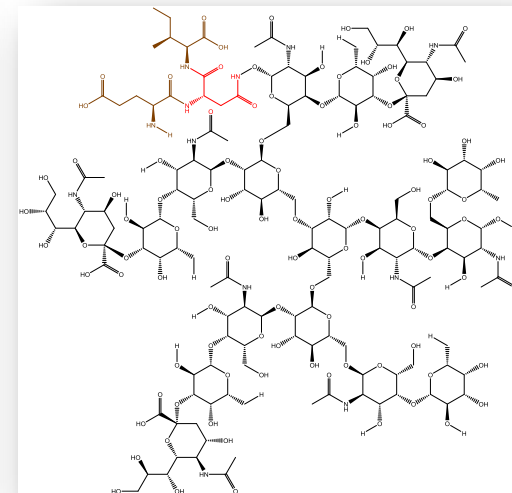
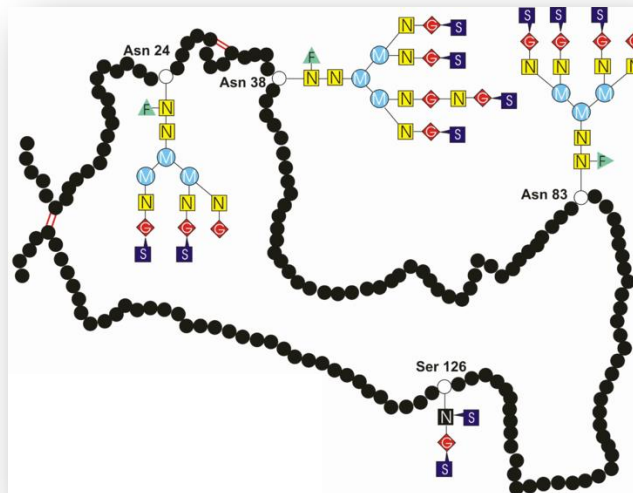
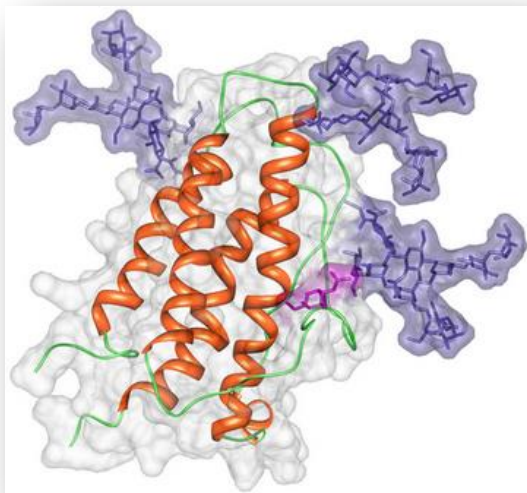


# Erythropoietin (EPO)



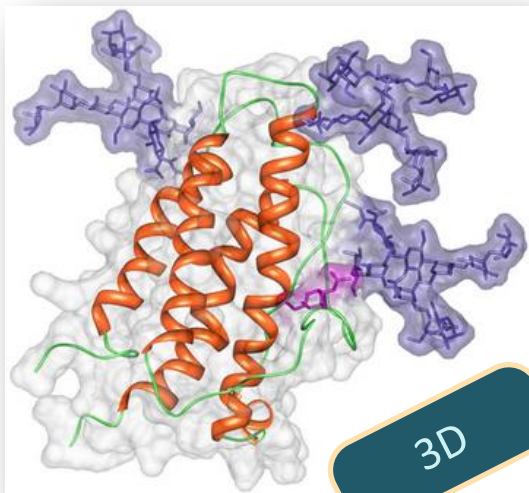
MGVHECPAWL WLLLSLLSLP LGLPVLGAPP RLICDSRVLE RYLLEAKEAE **N**ITGCAEHC  
 SLN**E**ITVDP TKVNFYAWKR MEVGQQAVEV WQGLALLSEA VLRGQALLV**N** SSQPWEPLQL  
 HVDKAVSGLR SLTLLRALG AQKEAISPPD AASAAPLRTI TADTFRKLFV VYSNFLRGKL  
 KLYTGEACRT GDR

