

Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

New

Open... Ctrl+O

Open File (Ctrl+O)

Insert

Display the Open dialog and select one or more files to open.

Close

Click here for help

See also: [Import preference](#), [Default File Types preference](#)

Save

Save

Page Setup...

Print Preview

Print... Ctrl+P

Change Server...

Refresh F5

Properties...

Recent Files

Exit Alt+F4

Predict Protein Formulation Properties

Calculate Protein Formulation Properties...

Calculate Aggregation Scores...

View Aggregation Sites

Define Protein <Undefined>

View at Cutoff Radius: Unset

Step through aggregation sites.

Show Details of Site

Show Surface

Reset Display

Analyze Transmembrane Domains

Done

Receptor-Ligand Interactions Pharmacophores Small Molecules X-ray My Tools

Search product features

Display Style

Discovery Studio 2020

What's New

Actions

My Recent Actions

- Calculate Protein Formulation ...
- Build Mutants...
- Open URL...
- Model Antibody Loops...
- Model Antibody Framework...
- Verify Protein (MODELER)...
- Build Homology Models...
- Analyze Ligand Poses...

My Recent Tool Panels

- Predict Protein Formulation Pr...
- Design Protein
- Build and Edit Protein
- Protein Report and Analysis
- Model Antibody

Data

My Recent Files

- 6Y84.mol2
- 6M03_active.dsv
- 6Y84_active.dsv
- DDT_ChEBI_161301.sd
- DDT_ChEBI_16130.sd
- DDT_ChEBI_16130.sdf
- Rloin_ChEBI_73222_aloevera.sdf
- Molecule.sd
- 6LU7_prep.dsv
- B-mycene.sd
- 6JIM_Many.dsv
- 6Y84_many1.dsv
- ursolic acid_ChEBI_9908_basil...
- 6VVO.dsv
- 6VVO_many1.dsv

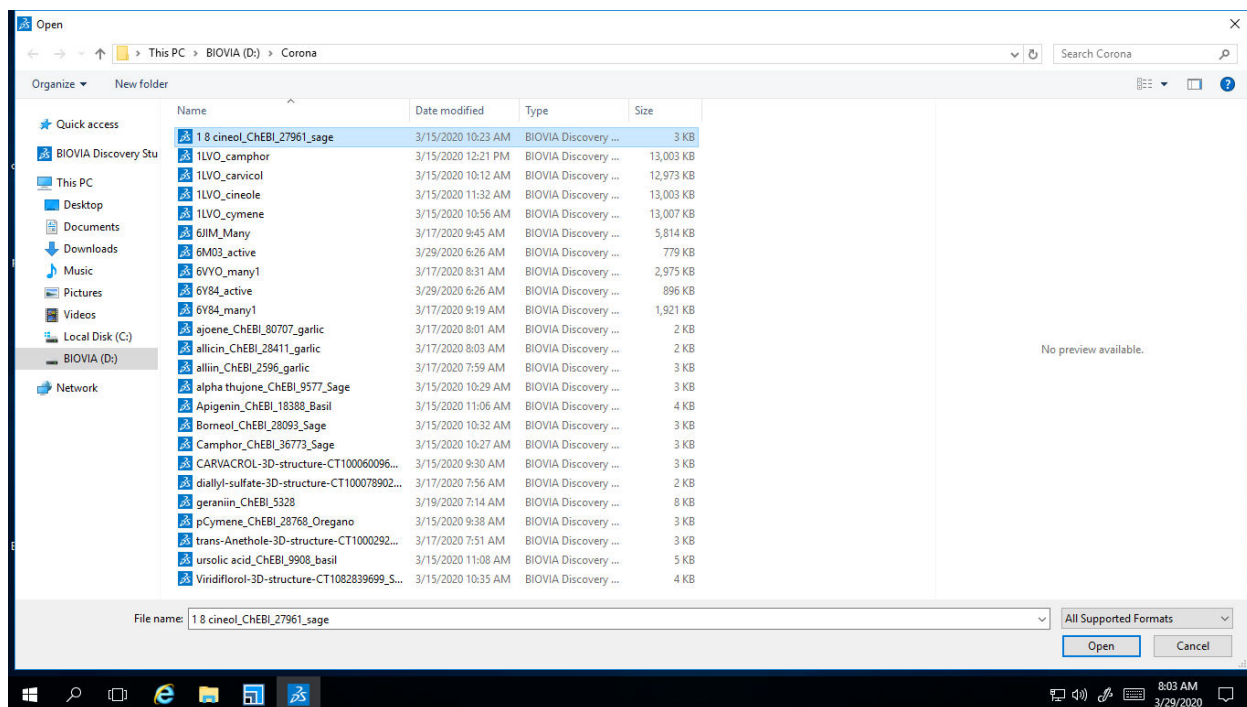
Notifications

My Recent Jobs

- Calculate Protein Formulation ...
- Error Mar-29 07:50
- Build Mutants
- Success Mar-29 07:47
- Model Antibody Loops
- Error Mar-29 07:36
- Verify Protein (MODELER)
- Success Mar-29 07:34
- Dock Ligands (CDOCKER)
- Success Mar-29 06:44
- Dock Ligands (CDOCKER)
- Success Mar-29 06:31
- Search 3D Database
- Success Mar-19 22:28

localhost:9943 20.1.0.19295

8:01 AM 3/29/2020



Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions Pharmacophores Small Molecules X-ray My Tools

Tools Files

Build and Edit Nucleic Acid
Build and Edit Protein
Query Online Databases
Protein Report and Analysis
Prepare Protein
Search Sequences by Similarity
Align Sequences and Structures
Analyze Sequences
Analyze Protein Conservation Pattern
Superimpose Proteins
Create Homology Models
Search Side-Chain Rotamers
Minimize and Refine Protein
Model Antibody
Dock and Analyze Protein Complexes
Design Protein
Predict Protein Formulation Properties
Calculate Protein Formulation Properties...
Calculate Aggregation Scores...
View Aggregation Sites
Define Protein <Undefined>
View at Cutoff Radius: Unset
Step through aggregation sites.
Show Details of Site
Show Surface
Reset Display
Analyze Transmembrane Domains

DS Welcome 1 8 cineol_CHEBI_27961_sage

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cineol	☒ Yes	☐ No	☐ No	CHEBI...	3	CHEBI:35814; CHEB...	Zineol; eucalypt...	1,3,3-trimethyl-2...	C ₁₀ H ₁₈ O	154.249	154.136

localhost:9943 20.1.0.19295

8:03 AM 3/29/2020

Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions Pharmacophores Small Molecules X-ray My Tools

Tools Files

Build and Edit Nucleic Acid
Build and Edit Protein
Query Online Databases
Protein Report and Analysis
Prepare Protein
Search Sequences by Similarity
Align Sequences and Structures
Analyze Sequences
Analyze Protein Conservation Pattern
Superimpose Proteins
Create Homology Models
Search Side-Chain Rotamers
Minimize and Refine Protein
Model Antibody
Dock and Analyze Protein Complexes
Design Protein
Predict Protein Formulation Properties
Calculate Protein Formulation Properties...
Calculate Aggregation Scores...
View Aggregation Sites
Define Protein <Undefined>
View at Cutoff Radius: Unset
Step through aggregation sites.
Show Details of Site
Show Surface
Reset Display
Analyze Transmembrane Domains

