

Next Generation Sequencing

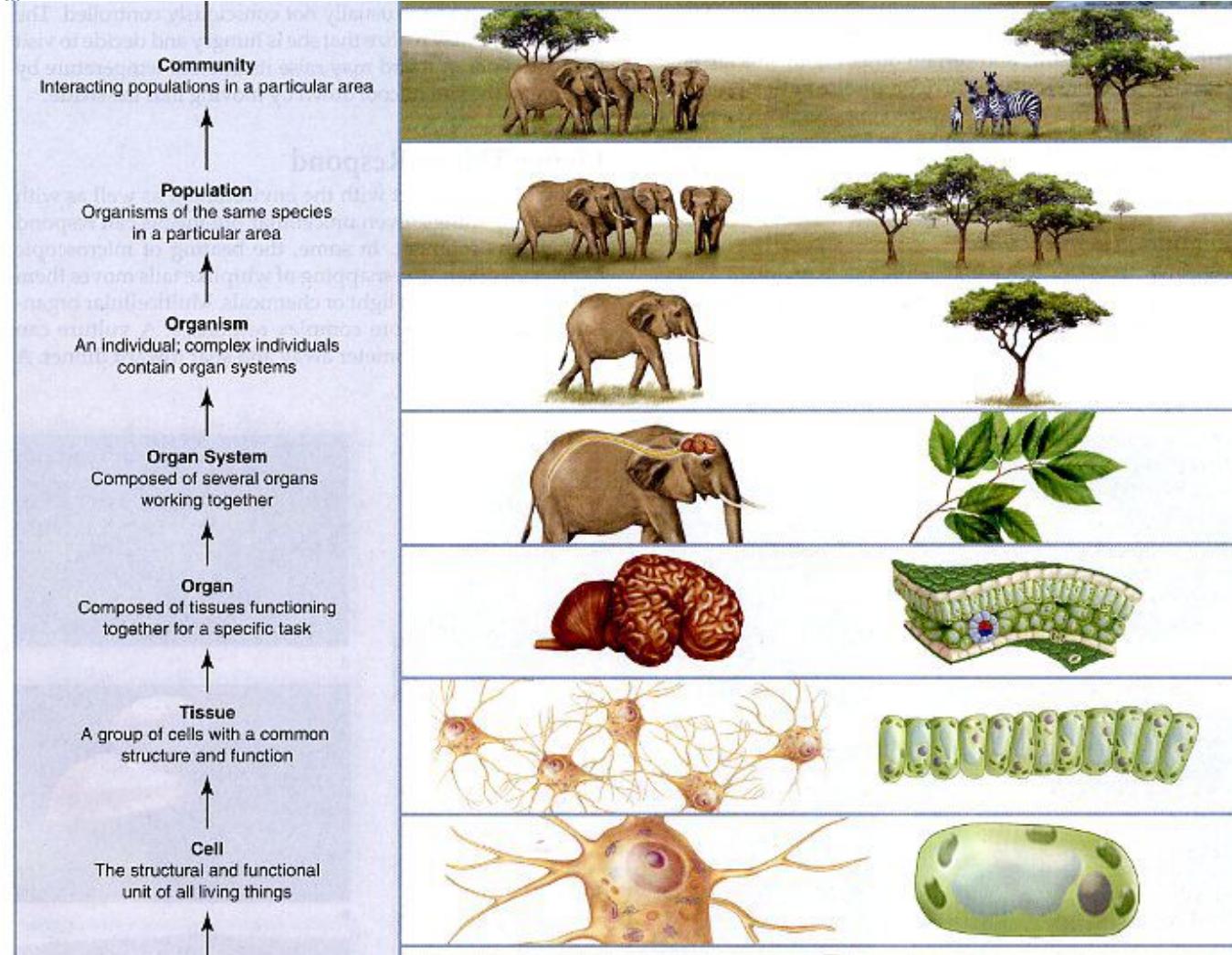
Applications

Niklaus Zemp

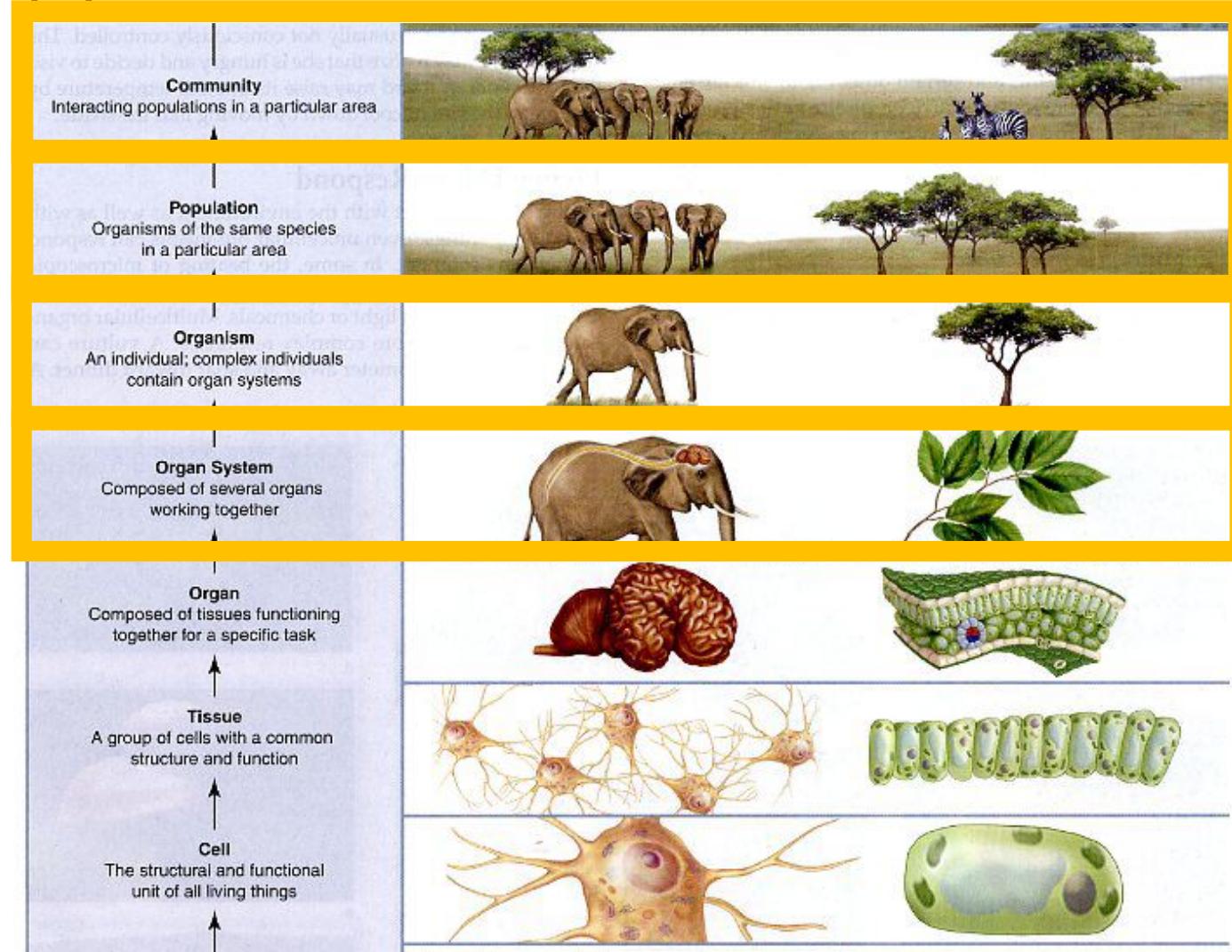
June 2020

Bioinformatics Genetic Diversity Centre (GDC) ETH Zurich

NGS applications

