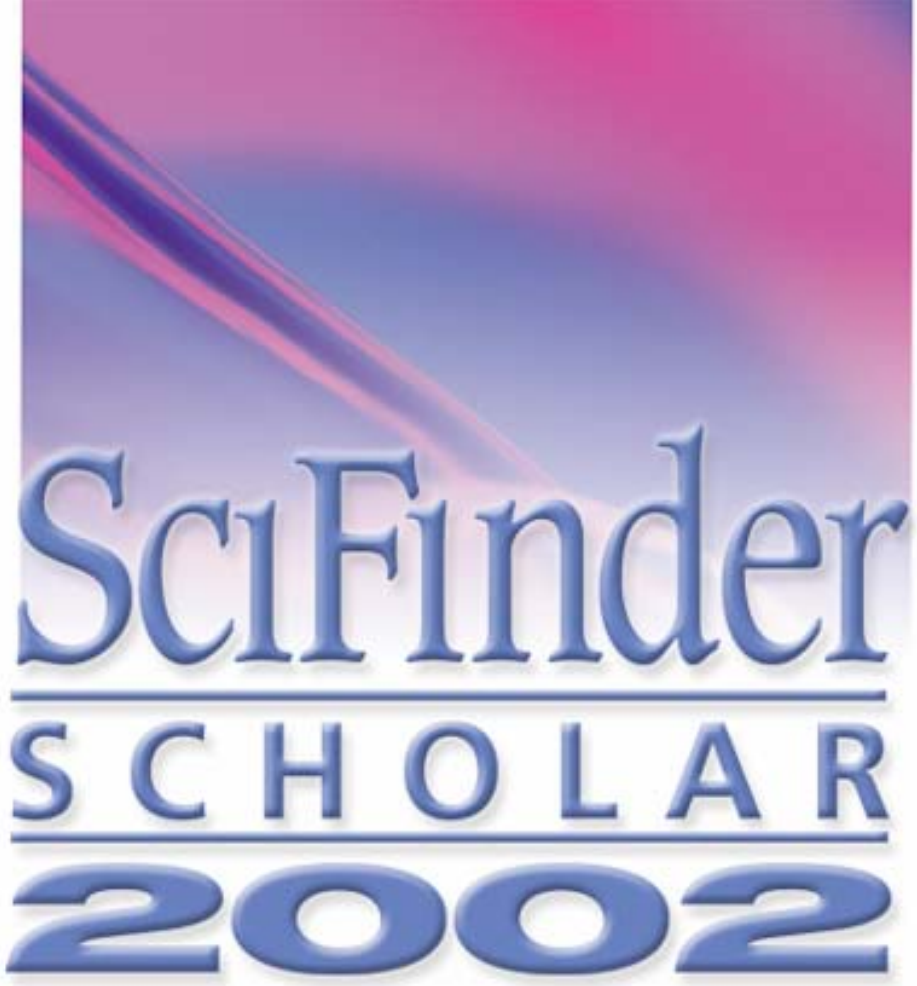

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Pressemitteilung: Der Nobelpr

9 Oktober 2002

Die Königlich Schwedische Akademie
beschlossen, den Nobelpreis des Jahr

„für die Entwicklung von Methoden zu
Strukturanalyse von biologischen Mak

zur einen Hälfte an

John B. Fenn
Virginia Commonwealth University, R

und

Koichi Tanaka
Shimadzu Corp., Kyoto, Japan

„für ihre Entwicklung von weichen Des
Methoden zur massenspektrometrisch
biologischen Makromolekülen“

und zur anderen Hälfte des Preises ar

Kurt Wüthrich
Eidgenössische Technische Hochschul
und The Scripps Research Institute, L

„für seine Entwicklung der kernmagne
Resonanzspektroskopie zur Bestimmu
dreidimensionalen Struktur von biolog
in Lösung“.