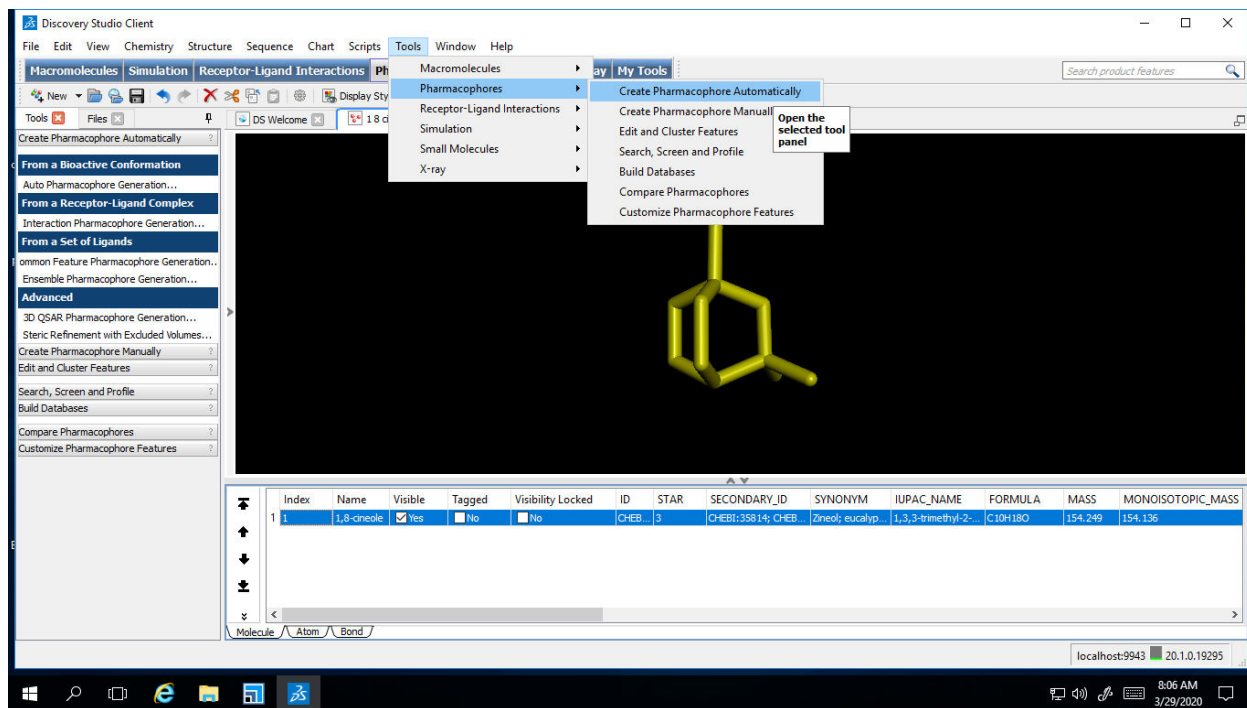
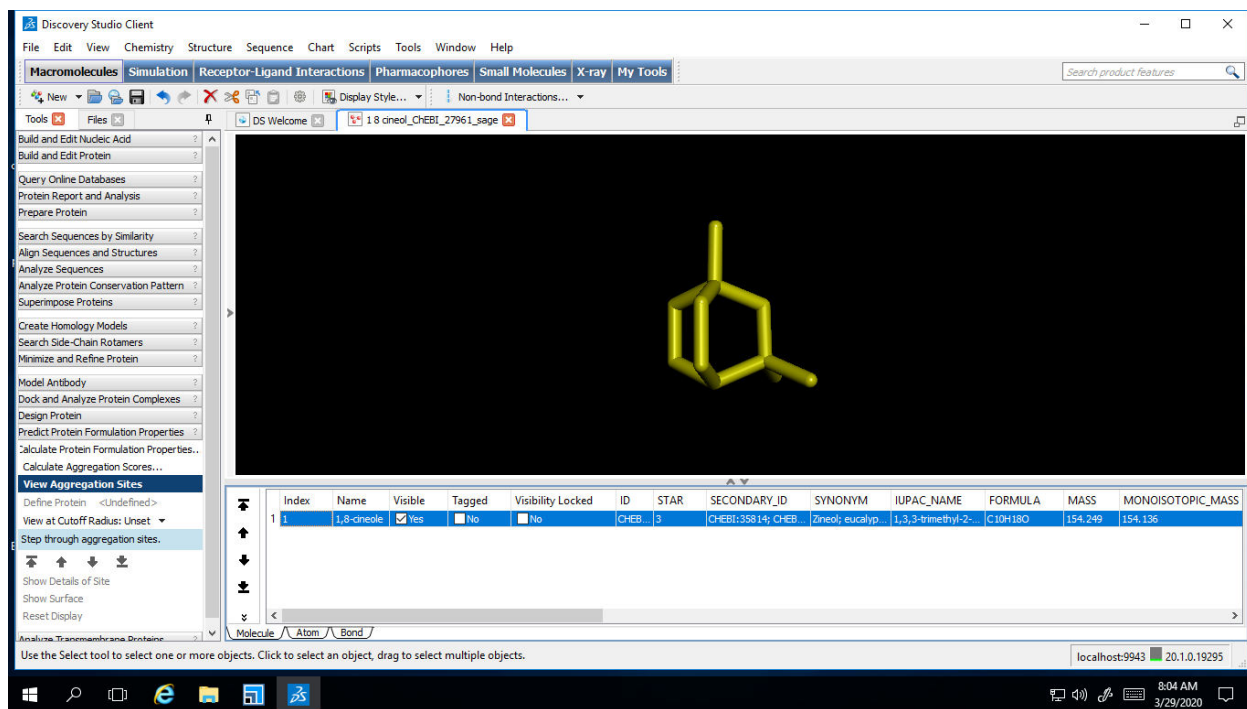
DS Welcome 1 8 cineol_CHEBI_27961_sage

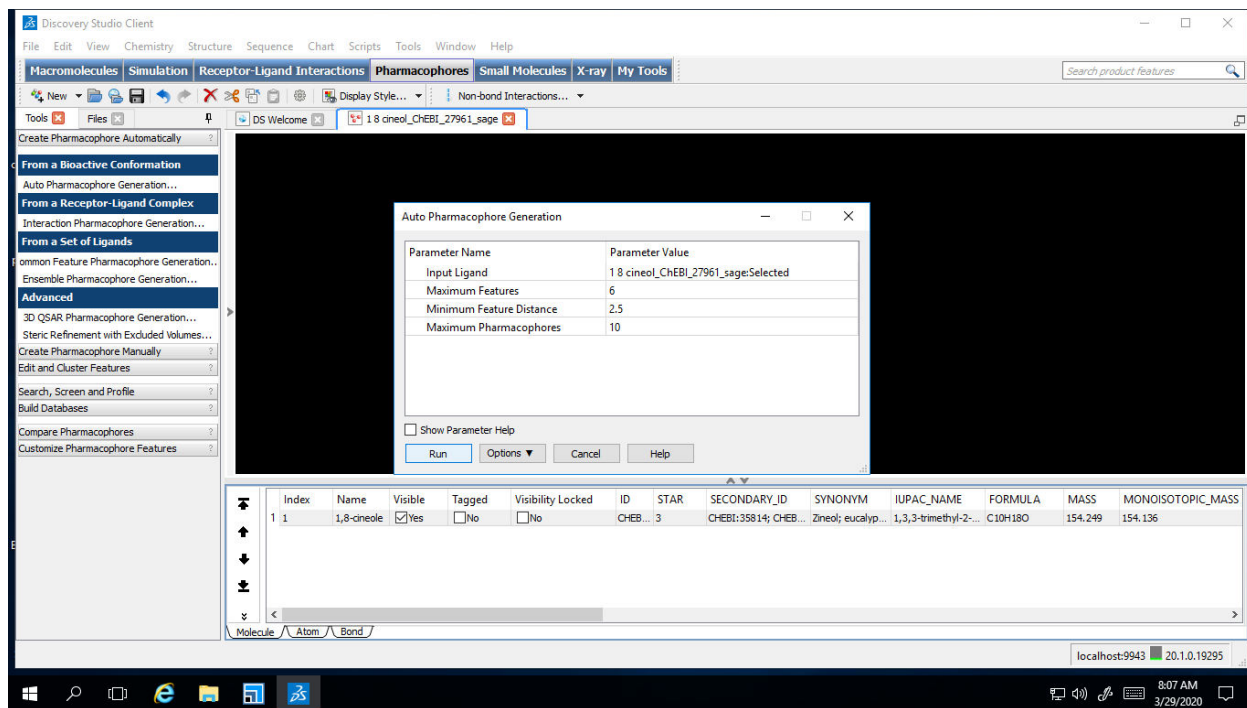
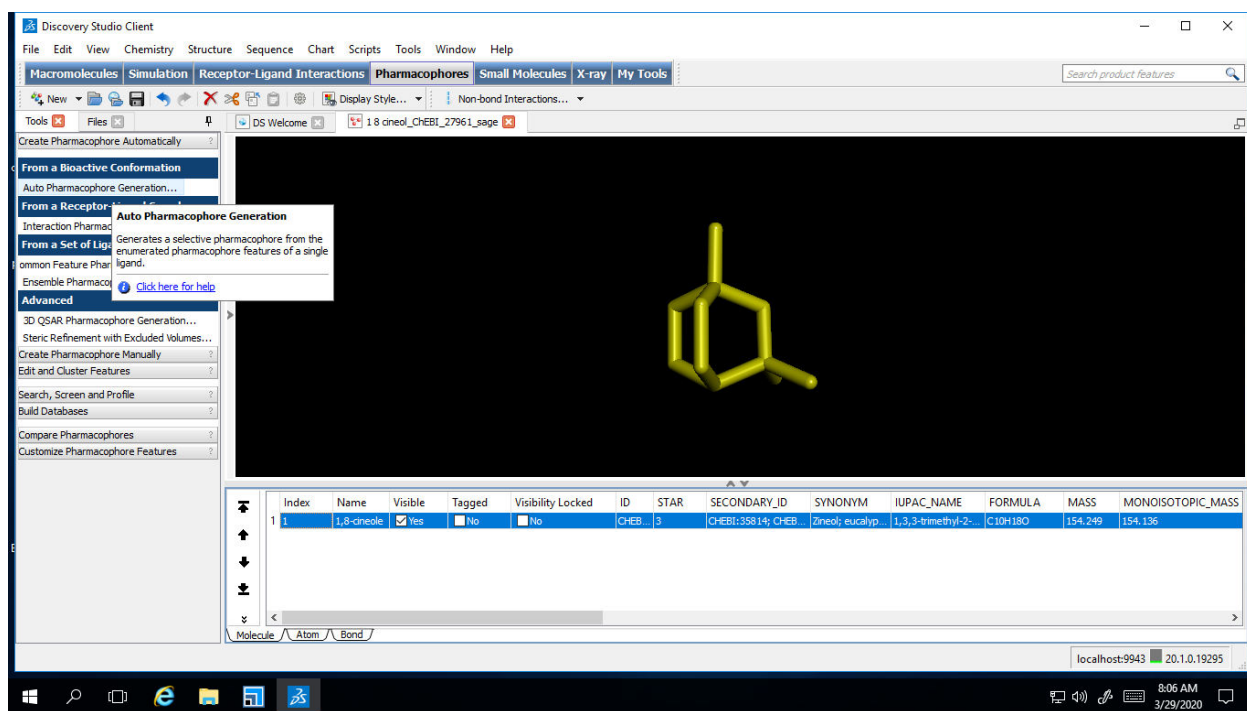
Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cineol	☒ Yes	☐ No	☐ No	CHEBI...	3	CHEBI:35814; CHEB...	Zineol; eucalypt...	1,3,3-trimethyl-2...	C ₁₀ H ₁₈ O	154.249	154.136