# Connecting dots...

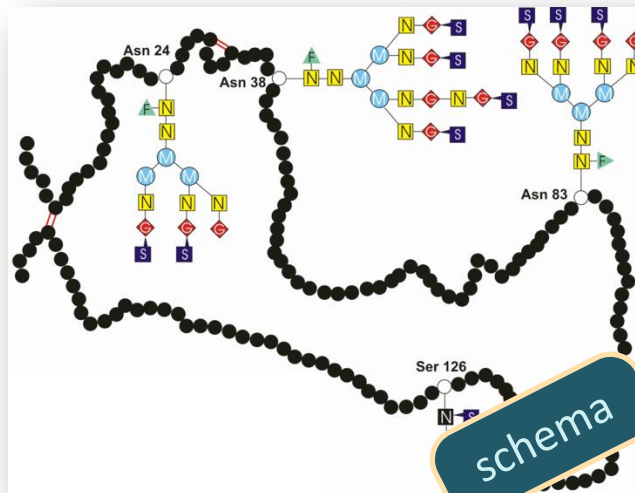
Bio-molecule registration using HELM and ChemAxon tools

Roland Knispel

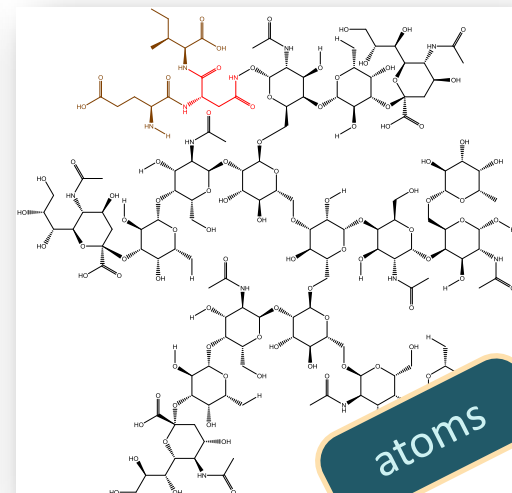
# Erythropoietin (EPO)



3D



schema



atoms

MGVHECPAWL WLLLSLLSLP LGLPVLGAPP RLICDSRVLE RYLLEAKEAE NITGCAEHC  
 SLNENITVPD TKVNFYAWKR MEVGQQAVEV WQGLALLSEA VLRGQALLVN SSQPWEPLQL  
 HVDKAVSGLR SLTLLRALG AQKEAISPPD AASAAPLRTI TADTFRKLFV VYSNFLRGKI  
 KLYTGEACRT GDR

