

# JSDraw

Unified Sketcher  
For Small Molecules and Macromolecules

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Chief Scientific Office, Scilligence Corp.



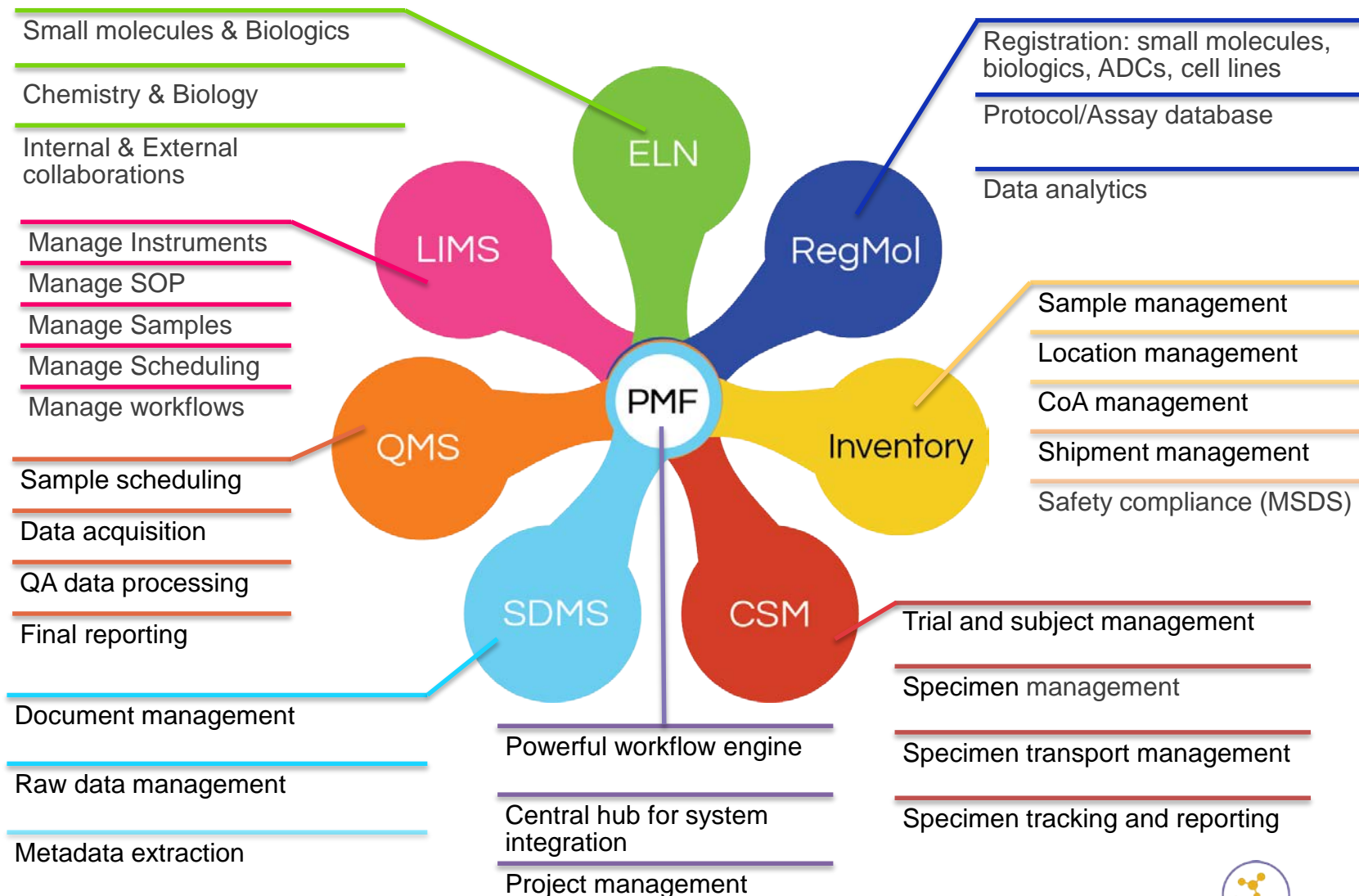
# Outline

- Overview of Scilligence Informatics Solutions
- The building blocks of Scilligence Informatics Solutions
- Why do we need a new structure sketcher/editor?
- JSDraw - A unified sketcher for both small molecules & macromolecules
- What's HELM and why do we need that?
- HELM Web Editor is based on JSDraw

# Company Overview

- **Boston Based and Founded in 2010**
- **ISO9001:2015 and ISO27001:2013 certified**
- **Informatics for Data Driven Discovery**
  - Web-based and no Java-dependency
  - A single platform bridging small molecules and biologics
  - Fully supports HELM Notation
  - Collaboration, workflow, and project management
  - Powerful components for data mining and analytics
- **Broad Customer Base**
  - Pharma, Biotech, and Chemical companies
  - Government Agencies
  - Universities, Hospitals, and Institutes

