

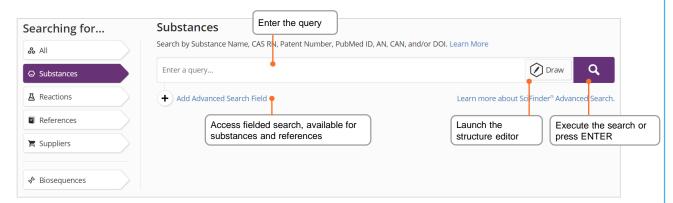
Quick Reference Guide

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Interface and Reference Search

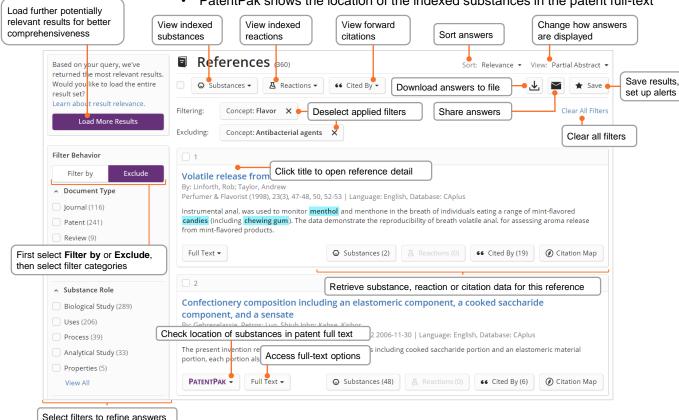


Search Interface SciFinderⁿ features a streamlined search interface.

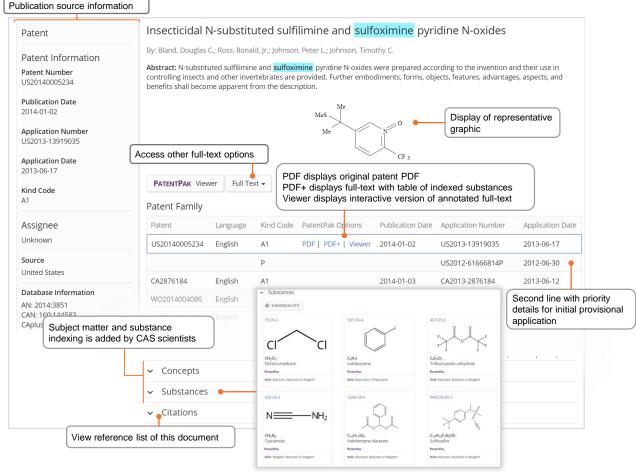


Reference Search The References display features visualizations, dynamic facets and an easy-to-use layout

- · References are ranked and sorted by relevance
- You may save your searches, send a link or set-up alerts
- · Filters allow you to focus the answers
- PatentPak shows the location of the indexed substances in the patent full-text



Reference detail and search



Boolean Operators Logical operators define precise text queries

Use parentheses to group logical expressions such as OR'ed synonyms, e.g.: (flavor or odor) and menthol

AND Requires both concepts to be present within the document



OR Requires either one or both concepts to be present Connect synonyms with OR



NOT Excludes documents from an answer set containing the word after NOT



Wildcards Wildcards allow for more comprehensive retrieval and more precision Use in reference and substance name searches

Internal and right-hand truncation is available

* Replaces 0 to any number of characters E.g.: polymorph* | immunoglobulin*conjugate*

? Replaces 0 or 1 character
E.g.: benzonorbornen?

Terms masked with double quotes will be searched as a phrase, e.g.: "Programmed cell death protein"



