

## TCR & BCR Repertoire Analysis — Hands-on

Develop an intuitive, analysis ready view of T cell and B cell receptor repertoires. This module covers biological foundations, core repertoire data structures, clonotype definitions, diversity metrics and conceptual workflows for tracking clonal dynamics across time, tissue and clinical groups.

# TCR & BCR Repertoire Analysis

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### Session 1

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## Repertoire Biology & Sequencing Overview

TCR and BCR biology and why repertoires matter

[V\(D\)J recombination concepts](#) [clonal selection and expansion](#) [links to infection, vaccination and cancer](#)

Sample types and repertoire sequencing strategies (conceptual)

[peripheral blood, tissue and tumour samples](#) [bulk versus single cell oriented view](#) [alpha beta TCR, gamma delta TCR and BCR heavy light chains](#)

High level repertoire data outputs and file structures (orientation)

annotated rearrangements table concept V, D, J calls  
and CDR3 sequences counts, frequencies and  
metadata columns

## Session 2

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### Clonotypes, CDR3 & Diversity Metrics

Defining clonotypes from rearrangements (conceptual options)

CDR3 sequence based grouping V and J gene usage  
considerations nucleotide versus amino acid level  
definitions

Basic repertoire summaries and diversity metrics (orientation)

richness and clonality ideas Shannon and Simpson  
style indices (conceptual) Gini and related inequality  
views

CDR3 length, composition and motif oriented summaries

length distributions and usage plots amino acid  
property based views motif level orientation for  
public specificities

## Session 3

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### Comparing Repertoires & Clonal Dynamics

Concepts for comparing repertoires across samples or groups

overlap and shared clonotypes distance based  
comparisons (orientation) public versus private  
repertoire concepts

Tracking clonal expansion and contraction over time (conceptual)

following selected clonotypes across visits link to  
vaccination or treatment time points visual motifs for

### **expansion trajectories**

Contextualising patterns in infection, vaccination and oncology examples

**acute versus chronic infection patterns** **vaccine responders versus non responders (conceptual)**  
**tumour infiltrating lymphocyte repertoires (orientation)**

### **Session 4**

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## **Mini Project: Repertoire Storyboard**

From annotated repertoire table to core plots and summaries (conceptual workflow)

**diversity and clonality overview** **CDR3 length and VJ usage style plots** **top clonotypes table with simple annotations**

Building a simple clonal dynamics storyboard (conceptual)

**selecting clonotypes of interest** **depicting change across time or conditions** **relating patterns to clinical or experimental events**

Summarising findings for immunology and translational teams

**short text narrative with key figures** **tables and plots suitable for slide decks** **handoff to neoantigen, biomarker and trial modules**