# Workflow Driven Integrated Platform



# Data Mining and Analytics

R-group  
Decomposition

SAR visualization

Gap analysis

Chrawler

Unstructured  
Data

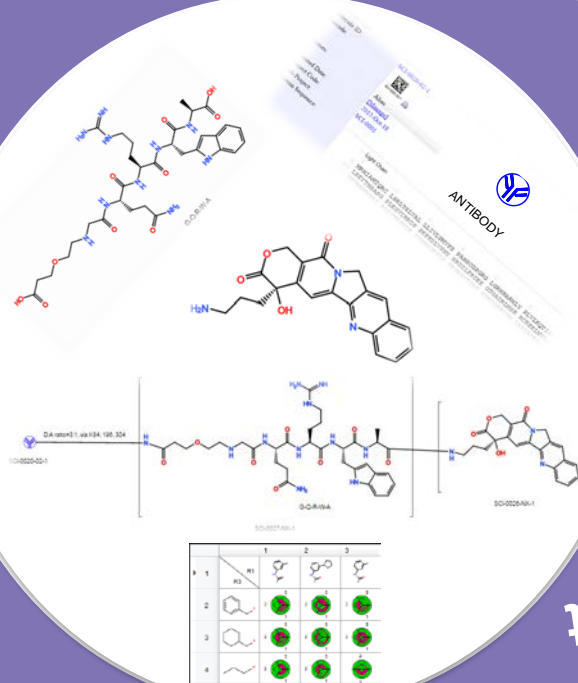
Network drive

Documentum

Database

SharePoint

SAR Analyzer



Chem4Sharepoint

Small  
Molecules  
&  
Biologics

Input

View

Search

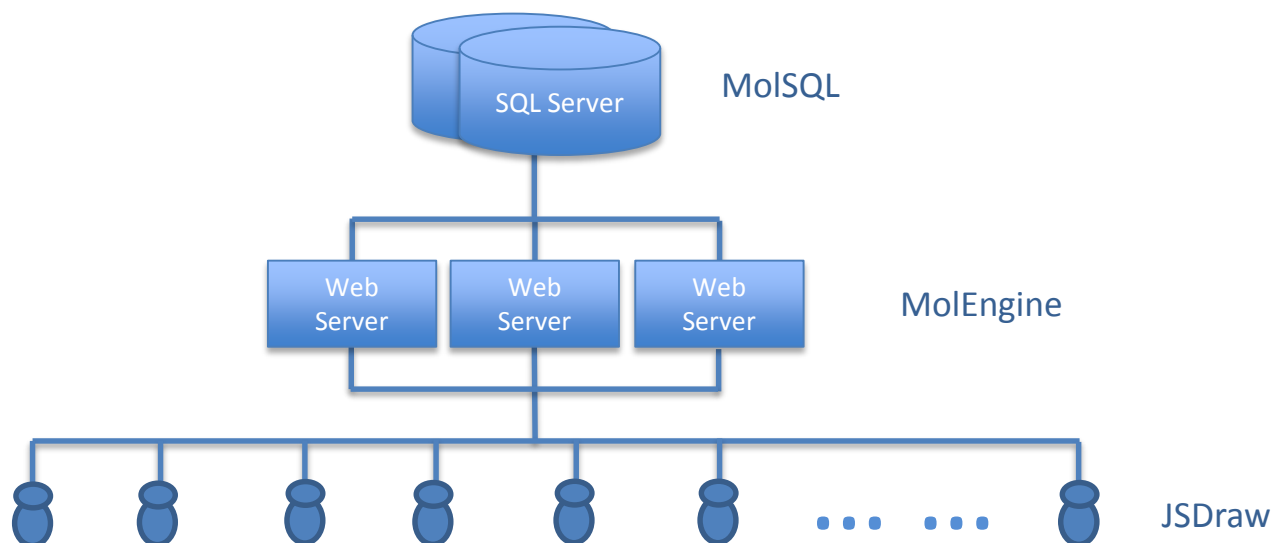
Cheminformatics inside  
Office

Informatics for Biologics &  
ADCs

Interface with databases

# Building Blocks for Scilligence Solutions

- **MoI SQL (Database Tier)**
  - Unified cartridge for small molecules and macromolecules
- **MolEngine (Middle Tier)**
  - Unified .NET toolkit for cheminformatics and bioinformatics
- **JSDraw (Client)**
  - Unified sketcher for small molecules and macromolecules