Massenspektrometrie ist eine sehr wichtige Analysenmethode – sie wird im allgemeinen in jedem Chemielaboratorium angewendet. Früher konnten nur kleinere Moleküle identifiziert werden, aber **John B. Fenn** und **Koichi Tanaka** haben Methoden entwickelt, die auch die Analyse biologischer Makromoleküle ermöglichen.

John B. Fenn veröffentlichte 1988 die sogenannte *ESI (electrospray ionisation)* Methode. Mit ihr werden zuerst kleine, geladene Tropfen einer Proteinlösung produziert, welche dann wegen des verdunstenden Wasseranteils zusammenschrumpfen. Langsam bleiben frei schwebende Proteinionen übrig, deren Massen man zum Beispiel dadurch ausrechnen kann, dass man sie beschleunigt und die Flugzeit über eine bekannte Strecke misst. Gleichzeitig führte Koichi Tanaka eine andere Technik ein, um Proteine frei schweben zu lassen, nämlich mittels *weicher Laserdesorption (soft laser desorption)*. Ein Laserpuls muss die Probe treffen, die in kleine Teile „gesprengt“ wird und somit die Moleküle freigibt.

Der zweite Teil des Preises zeichnet die Weiterentwicklung einer anderen Lieblingsmethode unter Chemikern, nämlich der *kernmagnetischen Resonanz, NMR,* aus. Mit NMR erhält man Information über die dreidimensionale Struktur und die Beweglichkeit des Moleküls. Durch seine Arbeit zu Beginn der 1980er Jahre hat **Kurt Wüthrich** die Anwendung von NMR auf Proteine ermöglicht. Er entwickelte zum einen eine generelle Methode zur systematischen Bestimmung gewisser Fixpunkte im Proteinmolekül, zum anderen ein Prinzip, um mittels der Abstände die dreidimensionale Struktur auszurechnen. Der Vorteil der NMR ist, dass man die Proteine in Lösung, also in einer den Verhältnissen in den Zellen gleichenden Umgebung studieren kann.

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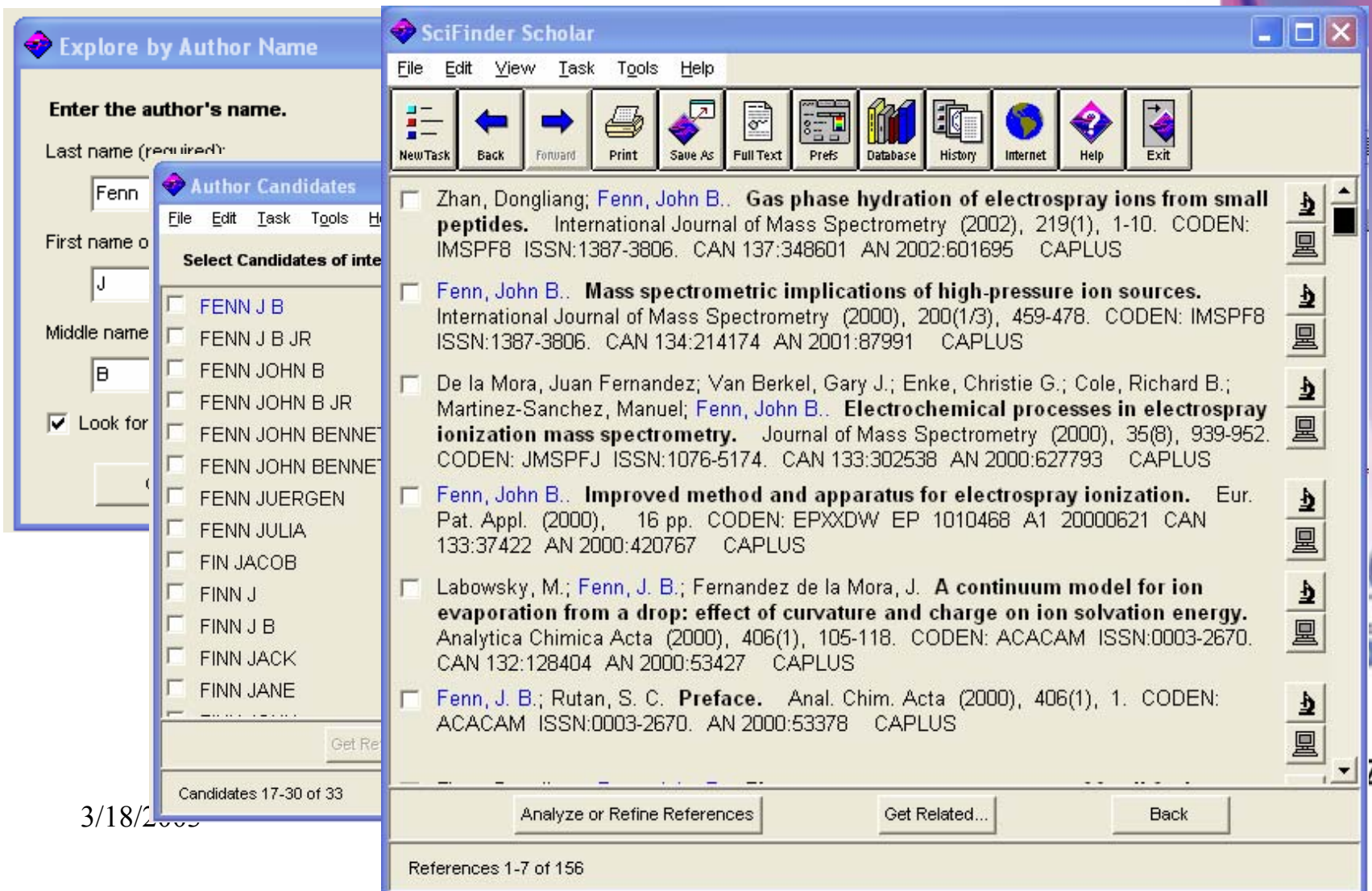