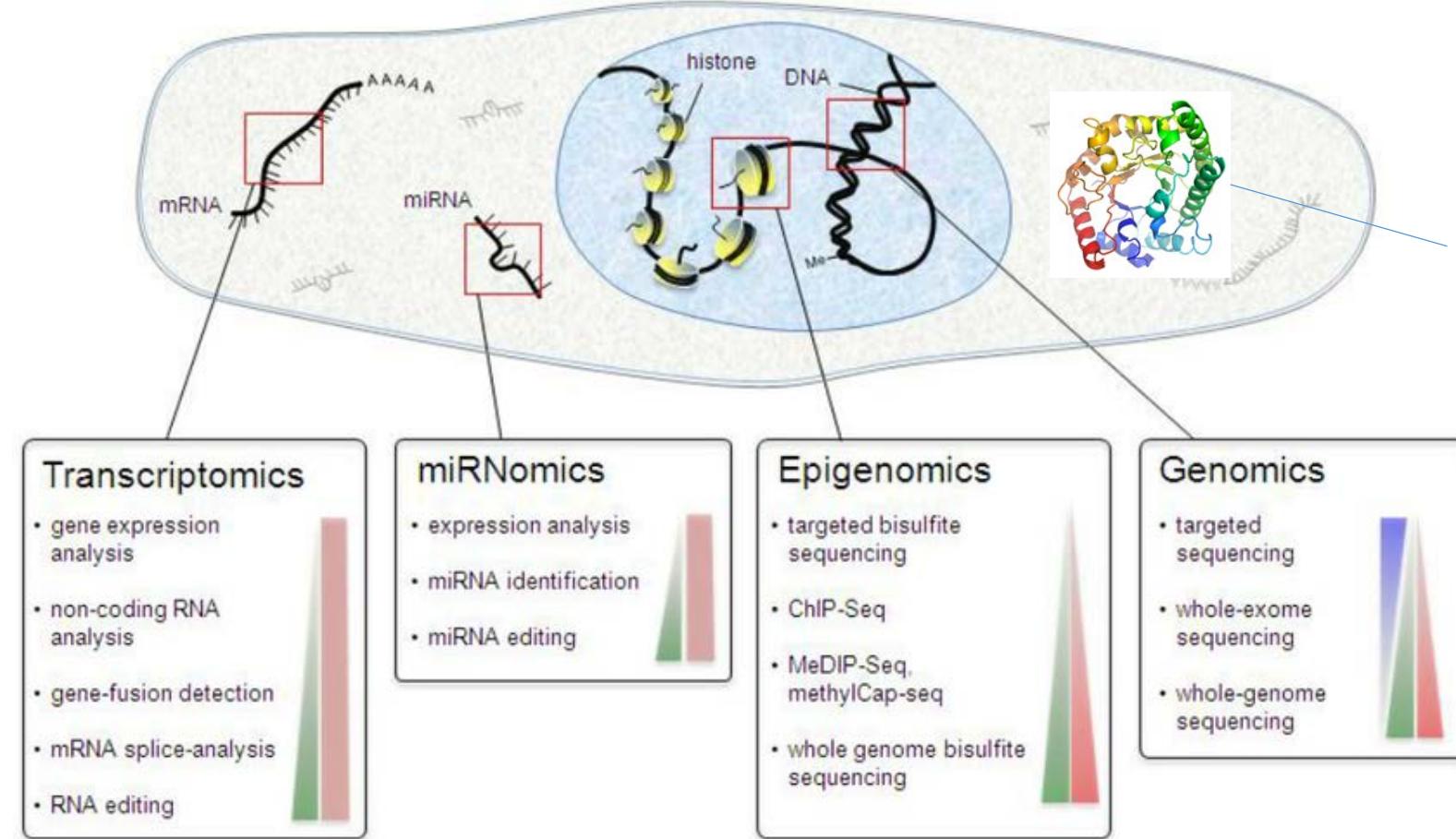


NGS applications

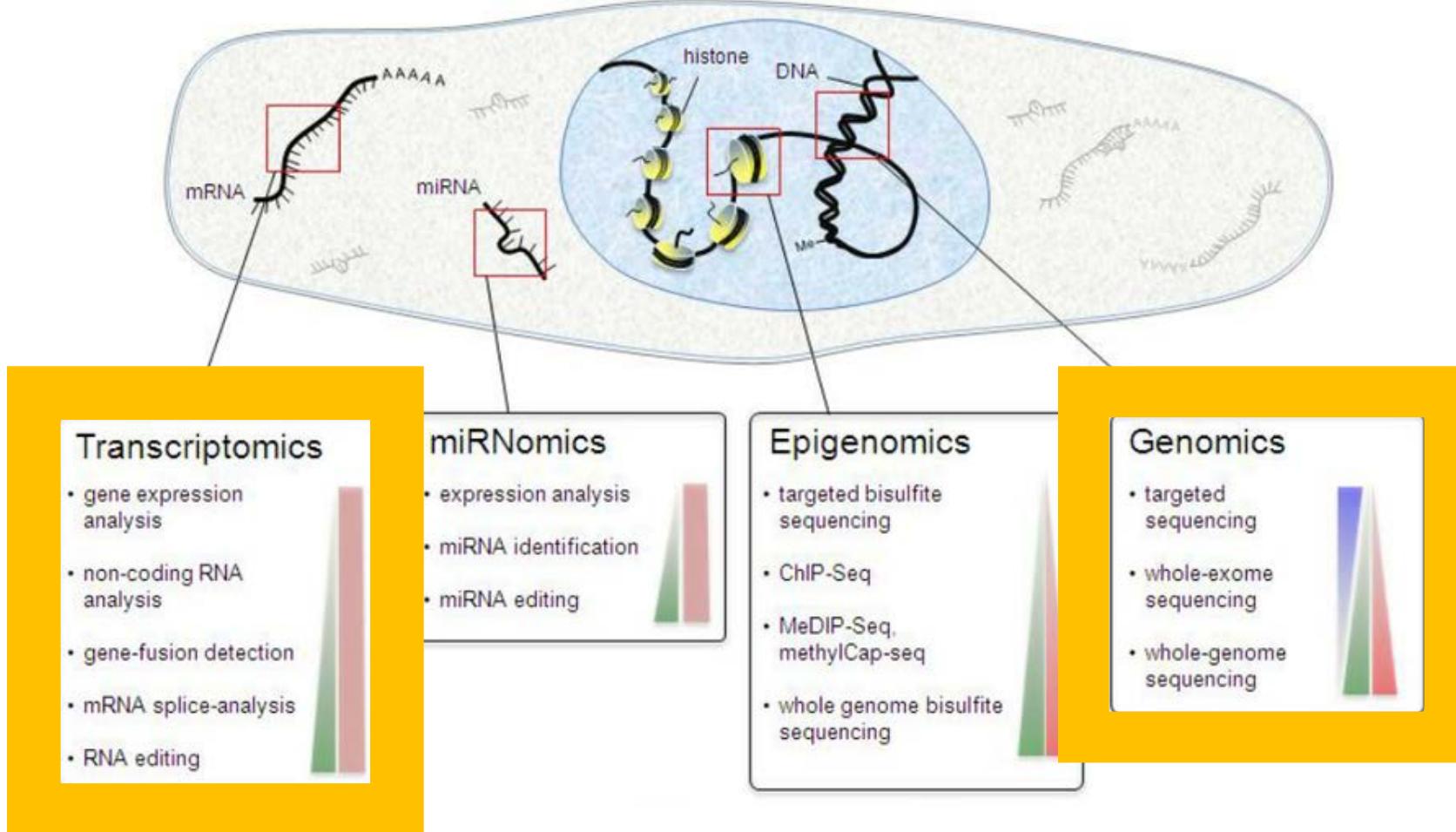


NGS applications

DNA → RNA → protein



NGS applications



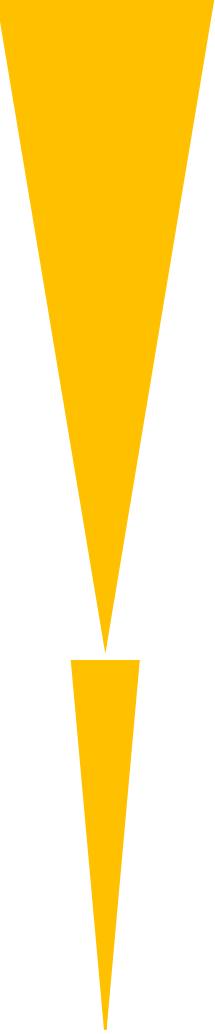
NGS applications/methods

- Genomics