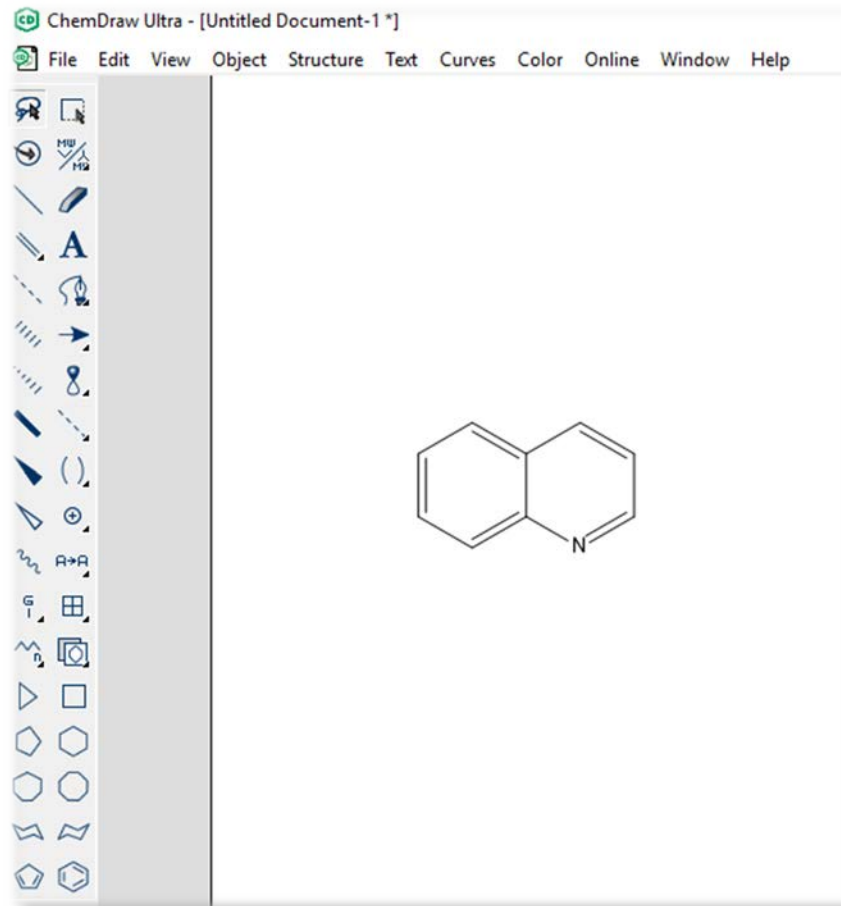


# Structure Editor Landscape before 2010

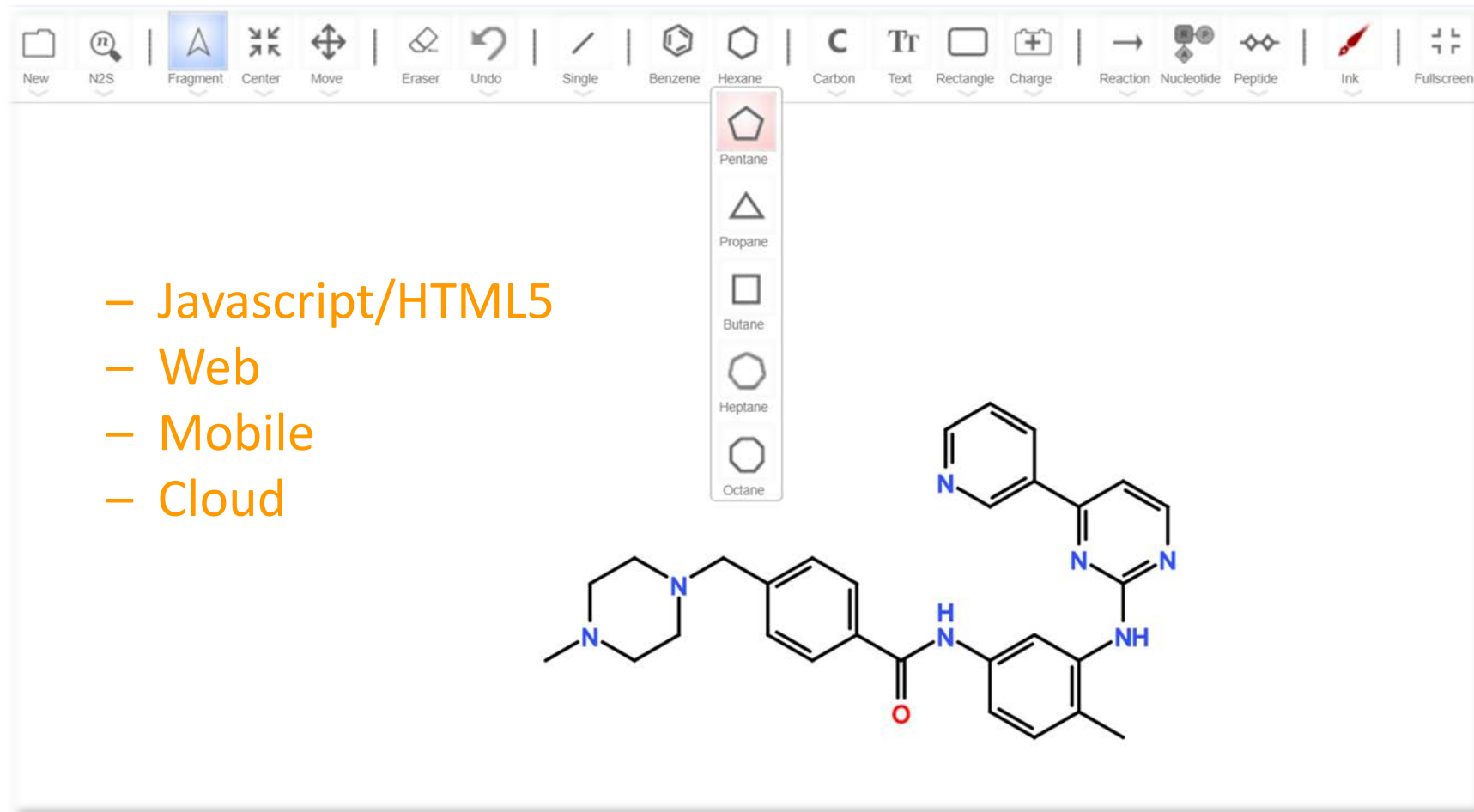
- **ISIS Draw**
- **ChemDraw**
- **MarvinSketch**
- ...