sequence

3D

sequence

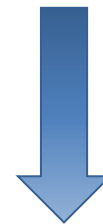
atoms

schema

-ligands  
-coordinates

-sequence  
+unnatural residues  
+ modifications

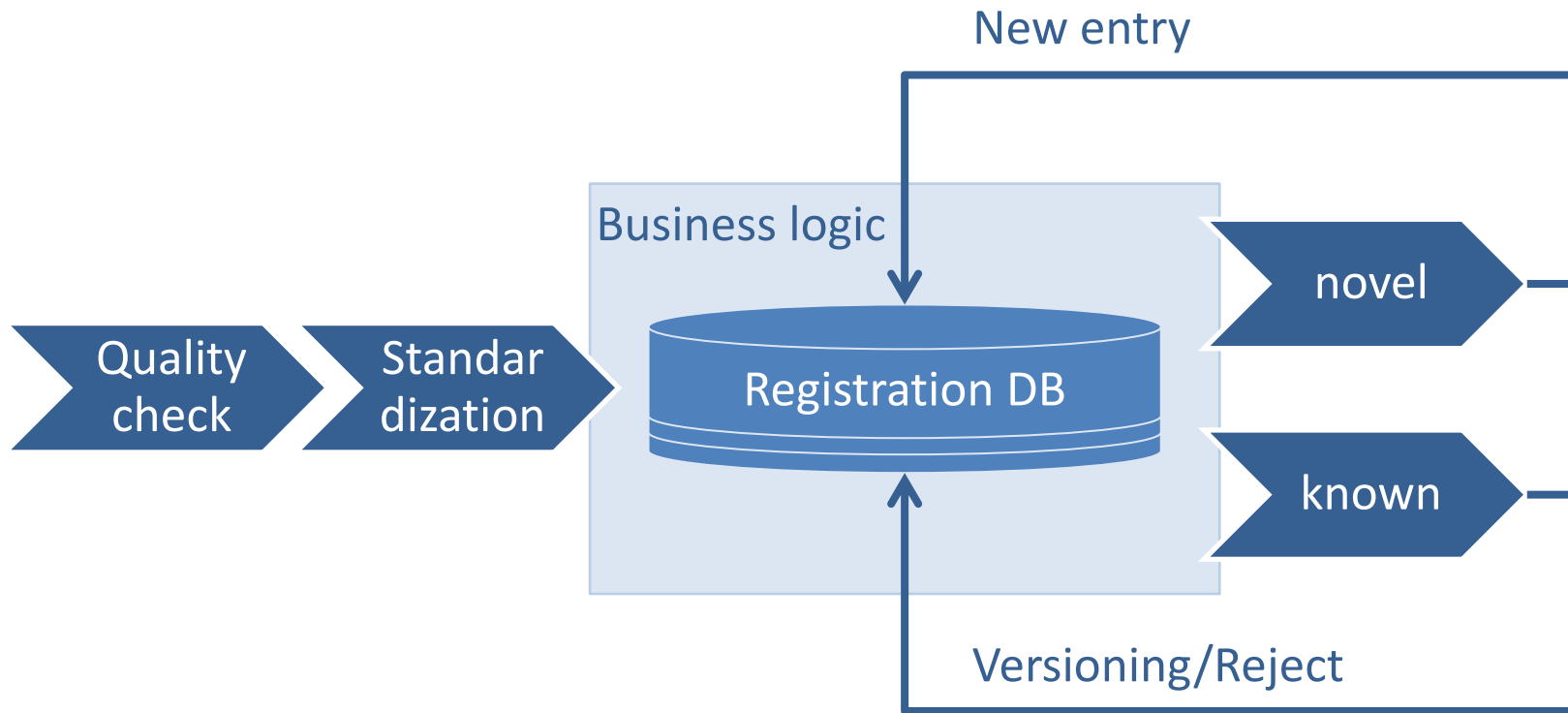
Custom non-std  
representation



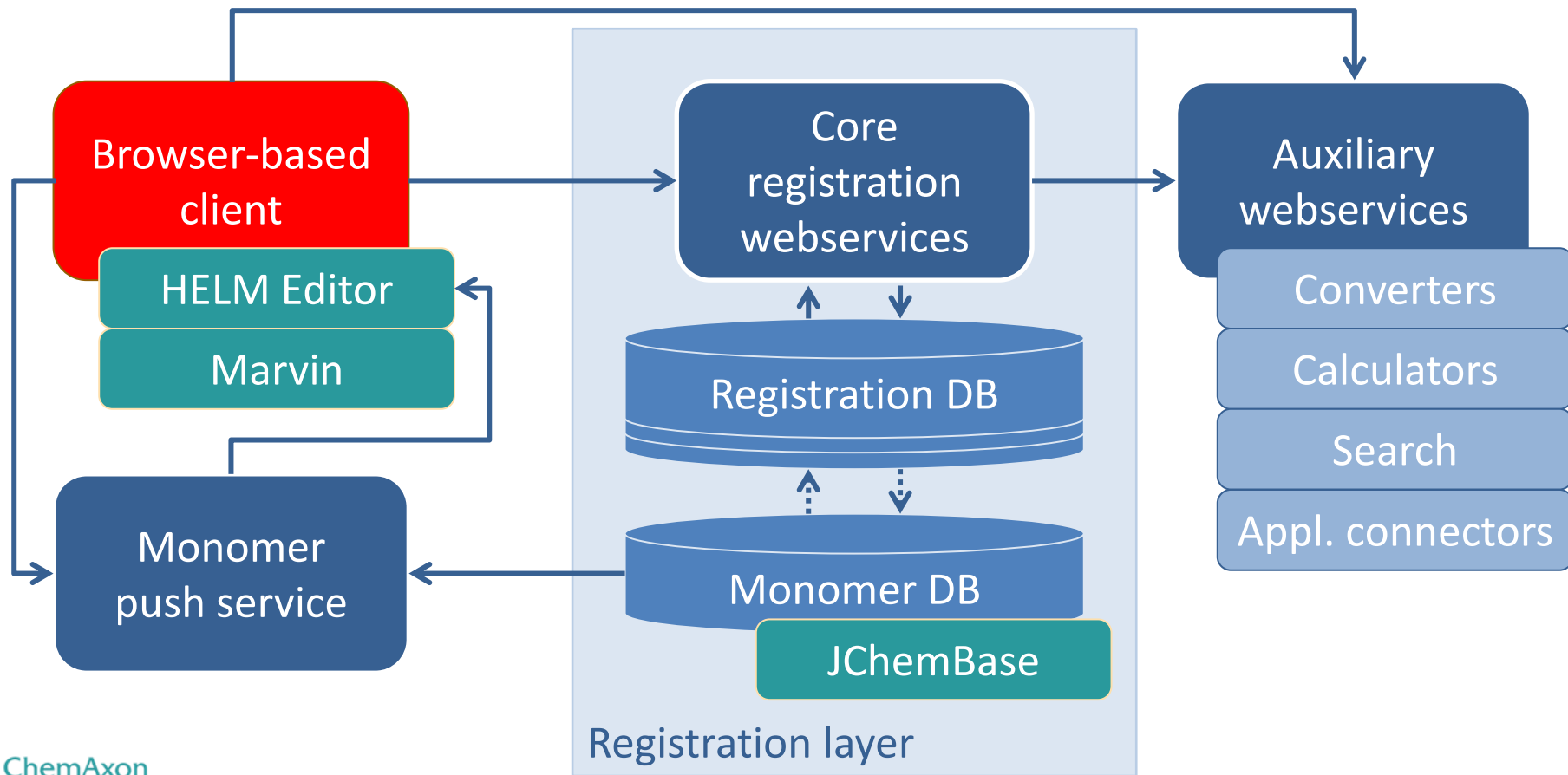
Search

Consumers  
DataMarts,  
Data Analysis  
applications

# Is it unique?



# Registration toolkit architecture



# List of available webservices

## HELM monomer registration

- Monomer Registration
- Monomer Updating
- Monomer Deletion
- GetByName
- GetByType
- GetAll

## Macromolecule registration

- Macromolecule Registration
- Macromolecule Updating
- Macromolecule Deletion
- GetByCid
- GetAll