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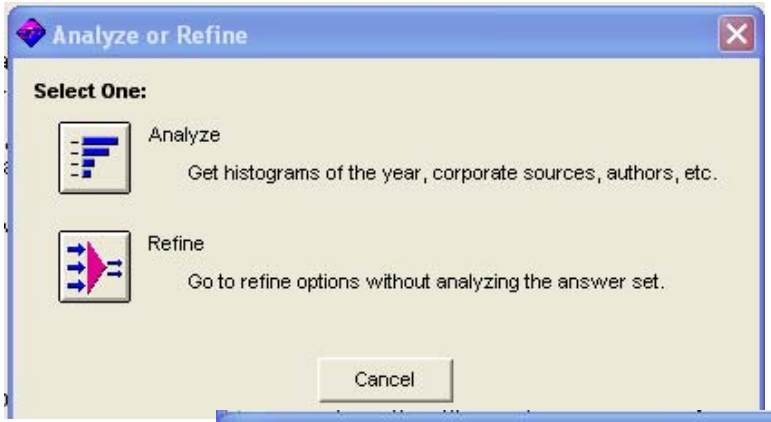

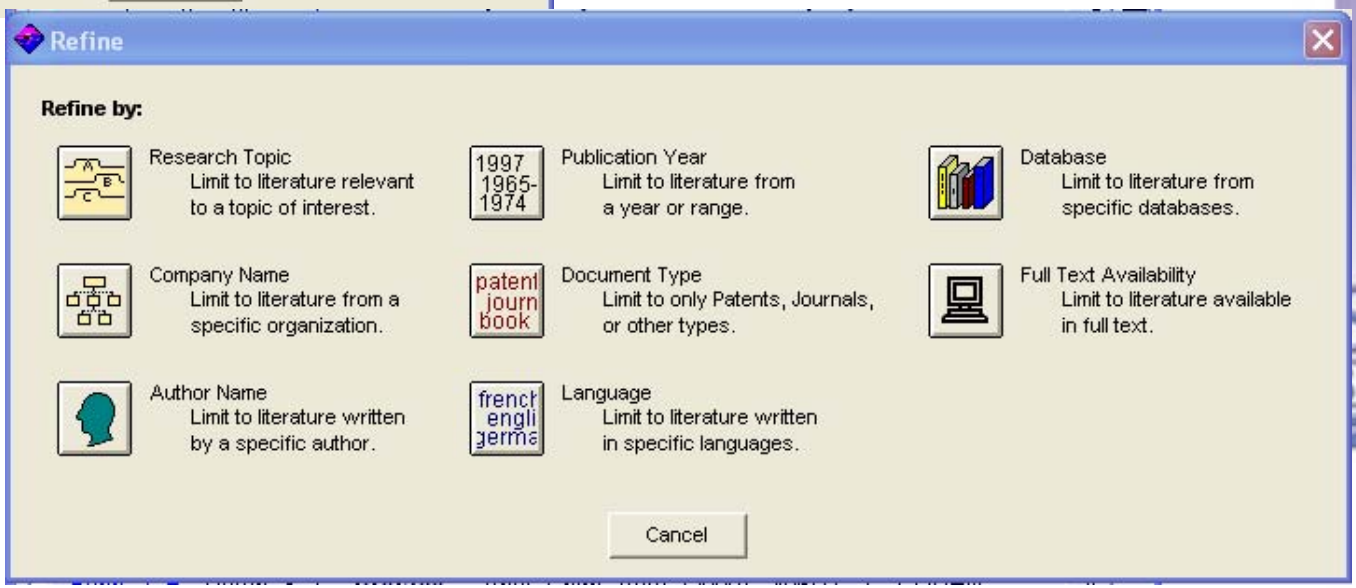

