

Description of Additional Supplementary Files

Supplementary Data 1. Example bash script demonstrating how to convert MiXCR output into a format compatible with the Immcantation R package.

Supplementary Data 2. Schematic overview of the CLUSTER consortium data access workflow, illustrating step-by-step navigation from the European Genome-phenome Archive (EGA) study and dataset pages to the CLUSTER consortium website and the formal data access request process.