Substance name and structure

Name searches

Search with one or more substance names or identifiers

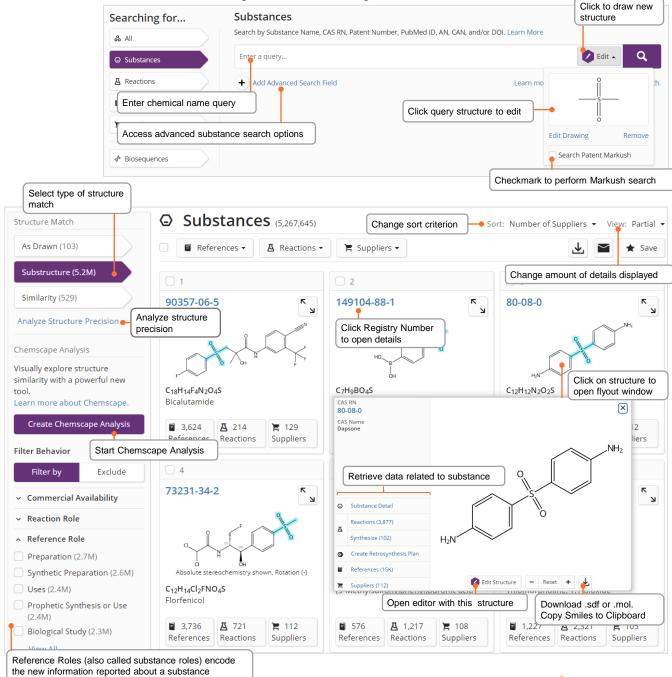
Streptomycin 57-92-1 Streptomycin sulfate "Streptomycin sulfate" Streptomycin Sulfoximin* WO2019234160

Finds Streptomycin record Finds Streptomycin record, uses CAS Registry number as identifier Finds 3 records: Streptomycin, Streptomycin sulfate and Sulfate

Finds 2 records: Streptomycin sulfate and Streptomycin Finds all names with the stem Sulfoximin Finds all indexed substances for this patent

Structure searches

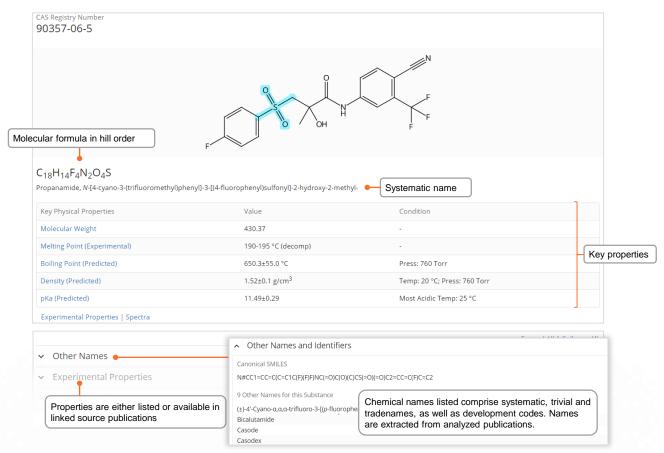
Substance searches returns results in an intuitive layout. The display highlights most relevant hits, critical property information and high-resolution images



Substance detail and structure editor

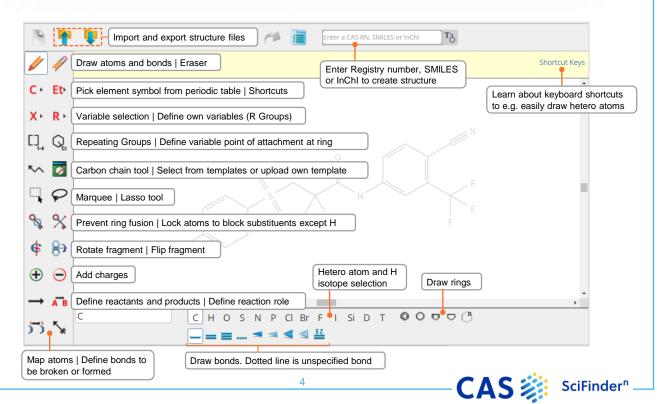
Substance detail

Click on the CAS Registry number to show substance details with structure, molecular formula, properties and further data



CAS Draw editor

Define structure and reaction queries with the structure editor



Advanced search query builder

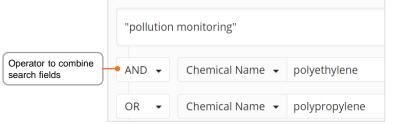
Advanced Search Query Builder

Provides specific reference and substance search fields from SciFinderⁿ's landing page

- · Operators are processed in this order: OR, AND, NOT
- · Operators are not allowed in a single adv. search field
- · Wildcards are allowed, e.g. peek*
- Up to 50 Advanced Search Fields (49 if also using the main search field)