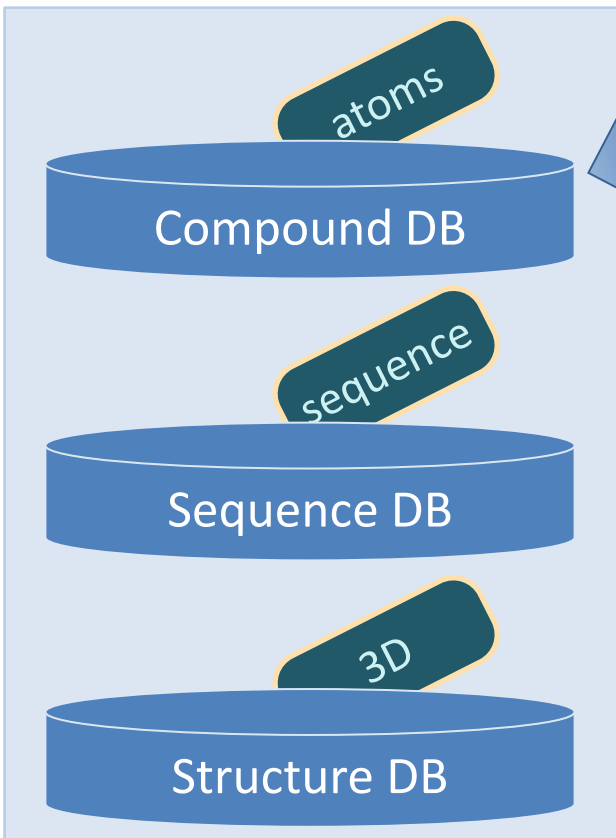
## Conversion

- Sequence/FASTA -> HELM
- HELM -> Natural Analogue sequence
- HELM -> canonical HELM
- HELM <-> SMILES
- HELM <-> Mol

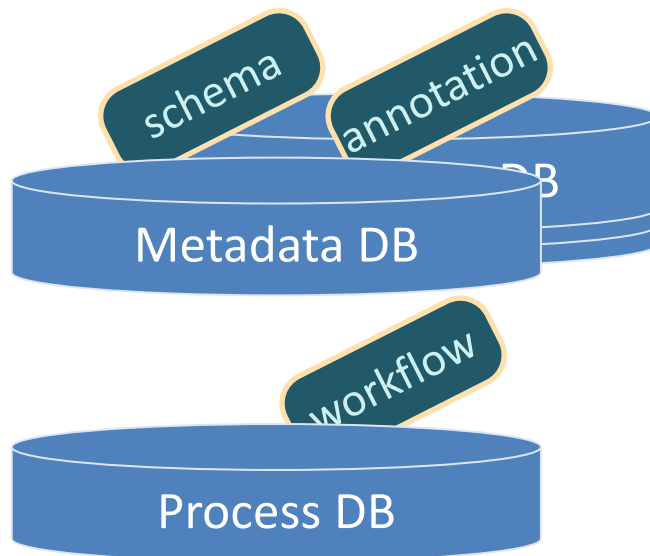
## Search

- General search
- QueryBuilder with filter types:
  - Full sequence
  - Subsequence
  - Any unnatural residue
  - Any chemical modification
  - ...

# Why not re-use existing systems?

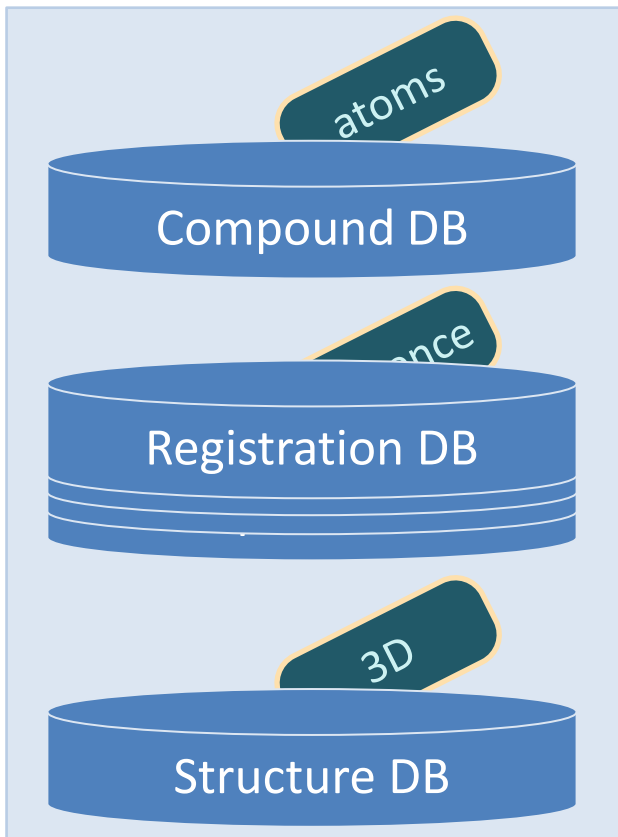


MGVHECPAWL WLLLSLLSLP LGLPVLGAPP RLIF  
 RYLLEAKEAE NITGCAEHC SLNENITVPD TK  
 MEVGQQAQEV WQGLALLSEA VLRGQALLVN SS  
 HVDKAVSGLR SLTLLRRLG AQKEAISPPD AA  
 TADTFRKLFR VYSNFLRGKL KLYTGEACRT GD





