



Biologics Design and Protein Engineering

- Structure-Based Biologics Design
- Analyze and Edit Protein Sequences
- \circ Protein Properties and Developability
- ADC and Fusion Protein Modeling
- $\circ\,$ Protein, Antibody and Peptide Modeling
- Protein Engineering
- Simulations



The Molecular Operating Environment (MOE) has a comprenhensive suite of applications for computer-aided biologics design packaged in a workflow oriented interface for streamlined visualization and analysis. These applications aid in identifying and prioritizing potential mutagenesis experiments for affinity maturation, property modulation and liability screening.

chemcomp.com

Biologics Design and Protein Engineering

Structure-Based Biologics Design

- Easy-to-Use Interface for Biologics Modeling
- Protein-protein Interface Visualization
- Identify Protein-Protein Contacts
- Surfaces and Property Maps for Analyzing Structures
- Solvent Analysis for Protein Surfaces

Analyze and Edit Protein Sequences

- Visualize, Manage and Edit Protein Sequences
- Automatic Annotation of Protein Family Motifs
- Align Multiple Sequences and Structures
- Calculate Residue Identity and Similarity
- Graft Sequences, Loops and Secondary Structures

Protein Properties and Developability

- Patch Analysis to Identify Aggregation Prone Regions
- Calculate Protein Property Descriptors
- Identify Surface Accessible Residues
- Liability Detection of Post-Translational Modifications
- Use Protein Descriptors for QSAR and QSPR Modeling

ADC and Fusion Protein Modeling

- Linker Modeler to Connect and Fuse Protein Domains.
- Domain Motif Searching to Identify Linkers
- Optimize Protein Mutations with Protein Builder
- Non-Natural Amino Acids for ADC Linkers
- Identify ADC Conjugation Sites and Reactive Motifs

Protein, Antibody and Peptide Modeling

- Specialized Protein Families and Antibody Databases
- Generate Diverse Antibody Models from Fv to Ig
- Protein Multimer Homology Modeling
- In Situ Modeling of Loops and Sidechains
- Peptide Library Design and Screening

Protein Engineering

- Ala, Cys, SNP, Residue Scanning
- Sequence Optimization and Mutagenesis
- Evaluate Thermostability and Affinity
- Generate Focused Mutant Libraries
- Assess Solubility, Viscosity, Aggregation and Liabilities

Simulations

- Protein-Protein Docking
- Generate, Refine and Score Ensembles
- Explicit and Implicit Solvent Models
- Run MD to Model Surfactants and Solvents
- Simulate Excipient Effects Formulation



Rank	1:Leu3	1:1165	1:1nr14	1:1010	T:Pneza	1:19131	
1	Leu	Val	Leu	Arg	Phe	Trp	~
2	Val	Leu	Arg	Val	Leu	Leu	C
3	Cys	Phe	Phe	Leu	Tyr	Phe	
4		Ala	Trp	Trp	Trp	Met	
5		Ser	Ser	Thr	(iet	Tyr	25
6		lle	Val	Ser	Ale	Val	2
7					Sen	SAC	X
		Tyr			Val	(Lys)	6h
					Oys	THE Y	4
					Thr	SIX	0
						Ser	



NORTH AMERICA

Corporate Headquarters 1010 Sherbrooke Street W., Suite 910 Montreal, Quebec, Canada H3A 2R7 Tel.: +1 514 393 1055

EUROPE

United Kingdom St John's Innovation Centre Cowley Road, Cambridge CB4 OWS Tel.: +44 1223 422320

EUROPE

Germany Kaiser-Wilhelm-Ring 11 50672 Köln Tel.: +49 221 337790 0

ASIA Japan MOLSIS Chuo-ku, Tokyo 104-0032 Tel.: +81-3-3553-8030