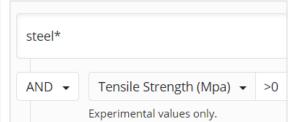
Examples Reference Search



Query interpretation:

"pollution monitoring" and (polyethylene or polypropylene)

Substance Search



Query interpretation:

Steel with tensile strength property information



Advanced Search Fields

The below advanced search fields are available

Reference

- Author Name
- Journal Name
- Organization Name
- Title
- Concepts
- Substances
- Publication Year
- Document Identifier
- · Patent Identifier

Substance

- · CAS Registry Number
- Chemical Name
- · Document Identifier
- Molecular Formula
- · Patent Identifier
- Experimental Spectra
- Biological
- Chemical Properties
- Density
- Electrical
- Lipinski
- Magnetic
- Mechanical
- Optical and Scattering
- · Structure Related
- Thermal



CAS Roles

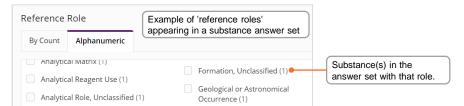
CAS Roles

Roles are linked to the substances and allow you to find focused publications connecting the substance of interest to its specific role within the scope of the publication.

- Super roles are broad categories and comprise all related specific roles. Examples are Analytical Study, Preparation or Occurrence
- Specific roles a more precise. They relate to aspects such as the use of the substance in an analytical study as an analyte (Analyte) or the occurrence of a compound in a plant (Natural Product Occurrence)

Roles in substance results

From a search on substance(s), the roles filter will indicate the type of roles that are connected to the substance(s) in the publications.

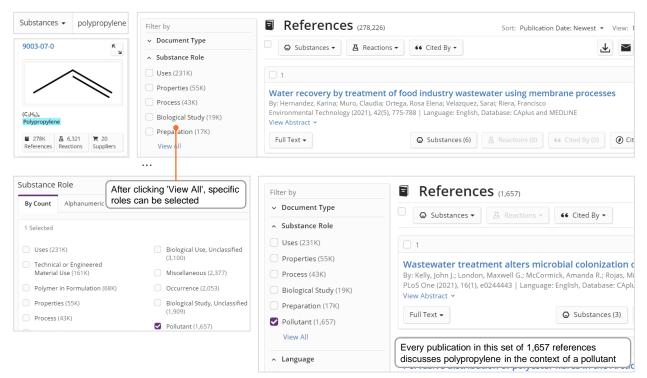


Roles in reference results

Roles will appear as a filter in reference result sets whenever you have retrieved hits in the substance indexing segment of the records, i.e. by retrieving substance names or performing a crossover after structure-based searches.

Example: I am interested in the subject of (marine) pollution, how can I find publications where polypropylene is specifically described as a pollutant?

The search for polypropylene retrieves a large number of references. The substance role window shows all roles that apply to Polypropylene in this answer set. The **Pollutant** role indicates there are 1648 publications that describe polypropylene as a pollutant.



Biosequence searching

Development

Sequence searching is developed in stages. June 2021 status:

- BLAST: Search similar sequences
- · CDR: Search antibodies via antigen binding sites
- Motif: Search conserved shorter sequence patterns
- Global crossover from sequences to patents

Implemented Implemented Implemented Implemented

· Crossover from sequences to scientific literature

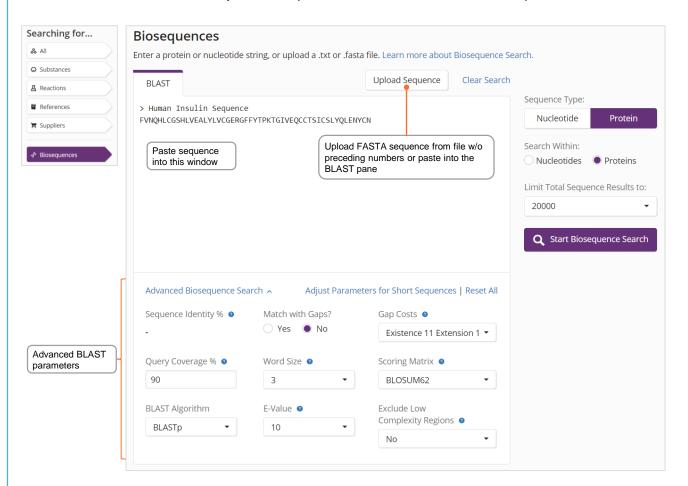
In development

BLAST similarity search

BLAST allows to search for similar nucleotide and amino acid sequences. Alignment results are shown in an intuitive graphical layout with easy-to-use precision filtering for identity and coverage percentages. Reference results are linked to the sequence hits.