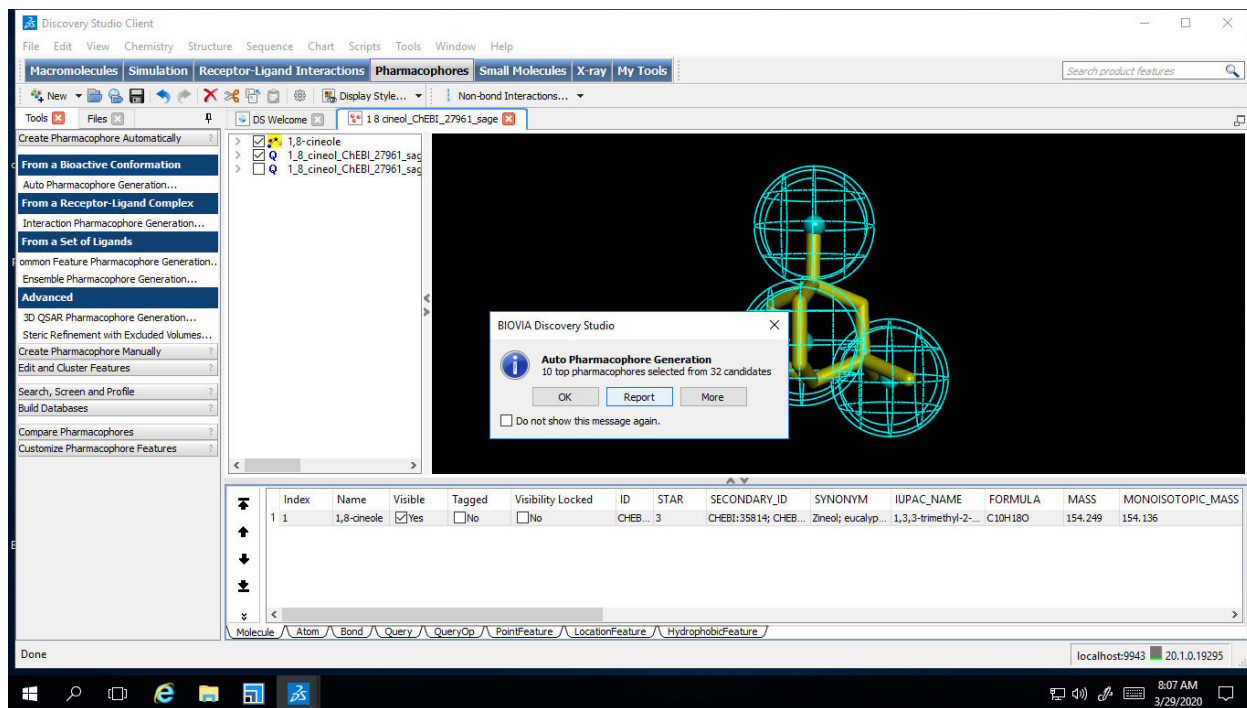
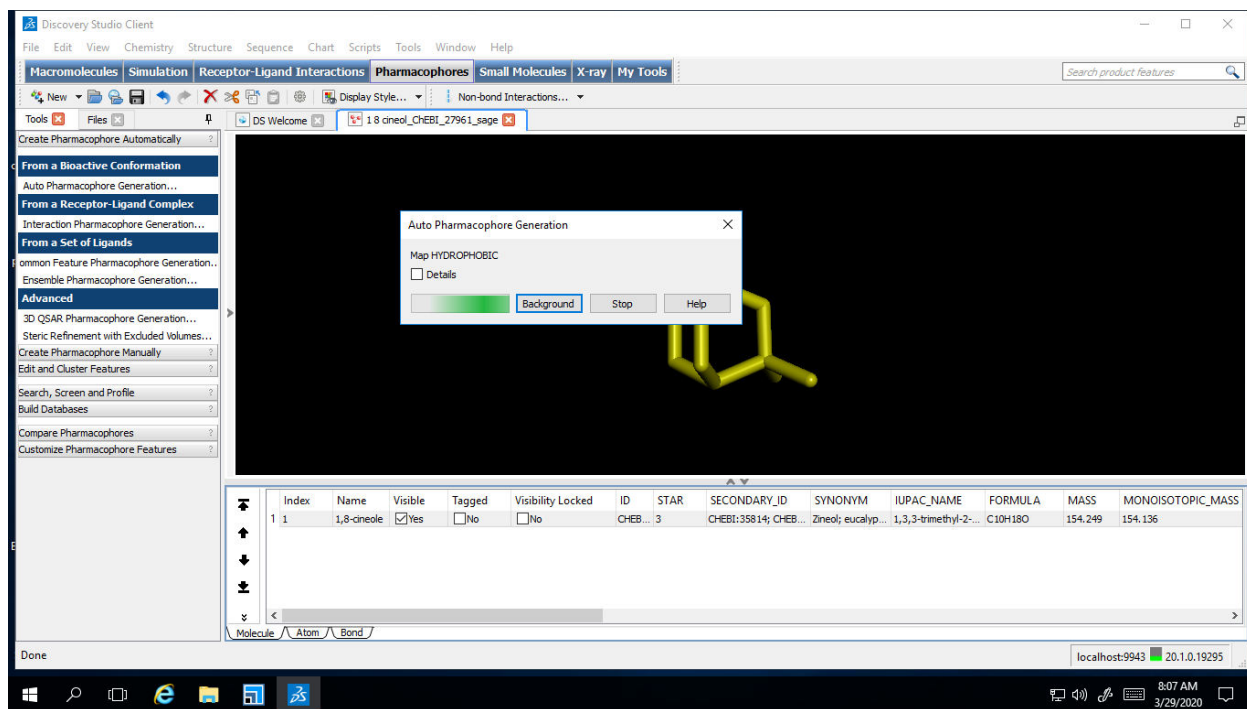
Show In Graphics View
Hide
Show
Show Selected Rows in the Graphics View
Show selected rows in the Graphics View.
Copy
Paste Ctrl+V
Export Rows...
Sort...
Unsort
Select Columns...
Add Attribute...
Rename Attribute...
Remove Attribute(s)...
Color By Activity
List Only Visible Objects
List Only Selected Objects
List Only Tagged Molecules
Tagging
Filter...
Remove Filters
Rearrange Molecules
View

localhost:9943 20.1.0.19295

8:04 AM 3/29/2020







Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

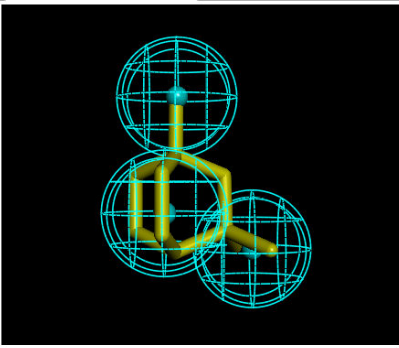
Tools Files

DS Welcome 1 8 cineol_ChEBI_27961_sage Auto Pharmacophore Generation

Create Pharmacophore Automatically

- > From a Bioactive Conformation
- > Auto Pharmacophore Generation...
- > From a Receptor-Ligand Complex
- > Interaction Pharmacophore Generation...
- > From a Set of Ligands
- > Common Feature Pharmacophore Generation...
- > Ensemble Pharmacophore Generation...
- > Advanced
- > 3D QSAR Pharmacophore Generation...
- > Steric Refinement with Excluded Volumes...
- > Create Pharmacophore Manually
- > Edit and Cluster Features
- > Search, Screen and Profile
- > Build Databases
- > Compare Pharmacophores
- > Customize Pharmacophore Features

