

excellence and accuracy since the field's inception. Honored in 2023 with the "Cristal collectif du CNRS" award for its high-impact support to research, IMGT® continues to be at the forefront of immunogenetic analysis.

We provide an integrated knowledge system for <u>sequences</u>, <u>genes</u> and <u>structures</u> of the immunoglobulins (**IG**) or antibodies, T cell receptors (**TR**) and major histocompatibility proteins (MH) of the adaptive immune responses, as well as of other proteins of the IG superfamily and MH superfamily of vertebrates and invertebrates. Our online resources, regularly updated to ensure the highest standards of excellence, are specifically designed to elucidate the complexities of the adaptive immune response







Databases

Online tools

Web pages

These resources are accessible to researchers, educators, and anyone with a scientific interest in this area.

Our Core Research Areas are:

- Deciphering of the IG and TR genes on the genomic loci of jawed vertebrates
- Exploring the expressed IG and TR repertoires
- Analysing the 2D and 3D structures of the adaptive immune proteins repertoires



















