

# Available BioIT tools

- [PhysBinder](#)
- [ConTra](#)
- [SitePrediction](#)
- [TFdiff](#)

## Other links

- [RegCreative](#) - The RegCreative Jamboree
- [ORegAnno](#) - The Open REGulatory ANNOtation database
- [JASPAR](#) - The high-quality transcription factor binding profile database

## Genome Browsers

- [Ensembl](#)
- [UCSC](#)
- [NCBI Map Viewer](#)

## Sequence databases

- [NCBI Entrez](#) - Nucleotide, Protein, Gene, ...
- [UniProt](#) - Swiss-Prot/TrEMBL
- [TIGR](#) - The Institute for Genomic Research
- [BioMart](#) - Generate output, including sequences and tabulated list data from Ensembl

[HUGO Gene Nomenclature Committee \(HGNC\)](#) - search approved gene names and symbols