## Not optimal for

- Web-based applications
- Java dependency
- Mobile platforms



# JSDraw: Designed for Web and Mobile





# JSDraw: JavaScript Framework for Cheminformatics

- **JSDraw core Cheminformatics Javascript library**
  - Edit and display chemical structures
  - Edit and annotate biological sequences
  - Chemistry spreadsheet
- **JSDraw.Form**
  - Web Form Framework for Cheminformatics
- **JSDraw.Page**
  - Web Application Framework for Cheminformatics

# UI for Proteins and Genes

The screenshot displays the Scilligence software interface for editing protein and gene sequences. The main window is titled "Edit Sequence" and contains two sections: "Light Chain" and "Heavy Chain". Each section shows a sequence of amino acids, with some residues highlighted in green and yellow. The "Light Chain" sequence is as follows:

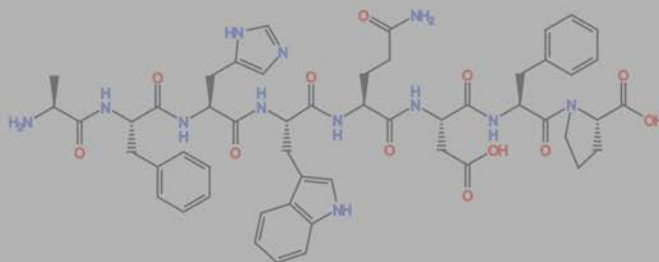
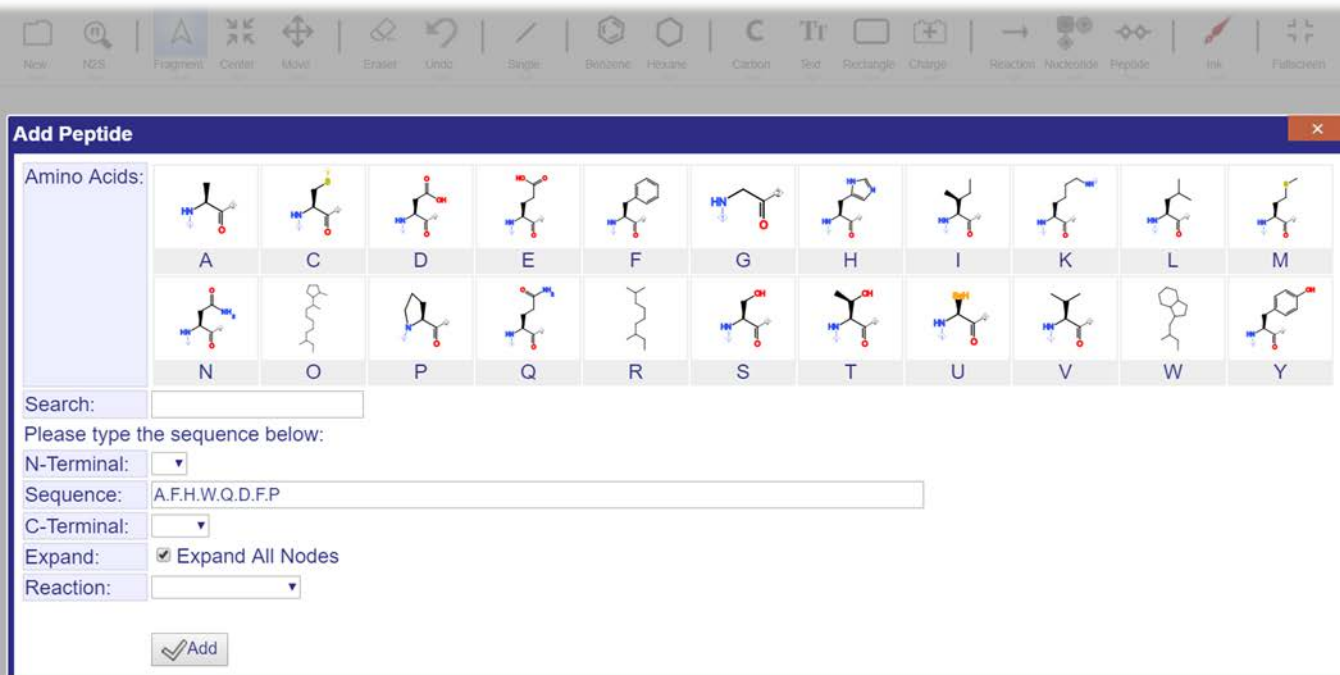
Position	1	2	3	4	5
0	PLVLKQTIPN	LS	EYTN	SASG	PLEGVIRRDS
50	GTGADRLMSK	RC	KEKLN	VLA	YSVMNEWPGI
100	YEGRAVTIAT	SDRDQSKYGM	LARLAVEAGF	DWVS	YVSRRH
150	ISSHVHGCFT	PE	TALL	ESG	VRKPLGELSI
200	FMDRNLEQMQ	NFVQLHTDGG	AVLTVTPAHL	VSVWQ	PESQK
250	EKNQVLVRDV	ETGELRP			

The "Heavy Chain" sequence is as follows:

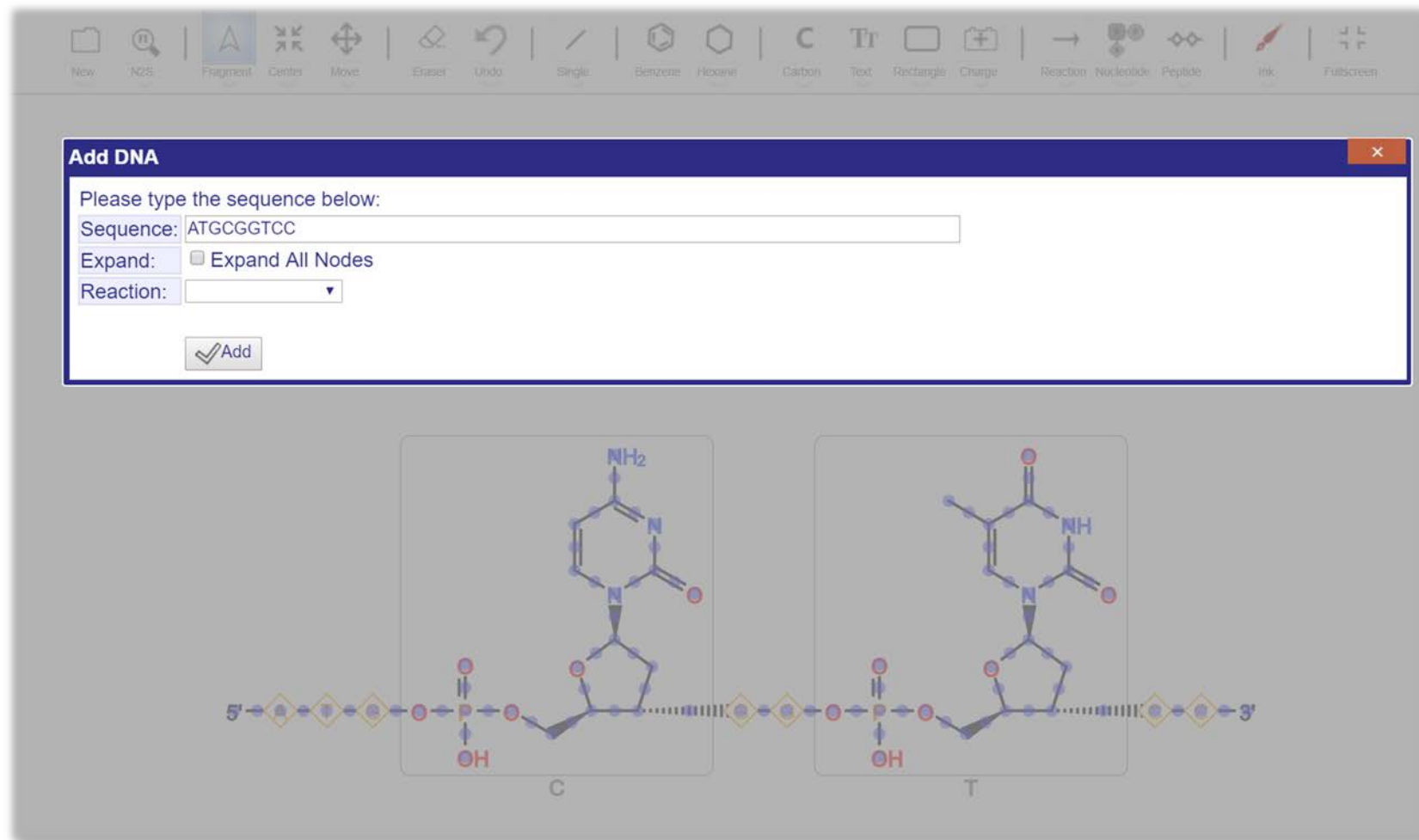
Position	1	2	3	4	5
0	TIPN	LS	EYTN	SASG	PLEGVI
50	LMSKRCKEKL	NVLAYSVMNE	WPGIRLLVTE	SWDEDYHHGQ	ESLHYEGRAV
100	TIATSDRDQS	KYGM	LARLAV	EAGFDWVS	YVSRRH
150	GCFTEPSTAL	LESGVRKPLG	ELSIGDRVLS	MTANGQAVYS	EVILFMDRNL
200	EQMQNFVQLH	TDGGAVLTVT	PAHLVSVWQP	ESQKLTFVFA	DRIEKNQVL
250	VRDVETGELR	PQRVVKVGSV	RSKGVVAPLT	REGTIVVNSV	AASCYAVINS

The interface also includes a top toolbar with various drawing and editing tools, and a left sidebar with icons for "GENE" and "ANTIBODY IgG". A "Save" button is located at the bottom right of the "Edit Sequence" window.