1,8-cin 1,8_cir 1,8_cir



Auto Pharmacophore Generation

Information

Summary

10 top pharmacophores selected from 32 candidates

Details

Results

Pharmacophore_01
Pharmacophore_02

View Results

Parameters

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Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SEC
1	1,8-cineol	☒ Yes	☐ No	☐ No	CHEB...	3	CHE

Molecule / Atom / Bond / Query / QueryOp / PointFeature / LocationFeature / Hydrophobicity

Done

localhost:9943 20.1.0.19295

8:08 AM 3/29/2020

Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

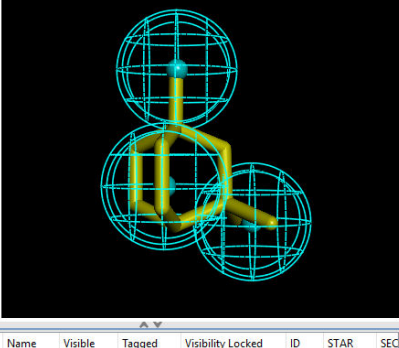
Tools Files

DS Welcome 1 8 cineol_ChEBI_27961_sage Auto Pharmacophore Generation

Create Pharmacophore Automatically

- > From a Bioactive Conformation
- > Auto Pharmacophore Generation...
- > From a Receptor-Ligand Complex
- > Interaction Pharmacophore Generation...
- > From a Set of Ligands
- > Common Feature Pharmacophore Generation...
- > Ensemble Pharmacophore Generation...
- > Advanced
- > 3D QSAR Pharmacophore Generation...
- > Steric Refinement with Excluded Volumes...
- > Create Pharmacophore Manually
- > Edit and Cluster Features
- > Search, Screen and Profile
- > Build Databases
- > Compare Pharmacophores
- > Customize Pharmacophore Features

1,8-cin 1,8_cir 1,8_cir



Auto Pharmacophore Generation

Information

Summary

10 top pharmacophores selected from 32 candidates

Details

Results

Pharmacophore_01
Pharmacophore_02

View Results

Parameters

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Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SEC
1	1,8-cineol	☒ Yes	☐ No	☐ No	CHEB...	3	CHE

Molecule / Atom / Bond / Query / QueryOp / PointFeature / LocationFeature / Hydrophobicity

File:///C:/Users/Deepankar/Documents/Discovery%20Studio/Results/AutoPharmacophoreGeneration_2020_03_29_080715_365/Output/1,8_cineol_ChEBI_27961_sage_01.ds_chm

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

Tools Files

Create Pharmacophore Automatically

- From a Bioactive Conformation
 - Auto Pharmacophore Generation...
 - From a Receptor-Ligand Complex
 - Interaction Pharmacophore Generation...
- From a Set of Ligands
 - Common Feature Pharmacophore Generation...
 - Ensemble Pharmacophore Generation...
- Advanced
 - 3D QSAR Pharmacophore Generation...
 - Steric Refinement with Excluded Volumes...
 - Create Pharmacophore Manually
 - Edit and Cluster Features
 - Search, Screen and Profile
 - Build Databases
 - Compare Pharmacophores
 - Customize Pharmacophore Features

1 8_cineol_CHEBI_27961_sage 1 8_cineol_CHEBI_27961_sage_01 Auto Pharmacophore Generation

Auto Pharmacophore Generation

Information

Summary

10 top pharmacophores selected from 32 candidates

Details

Results

Pharmacophore_01
Pharmacophore_02

View Results

Parameters

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Done localhost:9943 20.1.0.19295

Query: 6 columns (1 column hidden)

Query: \QueryOp \PointFeature \LocationFeature \HydrophobicFeature

Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

Tools Files

Create Pharmacophore Automatically

- From a Bioactive Conformation
 - Auto Pharmacophore Generation...
 - From a Receptor-Ligand Complex
 - Interaction Pharmacophore Generation...
- From a Set of Ligands
 - Common Feature Pharmacophore Generation...
 - Ensemble Pharmacophore Generation...
- Advanced
 - 3D QSAR Pharmacophore Generation...
 - Steric Refinement with Excluded Volumes...
 - Create Pharmacophore Manually
 - Edit and Cluster Features
 - Search, Screen and Profile
 - Build Databases
 - Compare Pharmacophores
 - Customize Pharmacophore Features

1 8_cineol_CHEBI_27961_sage 1 8_cineol_CHEBI_27961_sage_01 Auto Pharmacophore Generation

Auto Pharmacophore Generation

Information

Summary

10 top pharmacophores selected from 32 candidates

Details

Results

Pharmacophore_01
Pharmacophore_02

View Results

Parameters

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Done localhost:9943 20.1.0.19295

Query: 6 columns (1 column hidden)

Query: \QueryOp \PointFeature \LocationFeature \HydrophobicFeature

Name	ID	Visible	Color	Parent	X	Y	Z	Center of Mass	Type
1 Hydropho...	0	☒ Yes	■	HYDROPH...	-0.14	2.86	0.12	☐ No	Hyd
2 Hydropho...	0	☒ Yes	■	HYDROPH...	1.9	-1.22	-0.48	☐ No	Hyd
3 Hydropho...	0	☒ Yes	■	HYDROPH...	-0.42	-0.18	1.4	☐ No	Hyd

Pharmacophore feature

Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Tools Files

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features
Find Features
From an input ligand or ligands.
Feature Mapping...
From a binding site.
Interaction Generation...
Select Feature Category
Select the target feature category to be used for the commands below.
Current Feature: All Features
Show or hide the feature or the location constraints of the category selected above or any non-feature related objects (e.g., a protein).
Show/Hide All Features
Show/Hide Location Spheres
Show/Hide All Non-Features
Cluster
Perform a hierarchical clustering of the feature category selected above.
Cluster Current Feature
Show Only Cluster Centers
Keep Only Cluster Centers
Edit
Keep Only Current Feature Selections
Merge Hypotheses
Average Pharmacophore Features

1 8 cineol_CHEBI_27961_sage

Auto Pharmacophore Generation

Information
Summary
Details
Results
Parameters

10 top pharmacophores selected from 32 candidates

Pharmacophore_01
Pharmacophore_02
View Results

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Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SEC
1	1,8-cineol	☒ Yes	☐ No	☐ No	CHEB...	3	CHEB...

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Tools Files

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features
Find Features
From an input ligand or ligands.
Feature Mapping...
From a binding site.
Interaction Generation...
Select Feature Category
Select the target feature category to be used for the commands below.
Current Feature: All Features
Show or hide the feature or the location constraints of the category selected above or any non-feature related objects (e.g., a protein).
Show/Hide All Features
Show/Hide Location Spheres
Show/Hide All Non-Features
Cluster
Perform a hierarchical clustering of the feature category selected above.
Cluster Current Feature
Show Only Cluster Centers
Keep Only Cluster Centers
Edit
Keep Only Current Feature Selections
Merge Hypotheses
Average Pharmacophore Features

1 8 cineol_CHEBI_27961_sage

Auto Pharmacophore Generation

Information
Summary
Details
Results
Parameters

10 top pharmacophores selected from 32 candidates

Pharmacophore_01
Pharmacophore_02
View Results

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Feature Mapping

Parameter Name Parameter Value
Input Ligands 1 8 cineol_CHEBI_27961_sage:Visible
Features HB_ACCEPTOR,HB_DONOR,HYDROPHOBIC,NEG...
> Advanced

☐ Show Parameter Help

Run Options Cancel Help

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SEC
1	1,8-cineol	☒ Yes	☐ No	☐ No	CHEB...	3	CHEB...

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cineol_CHEBI_27961_sage

Auto Pharmacophore Generation

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Find Features
From an input ligand or ligands.
Feature Mapping...
From a binding site.

Interaction Generation...