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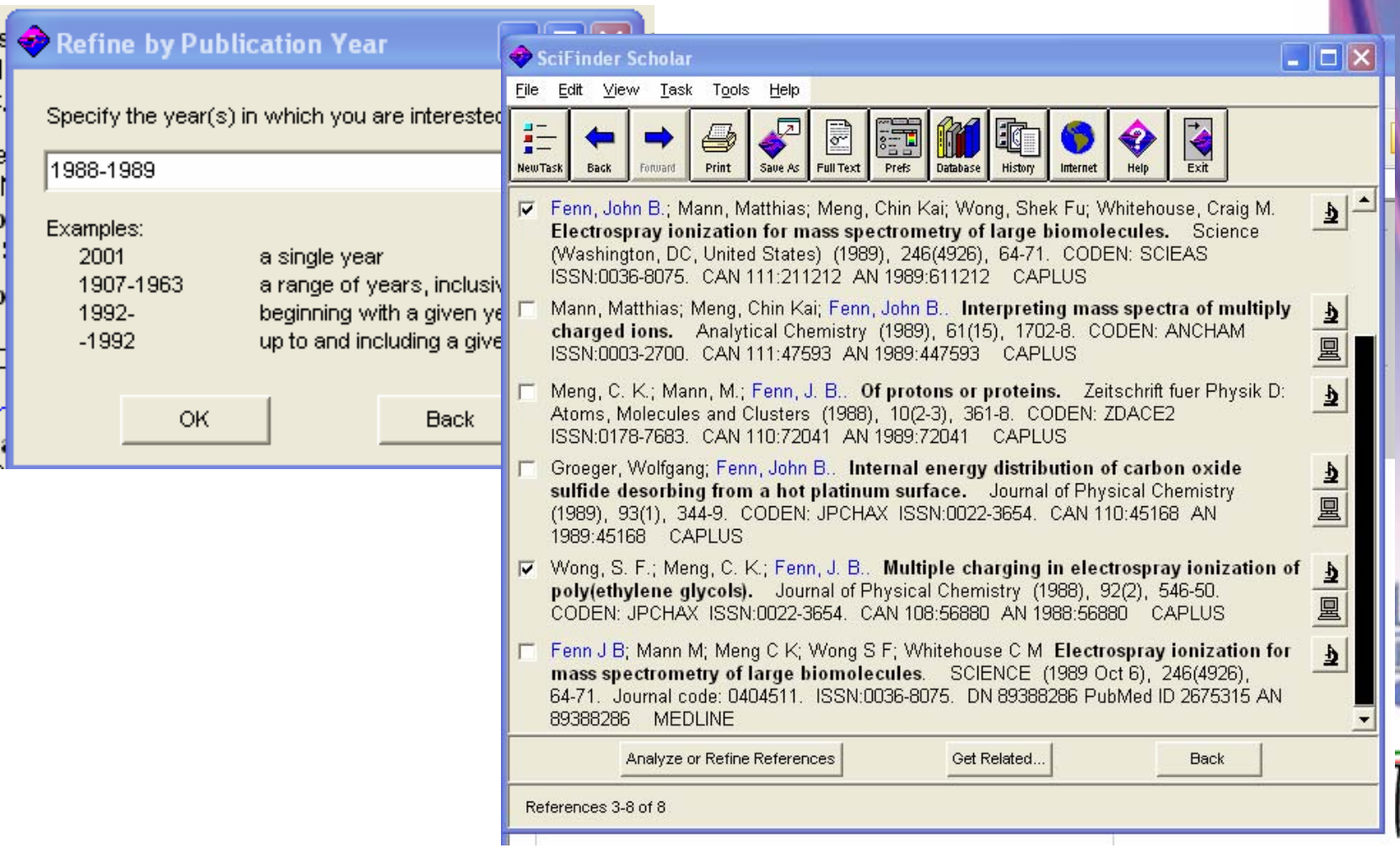

