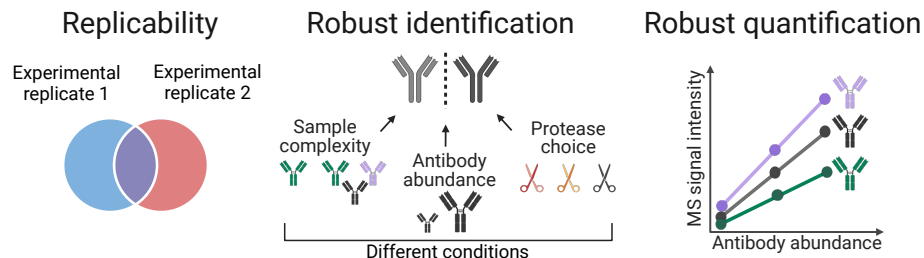


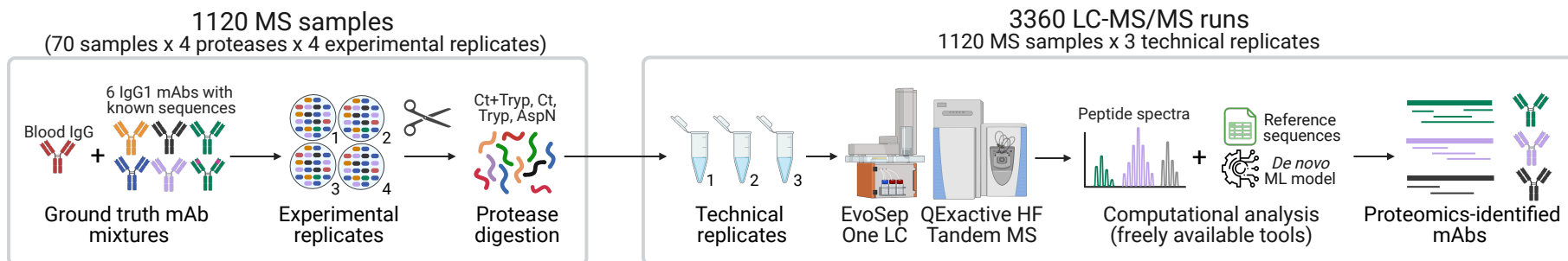
a Antibody repertoire sequencing



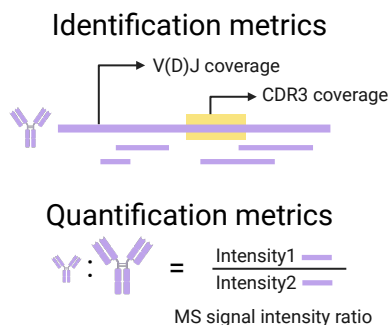
b Lack of Ab-seq benchmarking studies



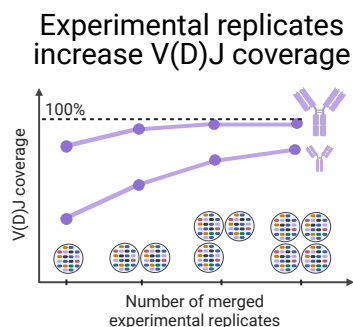
c Antibody LC-MS/MS benchmarking workflow



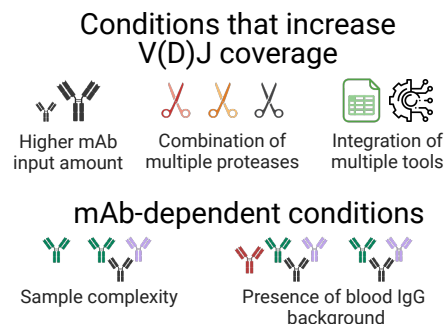
d Benchmark criteria



e Replicability



f Identification and quantification robustness



MS intensity-mAb input concordance is mAb- and protease-dependent