Select Feature Category
Select the target feature category to be used for the commands below.
Current Feature: All Features
Show or hide the feature or the location constraints of the category selected above or any non-feature related objects (e.g., a protein).
Show/Hide All Features
Show/Hide Location Spheres
Show/Hide All Non-Features

Cluster
Perform a hierarchical clustering of the feature category selected above.
Cluster Current Feature
Show Only Cluster Centers
Keep Only Cluster Centers

Edit
Keep Only Current Feature Selections
Merge Hypotheses
Average Pharmacophore Features

1 8 cineol_CHEBI_27961_sage

1,8-cin
1,8_cin
1,8_cin
hypo

Auto Pharmacophore Generation

Information
Summary
10 top pharmacophores selected from 32 candidates
Details
Results
Pharmacophore_01
Pharmacophore_02
View Results
Parameters

BIOVIA Discovery Studio
Feature Mapping
Found 6 features in ligand: 1_8_cineol_CHEBI_27961_sage
OK Report More
Do not show this message again.

Index Name Visible Tagged Visibility Locked ID STAR SEC
1 1 1,8-cineol [x] Yes [] No [] No CHEB... 3 CHE...

Molecule Atom Bond Query QueryOp PointFeature LocationFeature HydrophobicFeature

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cineol_CHEBI_27961_sage

Auto Pharmacophore Generation

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Find Features
From an input ligand or ligands.
Feature Mapping...
From a binding site.

Interaction Generation...

Select Feature Category
Select the target feature category to be used for the commands below.
Current Feature: All Features
Show or hide the feature or the location constraints of the category selected above or any non-feature related objects (e.g., a protein).
Show/Hide All Features
Show/Hide Location Spheres
Show/Hide All Non-Features

Cluster
Perform a hierarchical clustering of the feature category selected above.
Cluster Current Feature
Show Only Cluster Centers
Keep Only Cluster Centers

Edit
Keep Only Current Feature Selections
Merge Hypotheses
Average Pharmacophore Features

1 8 cineol_CHEBI_27961_sage

1,8-cin
1,8_cin
1,8_cin
hypo

Auto Pharmacophore Generation

Information
Summary
10 top pharmacophores selected from 32 candidates
Details
Results
Pharmacophore_01
Pharmacophore_02
View Results
Parameters

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1 Name ID Visible Color Parent X Y Z Center of Mass
1 Hydropho... 0 [] No HYDROPH... -0.14 2.86 0.12 [] No
2 Hydropho... 0 [] No HYDROPH... 1.9 -1.22 -0.48 [] No
3 Hydropho... 0 [] No HYDROPH... -0.42 -0.18 1.4 [] No
4 Hydropho... 0 [] No HYDROPH... -0.14 2.86 0.12 [] No
5 Hydropho... 0 [] No HYDROPH... 1.9 -1.22 -0.48 [] No
6 Hydropho... 0 [] No HYDROPH... -0.82 -0.14 0.4 [] No

Molecule Atom Query QueryOp PointFeature LocationFeature HydrophobicFeature

PointFeature: 11 columns (1 column hidden)

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Pharmacophore database search

Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cineol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Find Features
From an input ligand or ligands.
Feature Mapping...
From a binding site.

Interaction Generation...

Select Feature Category
Select the target feature category to be used for the commands below.
Current Feature: All Features
Show or hide the feature or the location constraints of the category selected above or any non-feature related objects (e.g., a protein).
Show/Hide All Features
Show/Hide Location Spheres
Show/Hide All Non-Features

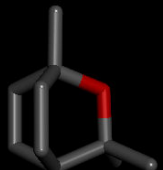
Cluster
Perform a hierarchical clustering of the feature category selected above.
Cluster Current Feature
Show Only Cluster Centers
Keep Only Cluster Centers

Edit
Keep Only Current Feature Selections
Merge Hypotheses
Average Pharmacophore Features

Use the Select tool to select one or more objects. Click to select an object, drag to select multiple objects.

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8:21 AM 3/29/2020



Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cineol	☒ Yes	☐ No	☐ No	CHEBI...	3	CHEBI:35814; CHEB...	Zneol; eucalyp...	1,3,3-trimethyl-2-...	C10H18O	154.249	154.136

Molecule Atom Bond

Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cineol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Find Features
From an input ligand or ligands.
Feature Mapping...
From a binding site.

Interaction Generation...

Select Feature Category
Select the target feature category to be used for the commands below.
Current Feature: All Features
Show or hide the feature or the location constraints of the category selected above or any non-feature related objects (e.g., a protein).
Show/Hide All Features
Show/Hide Location Spheres
Show/Hide All Non-Features

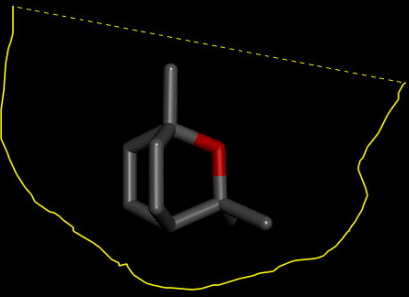
Cluster
Perform a hierarchical clustering of the feature category selected above.
Cluster Current Feature
Show Only Cluster Centers
Keep Only Cluster Centers

Edit
Keep Only Current Feature Selections
Merge Hypotheses
Average Pharmacophore Features

Use the Select tool to select one or more objects. Click to select an object, drag to select multiple objects.

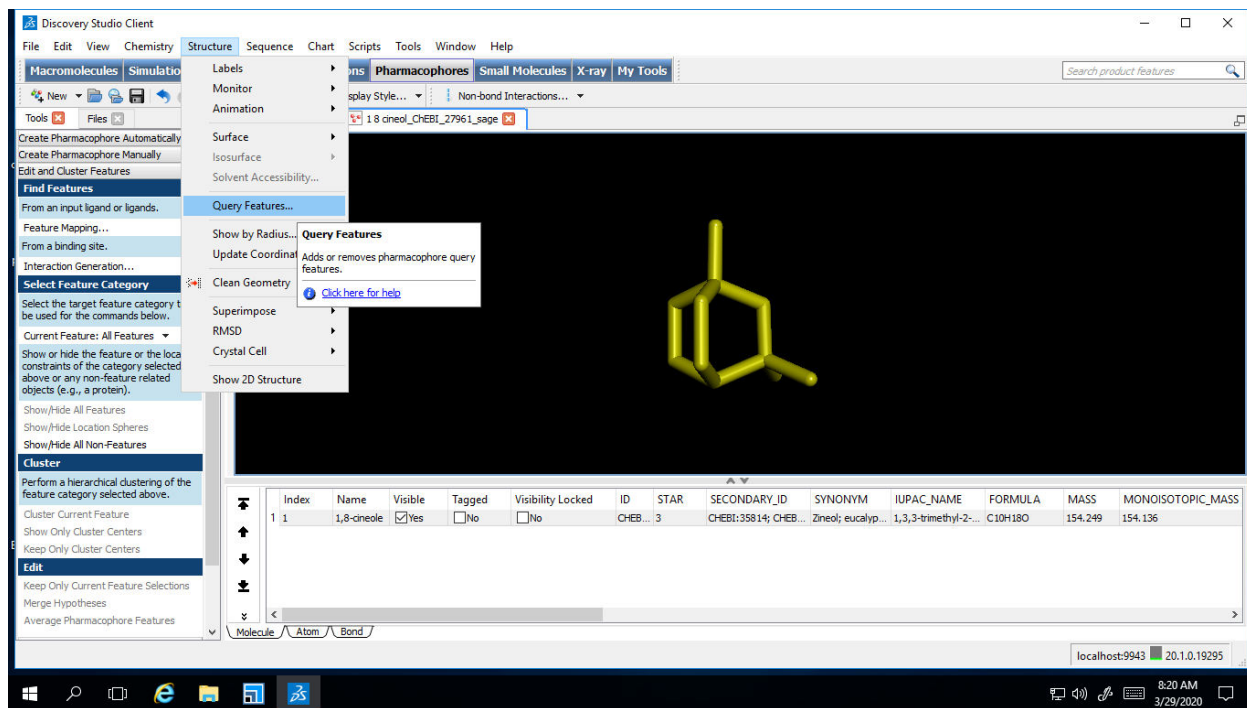
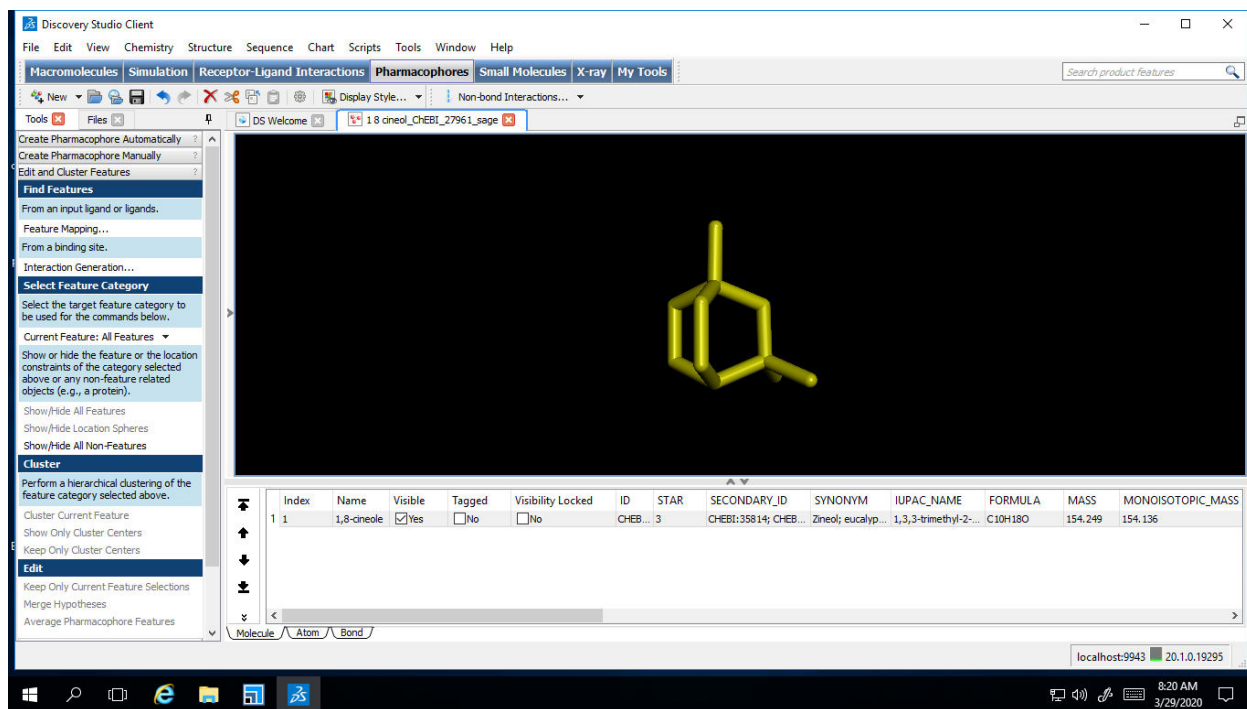
localhost:9943 20.1.0.19295

