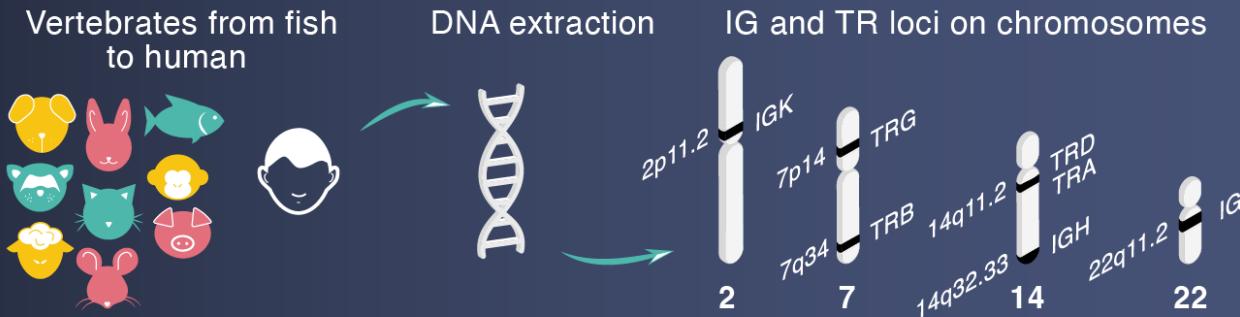


# Deciphering the IG and TR locus, genes and alleles in genomes of jawed vertebrates

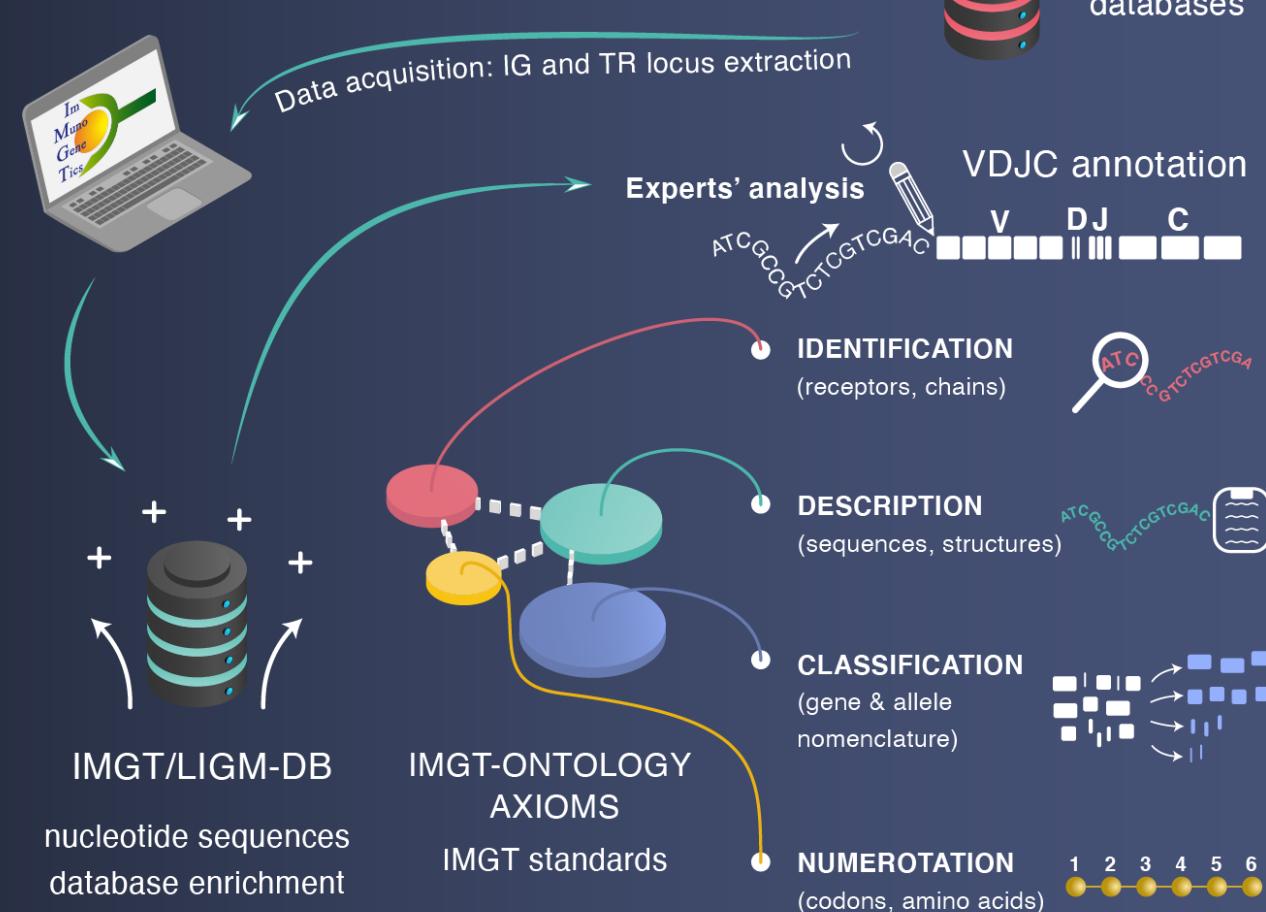


<http://www.imgt.org>

## General context: vertebrates genome assembly



## IMGT biocuration pipeline



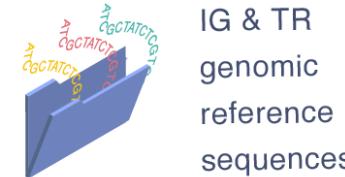
## Results: enriching IMGT core data sets

Databases enrichment, nomenclature assignment, standards enhancement

### IMGT repertoire

- Locus representations
- Gene tables
- Protein display
- Alignments of alleles

### IMGT reference directory



### IMGT/GENE-DB

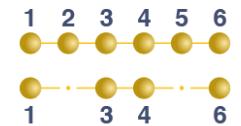


IG & TR genomic reference sequences

## IMGT tools and databases using IMGT reference directory



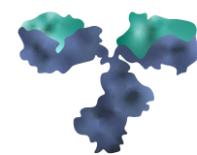
IMGT/V-QUEST &  
IMGT/HighV-QUEST



IMGT/  
DomainGapAlign



IMGT/  
Collier-de-Perles



IMGT/2Dstructure-DB  
& IMGT/3Dstructure-DB

## Applications

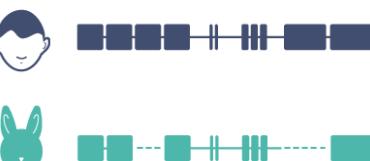
### - IG & TR sequence comparison

ATCGT Polymorphisms /  
ATGGT mutations



Haplotypes

### - Comparative genomics



Evolutionary  
studies



Newly sequenced  
species expertise