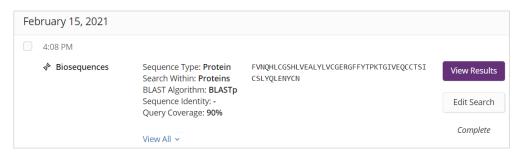
Perform a BLAST search

- Open the Biosequences module from the main SciFinderⁿ search page
- · Load sequence from file or paste sequence
- Supported formats: Sequences containing residues represented by singleletter codes, e.g. in the FASTA format. Leading numbers are not allowed.
- Sequence input may contain header line (starting with >). Sequences can be separated by (multiple) headers, thus allowing for batch processing
- Adjust BLAST parameters as desired and start biosequence search



BLAST result analysis

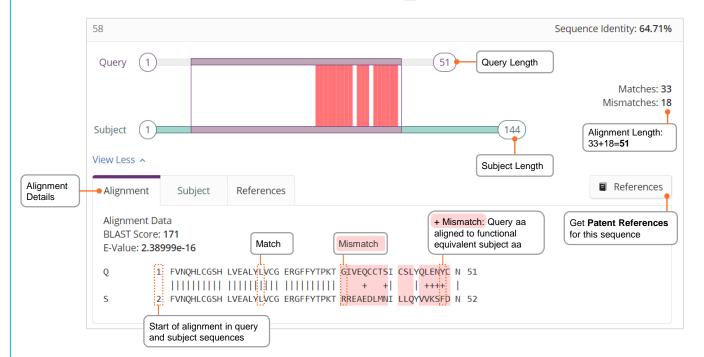
Access Results



View Results

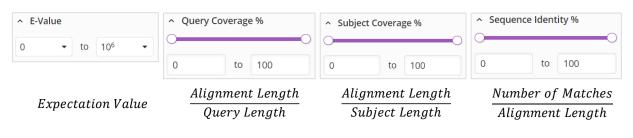
View BLAST sequence similarity results.

- · Alignments are sorted by Sequence Identity
- · Simplified graphical overview shows alignment quality
- · Mismatches are indicated by red lines
- · Detailed alignments can be viewed in 'Alignment' tab
- · Subject details and patent previews are available in separate tabs
- Click References to retrieve related patent references
- XLSX result download available ±



Filter Results

Filtering dynamically changes the result set

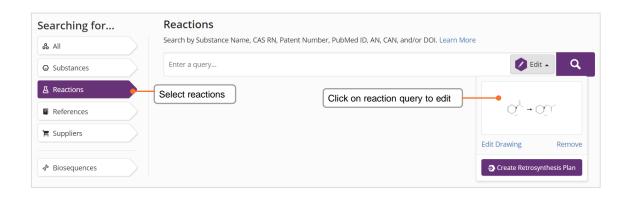


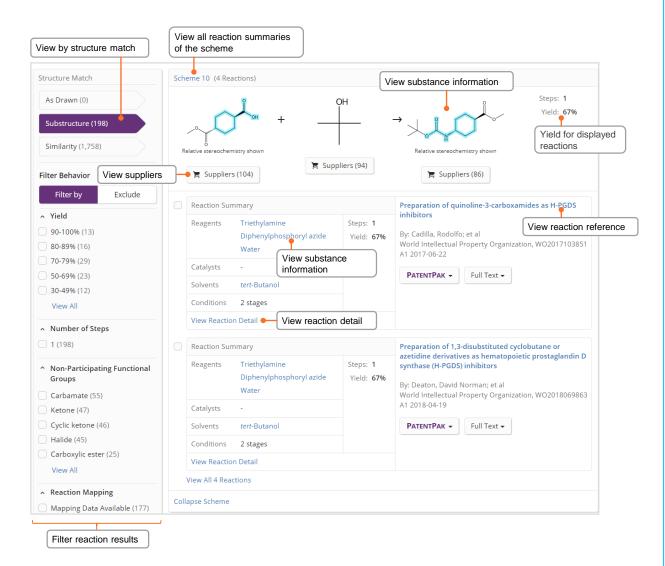
Reaction searching

Reaction searches

Reactions queries can be substance names, CAS Registry Numbers, document identifiers, or a chemical structure