8:19 AM 3/29/2020



Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cineol	☒ Yes	☐ No	☐ No	CHEBI...	3	CHEBI:35814; CHEB...	Zneol; eucalyp...	1,3,3-trimethyl-2-...	C10H18O	154.249	154.136

Molecule Atom Bond



Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions Pharmacophores Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cneol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Find Features
From an input ligand or ligands.
Feature Mapping...
From a binding site.

Interaction Generation...

Select Feature Category
Select the target feature category to be used for the commands below.
Current Feature: All Features
Show or hide the feature or the location constraints of the category selected above or any non-feature related objects (e.g., a protein).
Show/Hide All Features
Show/Hide Location Spheres
Show/Hide All Non-Features

Cluster
Perform a hierarchical clustering of the feature category selected above.
Cluster Current Feature
Show Only Cluster Centers
Keep Only Cluster Centers

Edit
Keep Only Current Feature Selections
Merge Hypotheses
Average Pharmacophore Features

Create Feature

Feature: Centroid
Type: Centroid
Name: Centroid

OK Cancel Help

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cneole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zneol; eucalyp...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions Pharmacophores Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cneol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Find Features
From an input ligand or ligands.
Feature Mapping...
From a binding site.

Interaction Generation...

Select Feature Category
Select the target feature category to be used for the commands below.
Current Feature: All Features
Show or hide the feature or the location constraints of the category selected above or any non-feature related objects (e.g., a protein).
Show/Hide All Features
Show/Hide Location Spheres
Show/Hide All Non-Features

Cluster
Perform a hierarchical clustering of the feature category selected above.
Cluster Current Feature
Show Only Cluster Centers
Keep Only Cluster Centers

Edit
Keep Only Current Feature Selections
Merge Hypotheses
Average Pharmacophore Features

Create Feature

Feature: Centroid
Centroid
ExclusionSphere
Line
Location
Normal
Plane
Point (Pharmacophore)
InteractionPoint (Docking)
Shape
QueryAtom

OK Cancel Help

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cneole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zneol; eucalyp...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond

localhost:9943 20.1.0.19295

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cneol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Find Features
From an input ligand or ligands.
Feature Mapping...
From a binding site.

Interaction Generation...

Select Feature Category
Select the target feature category to be used for the commands below.
Current Feature: All Features
Show or hide the feature or the location constraints of the category selected above or any non-feature related objects (e.g., a protein).
Show/Hide All Features
Show/Hide Location Spheres
Show/Hide All Non-Features

Cluster
Perform a hierarchical clustering of the feature category selected above.
Cluster Current Feature
Show Only Cluster Centers
Keep Only Cluster Centers

Edit
Keep Only Current Feature Selections
Merge Hypotheses
Average Pharmacophore Features

Create Feature

Feature: Point (Pharmacophore) OK
Type: Hydrophobe
Name: Point Cancel Help

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cneole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zneol; eucalyp...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond

localhost:9943 20.1.0.19295

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cneol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Find Features
From an input ligand or ligands.
Feature Mapping...
From a binding site.

Interaction Generation...

Select Feature Category
Select the target feature category to be used for the commands below.
Current Feature: All Features
Show or hide the feature or the location constraints of the category selected above or any non-feature related objects (e.g., a protein).
Show/Hide All Features
Show/Hide Location Spheres
Show/Hide All Non-Features

Cluster
Perform a hierarchical clustering of the feature category selected above.
Cluster Current Feature
Show Only Cluster Centers
Keep Only Cluster Centers

Edit
Keep Only Current Feature Selections
Merge Hypotheses
Average Pharmacophore Features

Hydrophobe

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cneole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zneol; eucalyp...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond Query QueryOp PointFeature HydrophobicFeature Label

Use the Select tool to select one or more objects. Click to select an object, drag to select multiple objects.

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions Pharmacophores Small Molecules X-ray My Tools

Tools Files

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Find Features
From an input ligand or ligands.
Feature Mapping...
From a binding site.

Interaction Generation...

Select Feature Category
Select the target feature category to be used for the commands below.
Current Feature: All Features
Show or hide the feature or the location constraints of the category selected above or any non-feature related objects (e.g., a protein).
Show/Hide All Features
Show/Hide Location Spheres
Show/Hide All Non-Features

Cluster
Perform a hierarchical clustering of the feature category selected above.
Cluster Current Feature
Show Only Cluster Centers
Keep Only Cluster Centers

Edit
Keep Only Current Feature Selections
Merge Hypotheses
Average Pharmacophore Features

Open the selected tool panel

Hydrophobe

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cineole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zineol; eucalypt...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond Query QueryOp PointFeature HydrophobicFeature Label

localhost:9943 20.1.0.19295

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions Pharmacophores Small Molecules X-ray My Tools

Tools Files

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Search, Screen and Profile

Mapping
Map multiple ligands to a single pharmacophore for detailed analysis of multiple mappings.
Ligand Pharmacophore Mapping...

Screening
Maps multiple ligands to a single pharmacophore with optional partial mapping, required features and group features.
Screen Library...

Profile Ligands using Pharmacophore
Map multiple ligands to multiple pharmacophores.
Ligand Profiler...

Search DB - Indexed for Speed
Search an indexed multi-conformer database with a single pharmacophore. This is optimized for speed and efficiency. Use [Build 3D Database](#) to build a database.

Search 3D Database...

Browse DB
Browse 3D Database...