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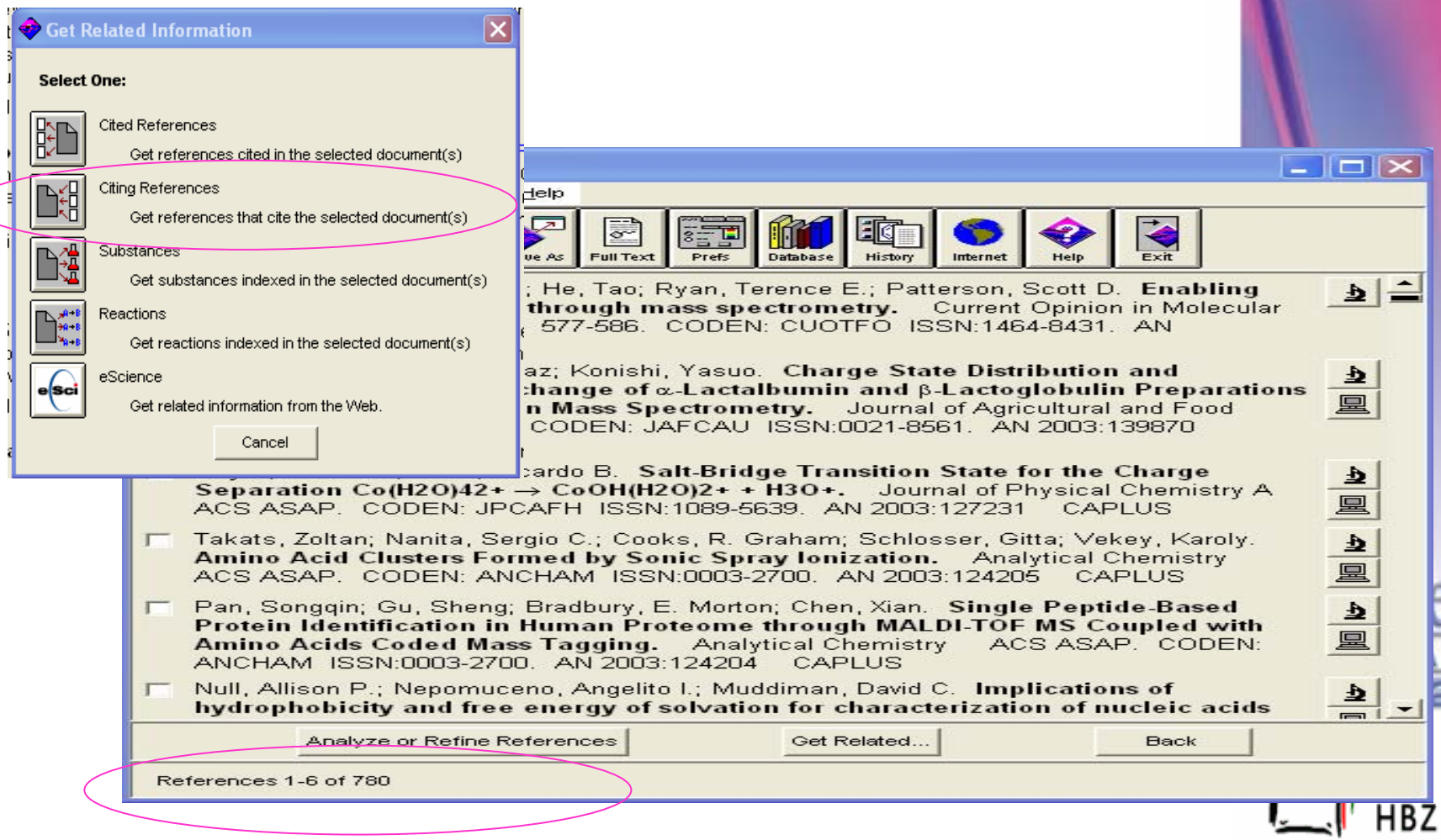