# ChemAxon's Biomolecule toolkit

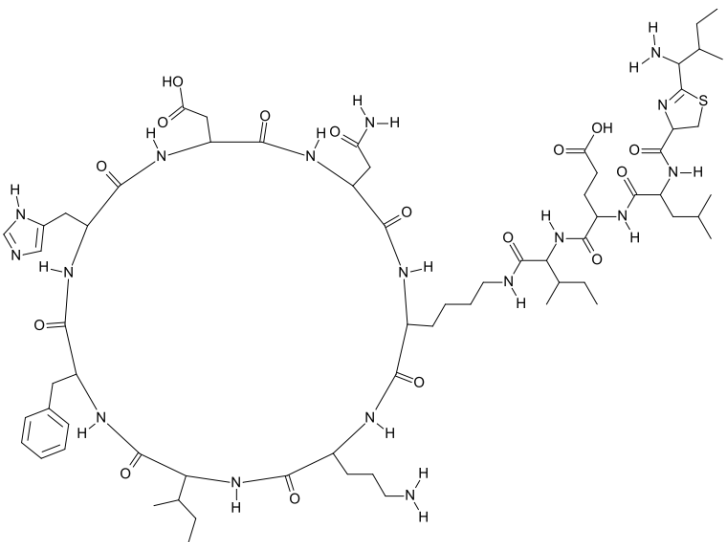


Canonical representation of small and large molecules

Registration of a chemically defined full IgG antibody in the „wink of an eye“ (100-400ms)



# Use case (Macrocyclic peptides)



## Requirements:

- Registration of **sequence and chemical structure**
- **Unambiguous representation** of peptide sequence
- Ability to quickly **enumerate library** with residue replacements chosen from a large set of peptide building blocks

POC implemented with ChemAxon's Bio-molecule registration toolkit

# Benefits



R&D  
 Synthesis  
 Bioinformatics  
 Biochemical Development  
 Open Access  
 Information Services

With this toolkit I can now:

- **Share** my molecules **unambiguously** internally and with external collaborators
- **Save time** on curating inconsistent representations received from collaborators
- **Save resources** on maintaining parallel systems for specimen registration
- **Remove bottlenecks** caused by legacy system limitations

## Edit biomolecule

## Query builder

Search

Clear query

Sequence 1

Match type: Subsequence

Sequence type: PEPTIDE

Sequence 1: MTG

Modification 1

Affects: Sequence 1

Modification type: Chemical

## Search result

Quickfilter

CID	Name	Helm
CXN43	random peptide (Cys-Cys bridge)	PEPTIDE1{M.T.G.C.R.L.C.Y.W.E.C}\$PEPTIDE1,PEPTIDE1,7:R3-4:R3\$\$\$
CXN44	random peptide (Cys-Cys bridge)	PEPTIDE1{M.T.G.C.R.L.C.Y.W.E.C}\$PEPTIDE1,PEPTIDE1,11:R3-4:R3\$\$\$
CXN45	random peptide (Cys-Cys bridge)	PEPTIDE1{M.T.G.C.R.L.C.Y.W.E.C}\$PEPTIDE1,PEPTIDE1,11:R3-7:R3\$\$\$
CXN46	random peptide (Acetylation)	CHEM1{Ac} PEPTIDE1{M.T.G.C.R.L.C.Y.W.E.C}\$CHEM1,PEPTIDE1,1:R1-1:R1\$\$\$

# Integration opportunities



SOAP and REST-ful WS toolkit



Compound Registration



JChem Base



Plexus



Instant JChem



JChem for Office

+ 3rd party applications

# Summary

- Template-based bio-molecule registration toolkit
- Fully compatible with the Open HELM standard for representing large molecules
- WS API for integration
- Shippable preview version



# How can we help you?



Explore our demo station