Build Databases

Compare Pharmacophores
Customize Pharmacophore Features

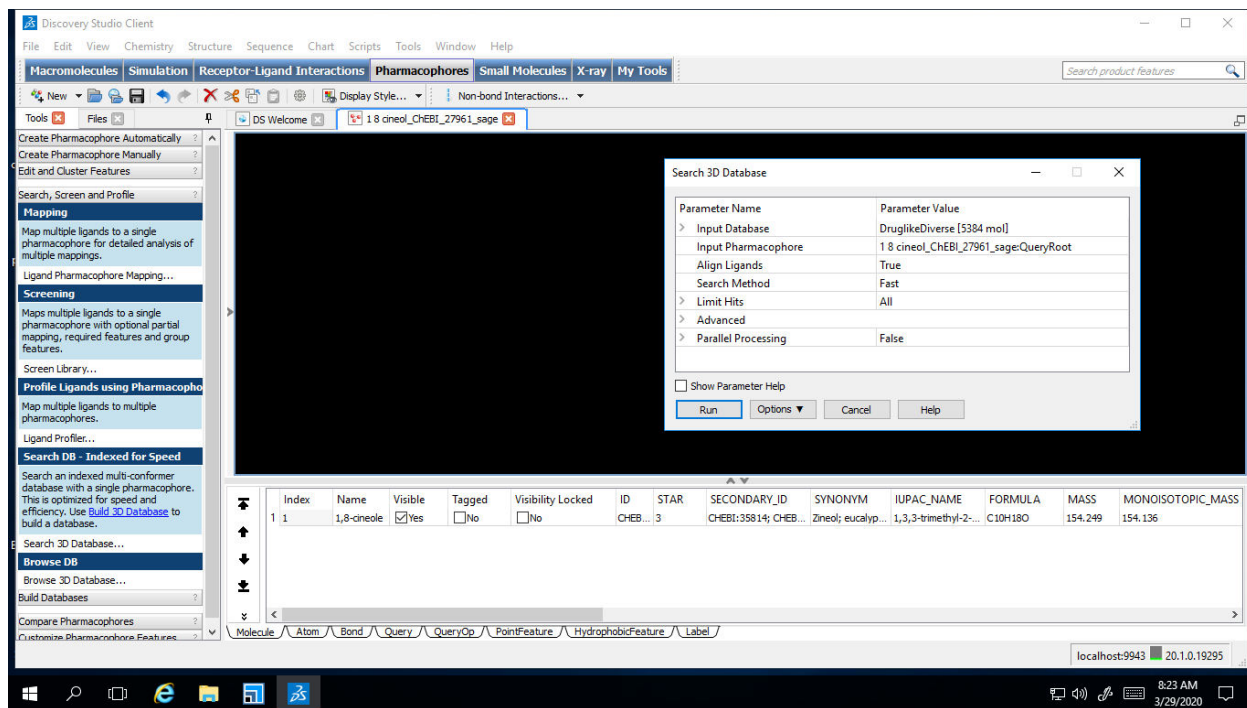
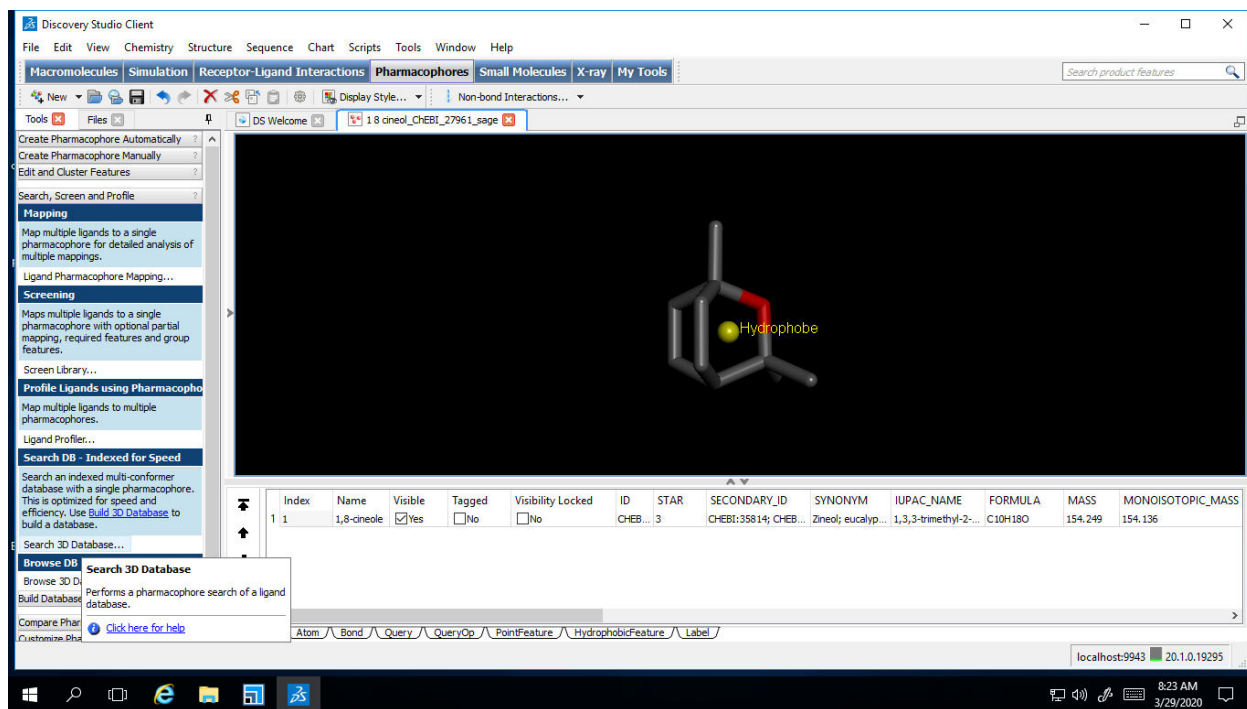
Hydrophobe

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cineole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zineol; eucalypt...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond Query QueryOp PointFeature HydrophobicFeature Label

localhost:9943 20.1.0.19295

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions Pharmacophores Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cineol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Search, Screen and Profile

Mapping
Map multiple ligands to a single pharmacophore for detailed analysis of multiple mappings.
Ligand Pharmacophore Mapping...

Screening
Maps multiple ligands to a single pharmacophore with optional partial mapping, required features and group features.
Screen Library...

Profile Ligands using Pharmacophores
Map multiple ligands to multiple pharmacophores.
Ligand Profiler...

Search DB - Indexed for Speed
Search an indexed multi-conformer database with a single pharmacophore. This is optimized for speed and efficiency. Use [Build 3D Database](#) to build a database.

Search 3D Database...
Browse DB
Browse 3D Database...
Build Databases

Compare Pharmacophores
Customize Pharmacophore Features...

Search 3D Database

Parameter Name	Parameter Value
> Input Database	DruglikeDiverse [5384 mol]
Input Pharmacophore	1 8 cineol_CHEBI_27961_sage:QueryRoot
Align Ligands	True
Search Method	Fast
> Limit Hits	All
> Advanced	First N
> Parallel Processing	Best N

☐ Show Parameter Help

Run Options Cancel Help

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cineole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zineol; eucalypt...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond Query QueryOp PointFeature HydrophobicFeature Label

localhost:9943 20.1.0.19295

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions Pharmacophores Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cineol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Search, Screen and Profile

Mapping
Map multiple ligands to a single pharmacophore for detailed analysis of multiple mappings.
Ligand Pharmacophore Mapping...

Screening
Maps multiple ligands to a single pharmacophore with optional partial mapping, required features and group features.
Screen Library...

Profile Ligands using Pharmacophores
Map multiple ligands to multiple pharmacophores.
Ligand Profiler...

Search DB - Indexed for Speed
Search an indexed multi-conformer database with a single pharmacophore. This is optimized for speed and efficiency. Use [Build 3D Database](#) to build a database.

Search 3D Database...
Browse DB
Browse 3D Database...
Build Databases

Compare Pharmacophores
Customize Pharmacophore Features...

Search 3D Database

Parameter Name	Parameter Value
> Input Database	DruglikeDiverse [5384 mol]
Input Pharmacophore	1 8 cineol_CHEBI_27961_sage:QueryRoot
Align Ligands	True
Search Method	Fast
> Limit Hits	Best N
Maximum Hits	300
> Advanced	
> Parallel Processing	False

☐ Show Parameter Help

Run Options Cancel Help

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cineole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zineol; eucalypt...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond Query QueryOp PointFeature HydrophobicFeature Label

localhost:9943 20.1.0.19295

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cineol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Search, Screen and Profile

Mapping
Map multiple ligands to a single pharmacophore for detailed analysis of multiple mappings.
Ligand Pharmacophore Mapping...

Screening
Maps multiple ligands to a single pharmacophore with optional partial mapping, required features and group features.
Screen Library...

Profile Ligands using Pharmacophores
Map multiple ligands to multiple pharmacophores.
Ligand Profiler...

Search DB - Indexed for Speed
Search an indexed multi-conformer database with a single pharmacophore. This is optimized for speed and efficiency. Use [Build 3D Database](#) to build a database.

Search 3D Database...
Browse DB
Browse 3D Database...
Build Databases

Compare Pharmacophores
Customize Pharmacophore Features...