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121 Journal Articles, 21 Reviews, 7 Conferences

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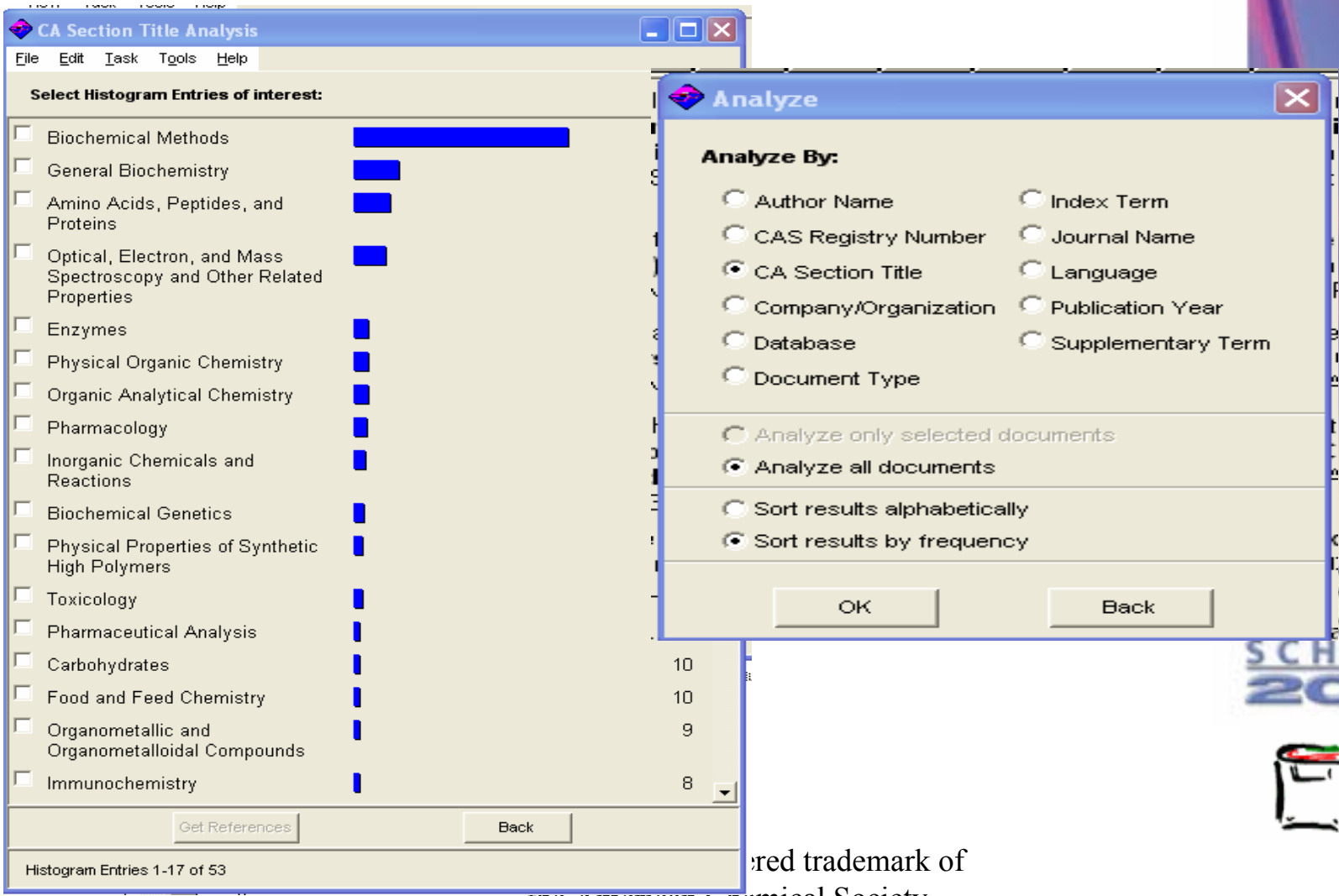