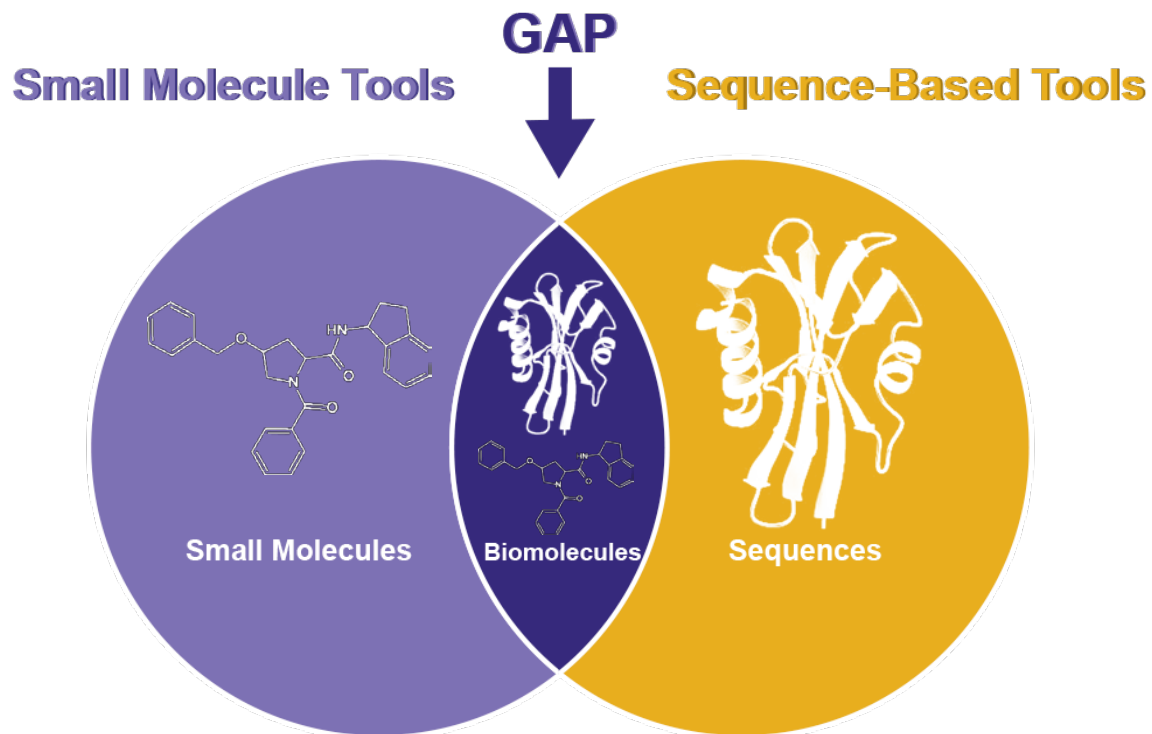
# UI for Peptides



# UI for Oligonucleotides/Oligonucleosides



# Why HELM?



**HELM:** The representation language and toolkit that remediate this gap

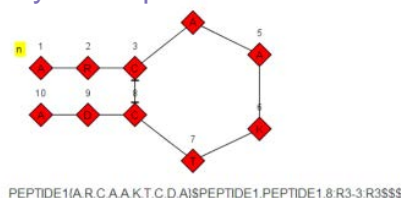
# What is HELM?

- **HELM stands for Hierarchical Editing Language for Macromolecules**
- **Invented by Pfizer and open-sourced by Pistoia Alliance**
- **HELM creates a standard for representing, exchanging, and manipulating macromolecules**
- **Improves data storage and knowledge sharing in R&D**

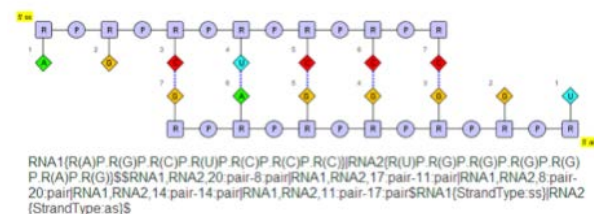
## Linear oligonucleotides



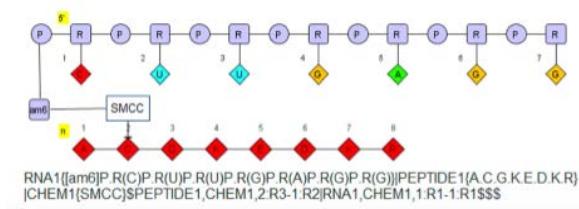
## Cyclic Peptide



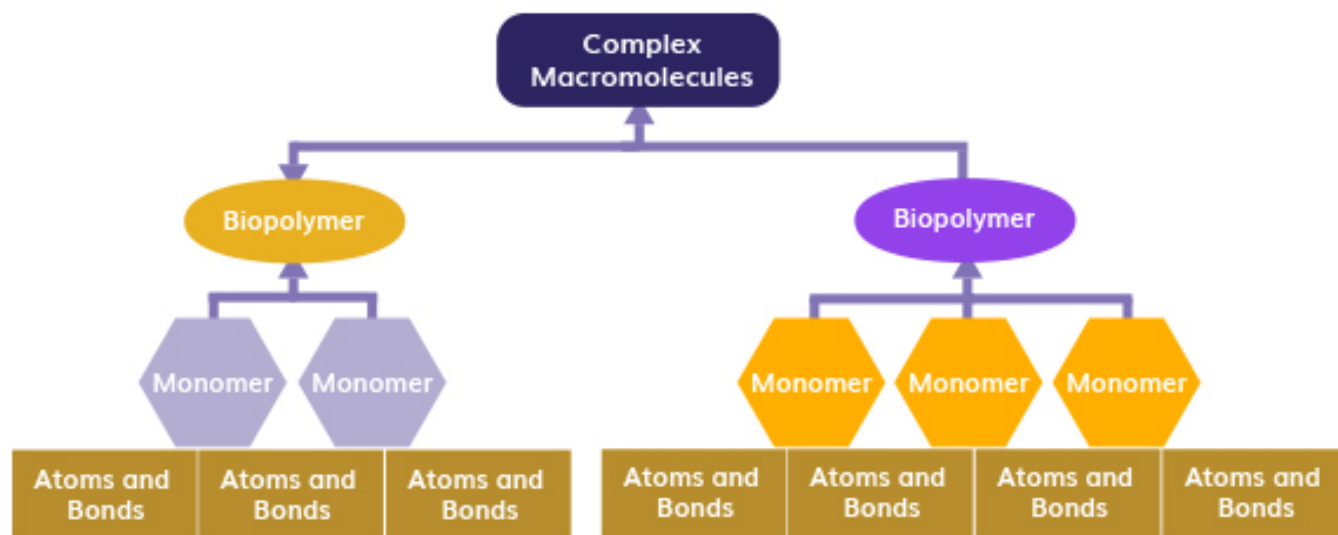
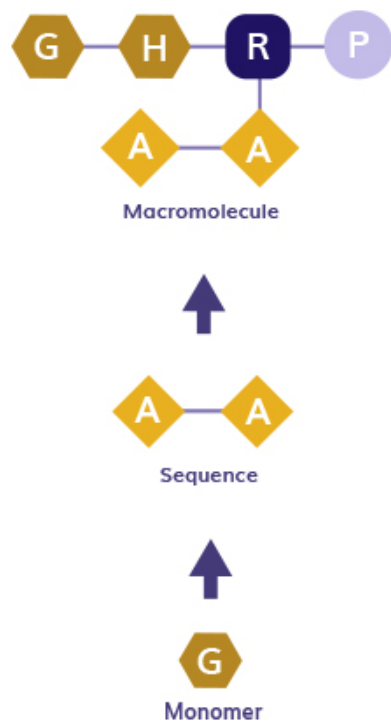
siRNA