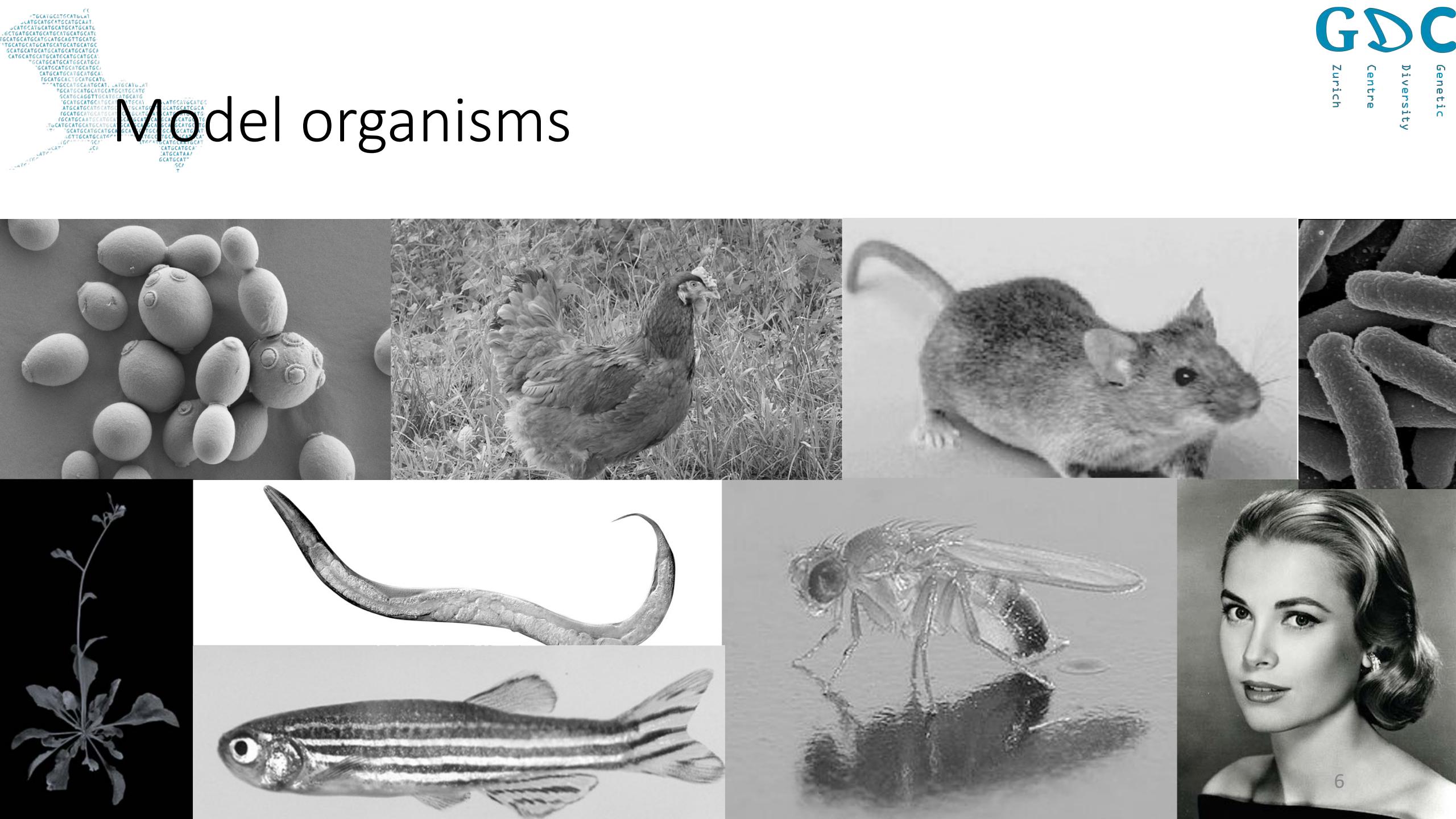
- Metagenomics
 - DNAseq (WGS)
 - Target enrichment sequencing
 - RNAseq
 - RADseq
 - Ampseq
- e.g. screenings, functional analysis
- e.g. signature of selection, GWAS
- e.g. population genomics, GWAS
- e.g. population stucture
- e.g. metabarcoding

- Transcriptomics

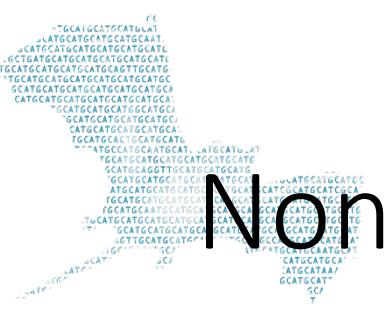
- Metatranscriptomics
 - RNAseq
 - Amplicon sequencing
- e.g. functional analysis
- e.g. genome expression anaylsis
- e.g. expression anaylsis



Model organisms



Non-model organisms



Take home massage

- Nearly universal applicable
 - Fastly developing
 - Revolutionised genomics of non-model organisms
 - Limitation in non-model organisms