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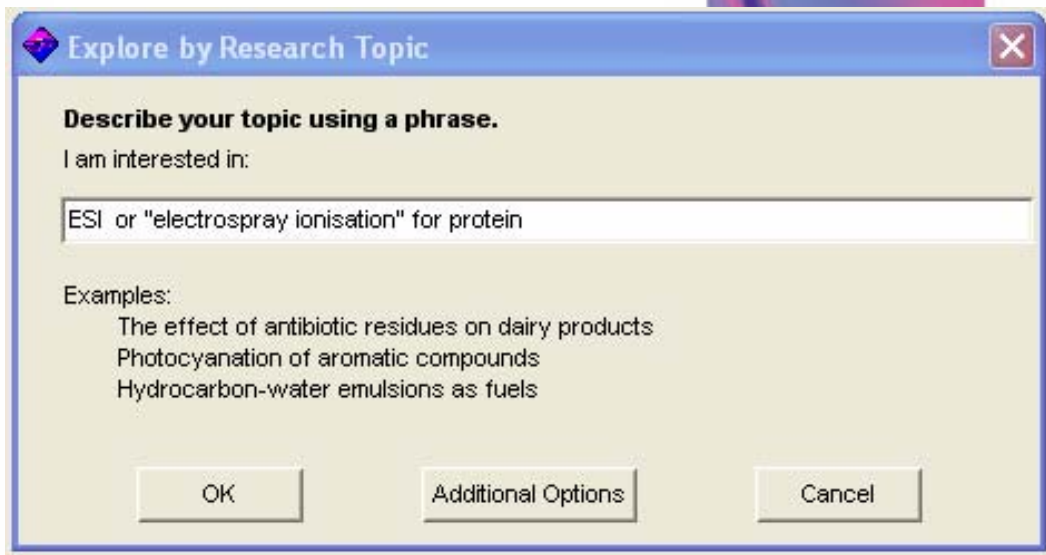