- Reactions are grouped into schemes with identical reactants and products
- · Reactions are sorted by yield within a scheme
- Find reactions by substance name, registry number, document identifier or chemical structure

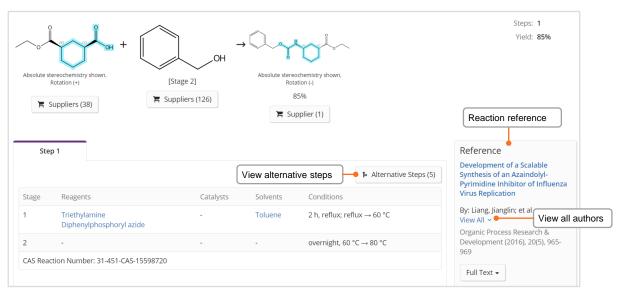


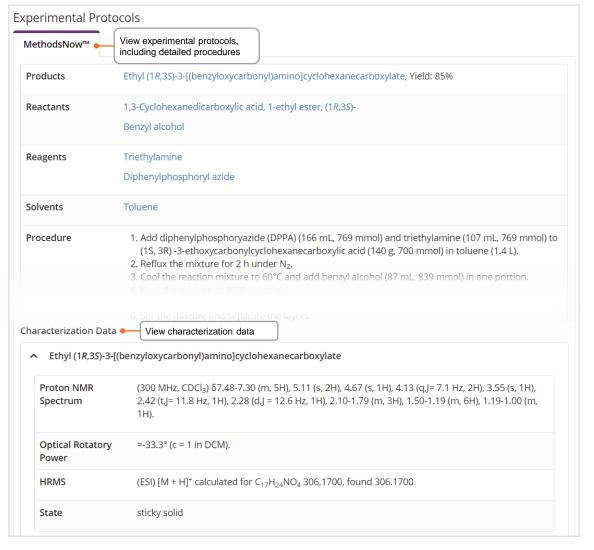


Reaction details

Reaction details

Details incl. solvents, catalysts, reagents, conditions and experimental protocols extracted from the publication and its supplement



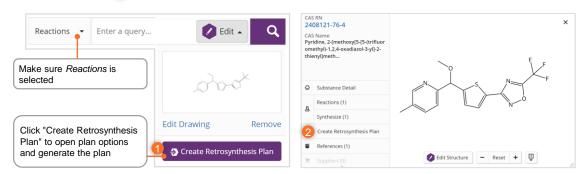


Retrosynthesis planner

Launch plan generation

There are two options to launch SciFindern's Retrosynthesis Planner

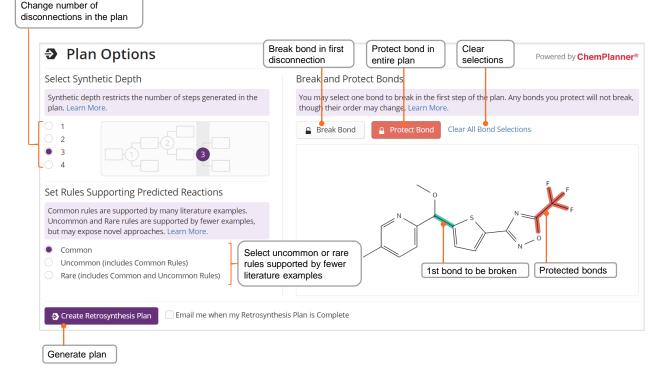
- Oraw reaction structure and create plan from Edit icon
- Open structure flyout window and start plan generation



Plan options

Edit plan options to...