## Oligonucleotide peptide conjugate



# The Power of HELM's Hierarchy



# HELM Web Editor is Based On JSDraw

- **HELM**

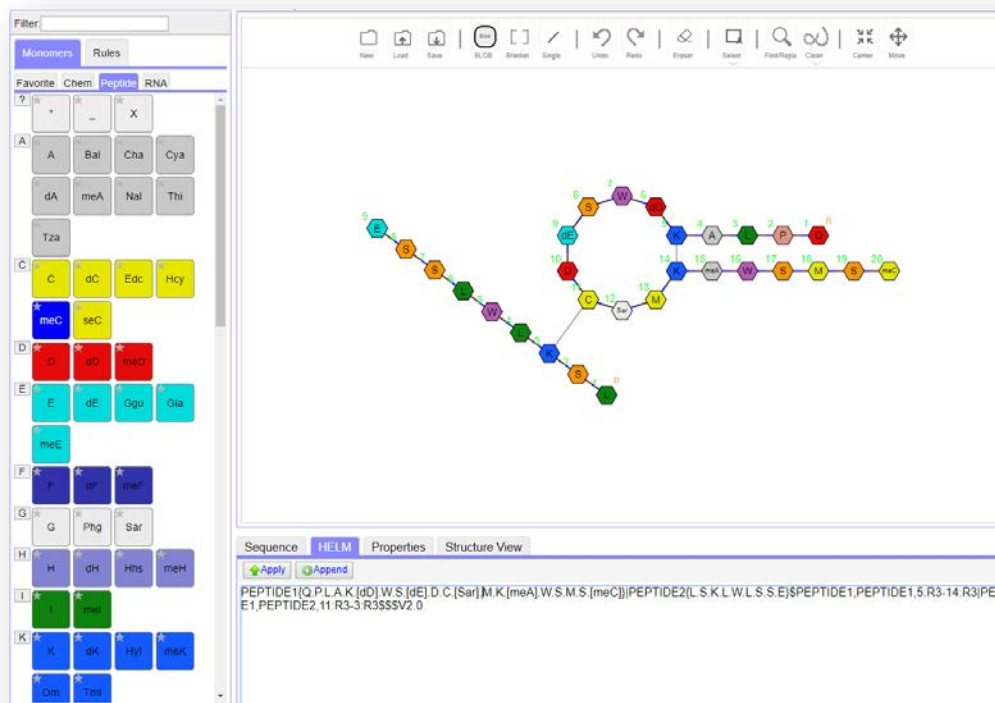
- Invented at Pfizer
- Open sourced by Pfizer
- Adopted by Pistoia

