

The future of immunogenicity screening

Introducing LENS^{ai} Integrated Intelligence Technology - an innovative platform that integrates advanced AI capabilities to provide unparalleled protein analysis and immunogenicity screening with lightning-fast throughput. LENS^{ai} Immunogenicity calculation combines HLA II binding and human proteome presence using BioStrand's proprietary HYFT[®] technology for early stage risk assessment.

High-throughput immunogenicity analysis

Built for high volume

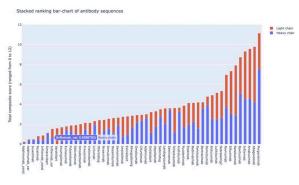
Virtually limitless quantity can be screened, compared and ranked

Flexible implementation

Ability to integrate in your own pipelines and workflows

Detailed reporting

Massive data package made transparent and instantly usable

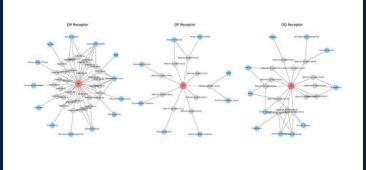


In-depth immunogenic profiling

Built for insight

- Detailed linkage between clone and target
- Geno- and phenotype binding distribution mapped to target indication profile

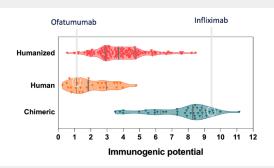
Connecting target, lead and clinical events MHCII allele phenotypes and genotypes associated with clinical events



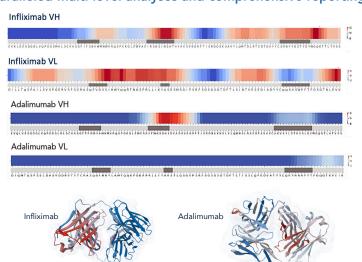
Introducing LENSai Immunogenicity Screening

Key features: unmatched throughput, speed, scalability, and accuracy

Reference your antibodies against a database of all therapeutic antibodies



Unparalleled multi-level analyses and comprehensive reporting



Immunogenicity composite scores of antibodies in the therapeutic structural Ab database (n \approx 2000):

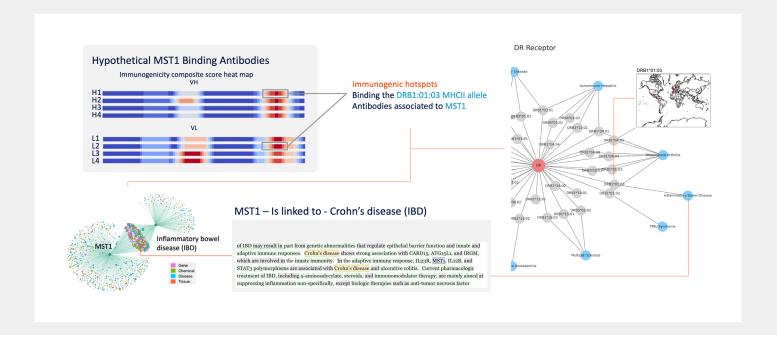
Parsed based on Nomenclature (n \approx 260) Statistical significance:

Significant difference between all means based on Kruksal-Wallis analysis [P-value: <0,0001]
Significant, Bonferonni corrected, difference between individual groups and Human (control group) based on Mann-Whitney [P-value: 0,0002 and better]

Immunogenic zones per sequence

Normalized score showing immunogenicity hotspots by combining HYFT Universal Fingerprint proteome screening and HLA II binding scores.

Dark Blue = 0 = low immunogenic potential
Dark Red = 100 = highest immunogenic potential
CDR annotations based on IMGT numbering





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