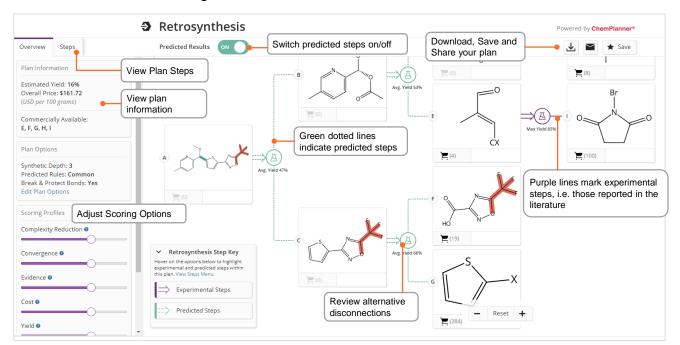
- · increase the synthetic depth
- · protect bonds through the entire synthetic route
- · define bonds to be broken in the first disconnection
- create a plan with more meaningful alternatives,
 e.g. for poly- or heterocyclic molecules



Retrosynthesis plan and alternative steps

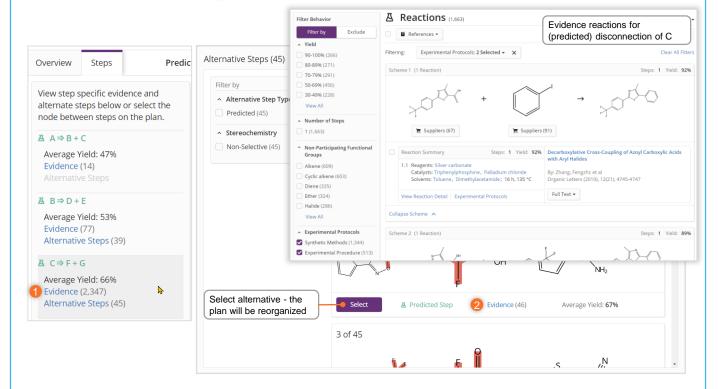
Open plan

The Experimental Plan is available within a few seconds, the calculation of the Predictive Retrosynthesis Plan will take a bit longer



Alternative steps Provide an overview of all experimental and predicted disconnections Evidence reactions are displayed as a reaction answer set

Access Evidence Reactions from the 1 link in the steps overview or 2 the alternative reaction scheme



Scoring Options

Scoring Options

For plans with **predicted** steps, you may increase or reduce the score assigned to steps and alternatives by each profile, which determines what is displayed in the plan/alternative steps.

- Each scoring profile may be set to Off (extreme left), Low, Medium, or High (extreme right)
- · The default setting for each profile is "Medium," as shown below
- Moving the slider all the way to the left turns that profile's scoring "Off," and it will
 not be a factor step selection or alternative ranking



Complexity Reduction

Reduces the complexity of a step's reactants compared to its product.

In retrosynthesis plans, you typically want high complexity reduction.

Convergence

Determines how "branched" the plan is; you typically want the plan to be as branched as possible (high convergence), rather than linear.

For a given step, the more precursors there are, and the closer their relative sizes are, the more it's considered convergent.

Increasing Convergence displays steps/alternatives with more reactants.

Evidence

Ranks plan steps/alternatives based on the number of evidence examples supporting the particular reaction type.

More evidence examples for a step means that the reaction type has more applications and is more versatile in terms of conditions and substrates, and hence predictions made based on it are probably more reliable.

Increasing Evidence displays steps/alternatives with more supporting examples.

Yield

Applies to the yield of each step in the plan, which contributes to the yield of the target molecule.

Increasing the Yield displays a higher yield target molecule and steps/alternatives.

Atom Efficiency

Reduces reactant parts not included in a plan step's product.

Increasing Atom Efficiency displays steps/alternatives with the least amount of reactant atoms that do not map to the product.

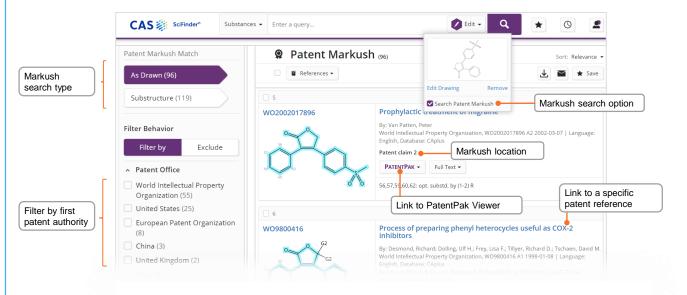
Clicking the Apply button redraws the retrosynthesis plan with the revised scoring profiles; clicking Reset Scoring restores the "Medium" default.