- **HELM Java Client**

- Built on Java
- Annotation is not supported
- Ambiguity is not supported

- **HELM Web Editor**

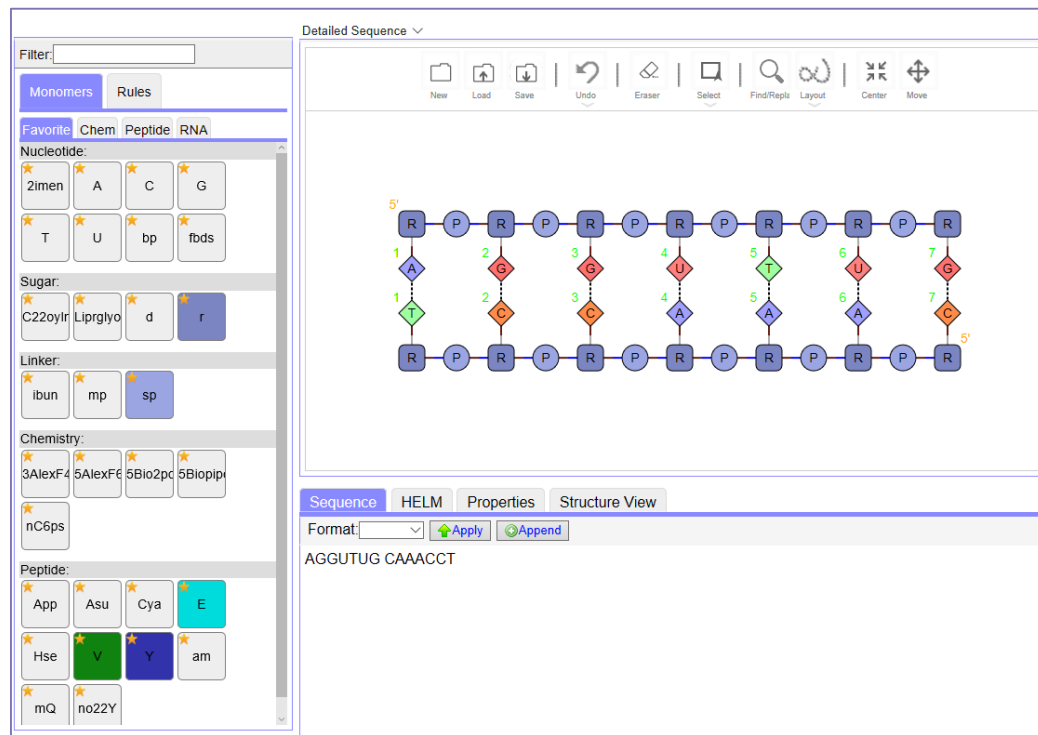
- Switch to JavaScript
- Support annotation
- Support Ambiguity





# HELM Web Editor Project

- **Scilligence was selected as the developer in 2016**
- **Three Phases**
  - Core HELM functionalities in HELM Web Editor 1.0
    - Released in February 2017
  - Ambiguity and Annotation Features
    - In Progress
  - Advance Annotation for Antibody
    - Planned



# HELM Web Editor 2.0 Modules

- HELM Editor Control
- HELM App
- HELM Monomer Manager
- HELM Rule Manager

Rule Set					
<div> <div> <div></div> <div></div> <div></div> </div> <div> <div>Category:</div> <div></div> </div> <div> <div>Count:</div> <div></div> </div> </div>					
ID	Category	Name	Description	Author	Created Date
1	General	Claire test 1	dzfd	Claire Bellamy	
2		Claire test 2	dsaa	Claire Bellamy	
5	General	Claire test 4		Claire Bellamy	2017-May-03
3	General	Micky test 1	nfhdgh	Micky Mouse	
6	General	P to sP	Replace Ps by sPs		2017-May-08
4	General	Test 3	fsaef	Claire Bellamy	

Rule	
ID:	1
Category:	General
Name:	Claire test 1
Description:	dzfd
Author:	Claire Bellamy
Javascript:	<pre>function(plugin) { }  //function(plugin) { //  scil.Utils.ajax('http://SERVER/youerservice', function(ret) { //    plugin.setHelm(ret.new_helm); //  }); //}</pre>

Monomer List

SymbolName:

Polymer Type:

Monomer Type:

Count:

Previous Page

1

...

9

10

11

12

13

14

15

16

17

18

19

Next Page

Symbol	Name	Natural Analog	Polymer Type	Monomer Type	R1	R2	R3	Author	Created Date
pnG	PNA Guanine	X	PEPTIDE	Backbone	H	OH			2017-Sep-29
pnT	PNA Thymine	X	PEPTIDE	Backbone	H	OH			2017-Sep-29
prpC	5-Propynyl-Cytosine	C	RNA	Branch	H				2017-Sep-29
prpU	5-propynyl Uracil	U	RNA	Branch	H				2017-Sep-29
qR	2-O-beta-hydroxy-ethoxy-methyl Ribose (Qiagen)	R	RNA	Backbone	H	H	OH		2017-Sep-29
sDBL	Symmetric Doubler	null	CHEM	Undefined	H	H	H		2017-Sep-29
sP	Phosphorothioate	P	RNA	Backbone	OH	OH			2017-Sep-29
seC	SelenoCysteine	C	PEPTIDE	Backbone	H	OH			2017-Sep-29
IR	Ribose	R	RNA	Backbone	H	H	OH		2017-Sep-29

Monomer

Symbol:

sDBL

Name:

Symmetric Doubler

Polymer Type:

CHEM

Monomer Type:

Undefined

Natural Analog:

null

Author:

SMILES:

[H]1JC(CNC(=O)CCCCO[H]2)JCNC(=O)CCCCO[H]3

Structure:

R1:

H

R2:

H

R3:

H

R4:

R5:

# Scilligence Solutions Fully Support HELM

- **Drawing HELM molecules in**
  - Scilligence ELN
  - RegMol (Registration System)
  - Scilligence Inventory
- **HELM Molecule Registration**
  - Molecule rules
  - Uniqueness check
  - Central monomer management
- **MoI SQL**
  - Chemistry Cartridge supporting HELM
- **TouchMol4Office**
  - The only Office add-on supporting HELM

# JSDraw – A Unified Sketcher/Editor

- **Web-based, Java-free**
- **Mobile compatible**
- **Cloud deployment**
- **Supports both small molecules and macromolecule**
- **Supports HELM**

Thank You!

[www.scilligence.com](http://www.scilligence.com)