Search 3D Database

Parameter Name Parameter Value

> Input Database DruglikeDiverse [5384 mol]

Input Pharmacophore 1 8 cineol_CHEBI_27961_sage:QueryRoot

Align Ligands True

Search Method Fast

Limit Hits Best N

Maximum Hits 5

> Advanced

> Parallel Processing False

☐ Show Parameter Help

Run Options Cancel Help

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cineole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zineol; eucalyp...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond Query QueryOp PointFeature HydrophobicFeature Label

localhost:9943 20.1.0.19295

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cineol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Search, Screen and Profile

Mapping
Map multiple ligands to a single pharmacophore for detailed analysis of multiple mappings.
Ligand Pharmacophore Mapping...

Screening
Maps multiple ligands to a single pharmacophore with optional partial mapping, required features and group features.
Screen Library...

Profile Ligands using Pharmacophores
Map multiple ligands to multiple pharmacophores.
Ligand Profiler...

Search DB - Indexed for Speed
Search an indexed multi-conformer database with a single pharmacophore. This is optimized for speed and efficiency. Use [Build 3D Database](#) to build a database.

Search 3D Database...
Browse DB
Browse 3D Database...
Build Databases

Compare Pharmacophores
Customize Pharmacophore Features...

Search 3D Database

Database Search

☐ Details

Background Stop Help

Hydrophobe

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cineole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zineol; eucalyp...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond Query QueryOp PointFeature HydrophobicFeature Label

localhost:9943 20.1.0.19295

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Search 3D Database has been launched

Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions Pharmacophores Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cneol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features
Search, Screen and Profile
Mapping
Map multiple ligands to a single pharmacophore for detailed analysis of multiple mappings.
Ligand Pharmacophore Mapping...
Screening
Maps multiple ligands to a single pharmacophore with optional partial mapping, required features and group features.
Screen Library...
Profile Ligands using Pharmacophore
Map multiple ligands to multiple pharmacophores.
Ligand Profiler...
Search DB - Indexed for Speed
Search an indexed multi-conformer database with a single pharmacophore. This is optimized for speed and efficiency. Use [Build 3D Database](#) to build a database.
Search 3D Database...
Browse DB
Browse 3D Database...
Build Databases
Compare Pharmacophores
Customize Pharmacophore Features

Search 3D Database

100% complete
☐ Details
Background Stop Help

Hydrophobe

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cneole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zneol; eucalyp...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond Query QueryOp PointFeature HydrophobicFeature Label

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions Pharmacophores Small Molecules X-ray My Tools

Search product features

Tools Files

DS Welcome 1 8 cneol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features
Search, Screen and Profile
Mapping
Map multiple ligands to a single pharmacophore for detailed analysis of multiple mappings.
Ligand Pharmacophore Mapping...
Screening
Maps multiple ligands to a single pharmacophore with optional partial mapping, required features and group features.
Screen Library...
Profile Ligands using Pharmacophore
Map multiple ligands to multiple pharmacophores.
Ligand Profiler...
Search DB - Indexed for Speed
Search an indexed multi-conformer database with a single pharmacophore. This is optimized for speed and efficiency. Use [Build 3D Database](#) to build a database.
Search 3D Database...
Browse DB
Browse 3D Database...
Build Databases
Compare Pharmacophores
Customize Pharmacophore Features

Search 3D Database

100% complete
☐ Details
Background Stop Help

Hydrophobe

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MASS
1	1,8-cneole	☒ Yes	☐ No	☐ No	CHEB...	3	CHEBI:35814; CHEB...	Zneol; eucalyp...	1,3,3-trimethyl-2...	C10H18O	154.249	154.136

Molecule Atom Bond Query QueryOp PointFeature HydrophobicFeature Label

localhost:9943 20.1.0.19295

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Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Tools Files

DS Welcome 1 8 cineol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features
Search, Screen and Profile

Mapping
Map multiple ligands to a single pharmacophore for detailed analysis of multiple mappings.
Ligand Pharmacophore Mapping...

Screening
Maps multiple ligands to a single pharmacophore with optional partial mapping, required features and group features.
Screen Library...

Profile Ligands using Pharmacophore
Map multiple ligands to multiple pharmacophores.
Ligand Profiler...

Search DB - Indexed for Speed
Search an indexed multi-conformer database with a single pharmacophore. This is optimized for speed and efficiency. Use [Build 3D Database](#) to build a database.
Search 3D Database...
Browse DB
Browse 3D Database...
Build Databases

Compare Pharmacophores
Customize Pharmacophore Features

Done

localhost:9943 20.1.0.19295

BIOVIA Discovery Studio

Search 3D Database
5 hits returned from search of DrugLikeDiverse database using 1_8_cineol_CHEBI_27961_sage.ds_dhm The results have been added to the window.

☐ Do not show this message again.

OK Report

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MA
1	1,8-cineole	☒ Yes	☐ No	☐ No								
2	CAP0000...	☐ No	☐ No	☐ No								
3	CAP0000...	☐ No	☐ No	☐ No								
4	CAP0000...	☐ No	☐ No	☐ No								
5	CAP0000...	☐ No	☐ No	☐ No								
6	CAP0000...	☐ No	☐ No	☐ No								

Molecule Atom Bond Query QueryOp PointFeature HydrophobicFeature Label

Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

Tools Files

DS Welcome 1 8 cineol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features
Search, Screen and Profile

Mapping
Map multiple ligands to a single pharmacophore for detailed analysis of multiple mappings.
Ligand Pharmacophore Mapping...

Screening
Maps multiple ligands to a single pharmacophore with optional partial mapping, required features and group features.
Screen Library...

Profile Ligands using Pharmacophore
Map multiple ligands to multiple pharmacophores.
Ligand Profiler...

Search DB - Indexed for Speed
Search an indexed multi-conformer database with a single pharmacophore. This is optimized for speed and efficiency. Use [Build 3D Database](#) to build a database.
Search 3D Database...
Browse DB
Browse 3D Database...
Build Databases

Compare Pharmacophores
Customize Pharmacophore Features

Use the Select tool to select one or more objects. Click to select an object, drag to select multiple objects.

localhost:9943 20.1.0.19295

Hydrophobe

Index	Name	Visible	Tagged	Visibility Locked	ID	STAR	SECONDARY_ID	SYNONYM	IUPAC_NAME	FORMULA	MASS	MONOISOTOPIC_MA
1	1,8-cineole	☒ Yes	☐ No	☐ No								
2	CAP0000...	☐ No	☐ No	☐ No								
3	CAP0000...	☐ No	☐ No	☐ No								
4	CAP0000...	☐ No	☐ No	☐ No								
5	CAP0000...	☐ No	☐ No	☐ No								
6	CAP0000...	☐ No	☐ No	☐ No								

Molecule Atom Bond Query QueryOp PointFeature HydrophobicFeature Label

Discovery Studio Client

File Edit View Chemistry Structure Sequence Chart Scripts Tools Window Help

Macromolecules Simulation Receptor-Ligand Interactions **Pharmacophores** Small Molecules X-ray My Tools

New Open Recent Save Undo Redo Copy Paste Delete Display Style... Non-bond Interactions...

Tools Files

DS Welcome 1 8 cineol_CHEBI_27961_sage

Create Pharmacophore Automatically
Create Pharmacophore Manually
Edit and Cluster Features

Search, Screen and Profile

Mapping
Map multiple ligands to a single pharmacophore for detailed analysis of multiple mappings.
Ligand Pharmacophore Mapping...

Screening
Maps multiple ligands to a single pharmacophore with optional partial mapping, required features and group features.
Screen Library...

Profile Ligands using Pharmacophores
Map multiple ligands to multiple pharmacophores.
Ligand Profiler...

Search DB - Indexed for Speed
Search an indexed multi-conformer database with a single pharmacophore. This is optimized for speed and efficiency. Use [Build 3D Database](#) to build a database.
Search 3D Database...

Browse DB
Browse 3D Database...
Build Databases

Compare Pharmacophores
Customize Pharmacophore Features

1 8 cineol_CHEBI_27961_sage

Index Name Visible Tagged Visibility Locked ID STAR SECONDARY_ID SYNONYM IUPAC_NAME FORMULA MASS MONOISOTOPIC_MA

1	1,8-cineole	☐ No	☐ No	☐ No	CHEBI:35814; CHEBI:35814	3		Zineol; eucalyptol	1,3,3-trimethyl-2-methyl-5-propenylcyclohex-2-ene	C10H18O	154.249	154.136
2	CAP00000...	☒ Yes	☐ No	☐ No								
3	CAP00000...	☐ No	☐ No	☐ No								
4	CAP00000...	☐ No	☐ No	☐ No								
5	CAP00000...	☐ No	☐ No	☐ No								
6	CAP00000...	☐ No	☐ No	☐ No								

Molecule / Atom / Bond / Query / QueryOp / PointFeature / HydrophobicFeature / Label

localhost:9943 20.1.0.19295

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