DESIGNS, DATA & CODE for Serrano-Blesa, E., Porter, A., Lendrem, D.W., Pitzalis, C., Barton, A., Treumann, A. and Isaacs, J.D., 2021. Robust Optimization of SWATH-MS workflow for human blood serum proteome analysis using a Quality by Design approach.

<https://www.researchsquare.com/article/rs-347514/v1>

* Screening design



* Data and code generating Bayesian Estimates.jmp



* Code generating Figure 1.



* Data and code generating Figure 2 and Figure 3.



* Validation data generating Figure 4.