Markush Searching and PatentPak

Markush searching

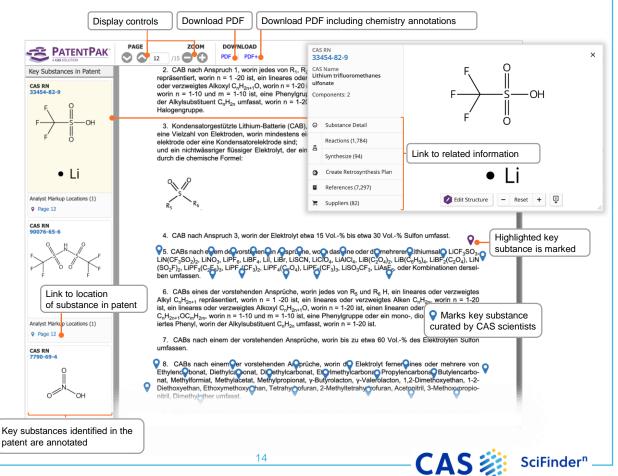
Markush structure searches can be performed by using the Search Patent Markush option while in Substances search mode



PatentPak

Up to three PatentPak Options for viewing a patent PDF:

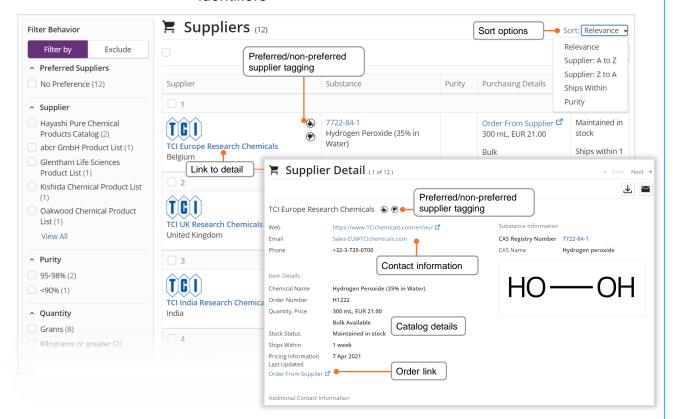
- PDF: Full-text patent PDF only; text-searchable PDF
- PDF+: Full-text patent PDF with marked-up Key Substances; text-searchable PDF
- Viewer: Patent PDF with linked markups of Key Substances; see below:



Supplier Searching and ChemDoodle®

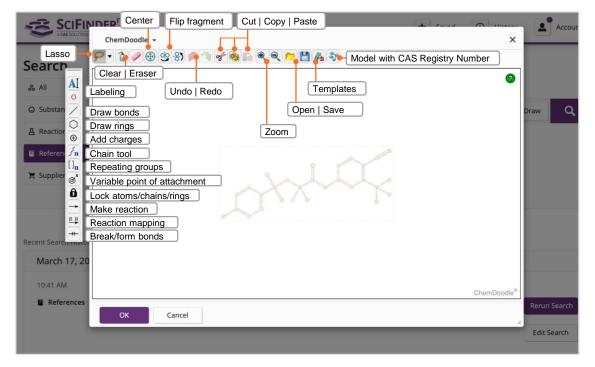
Suppliers searching

Suppliers searching allows for direct access to chemical catalog information based on chemical structure, names or other identifiers



ChemDoodle®

ChemDoodle structure editor is available in addition to the standard CASdraw editor. ChemDoodle is useful for tablets and mobile devices.



Login, Feedback and Support

Login Details

- · Login at http://scifinder-n.cas.org
- Use your existing SciFinder username and password

Feedback Button

Provide direct feedback to CAS



Learn More

SciFinderⁿ Training Topics:

https://www.cas.org/support/training/scifinder-n

Upcoming and recorded SciFinderⁿ webinars:

https://www.cas.org/about/events/scifinder-webinars

Contact Customer Support

Email help@cas.org to speak with a CAS Customer Center representative

