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As of March 2021, SciFinderⁿ

patents by normal or Markush structure, and also by reaction. SciFinderⁿ has powerful search functions to find scientific and engineering patents by Keyword and Boolean searches.

More about sequence searching:

SciFinderⁿ contains more than 500 million nucleotide and protein sequences, and allows you to search via BLAST, Motif or CDR (Complimentary Determining Region) search techniques. You will find alignments with misalignments highlighted and you can easily retrieve the relevant journal and patent references to you sequence query. The visualization platform BioScape helps to enable analyzing results.

If you need further assistance, contact a librarian:

Phone 0800 MASSEY (0800 627 739) and ask for the Library

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