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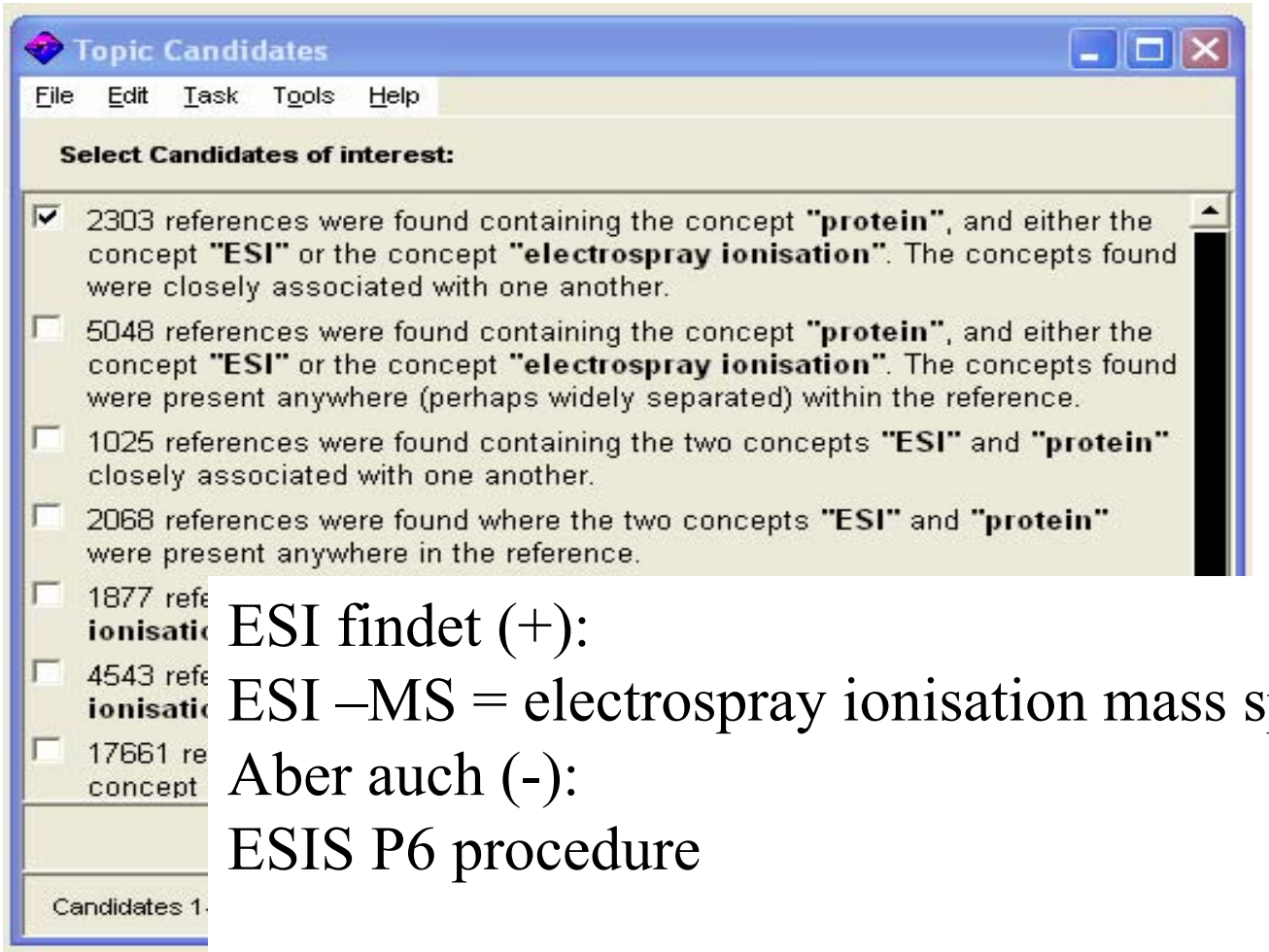

ESI findet (+):
ESI –MS = electrospray ionisation mass spectrometry
Aber auch (-):
ESIS P6 procedure

=> Abkürzungen stets überprüfen!

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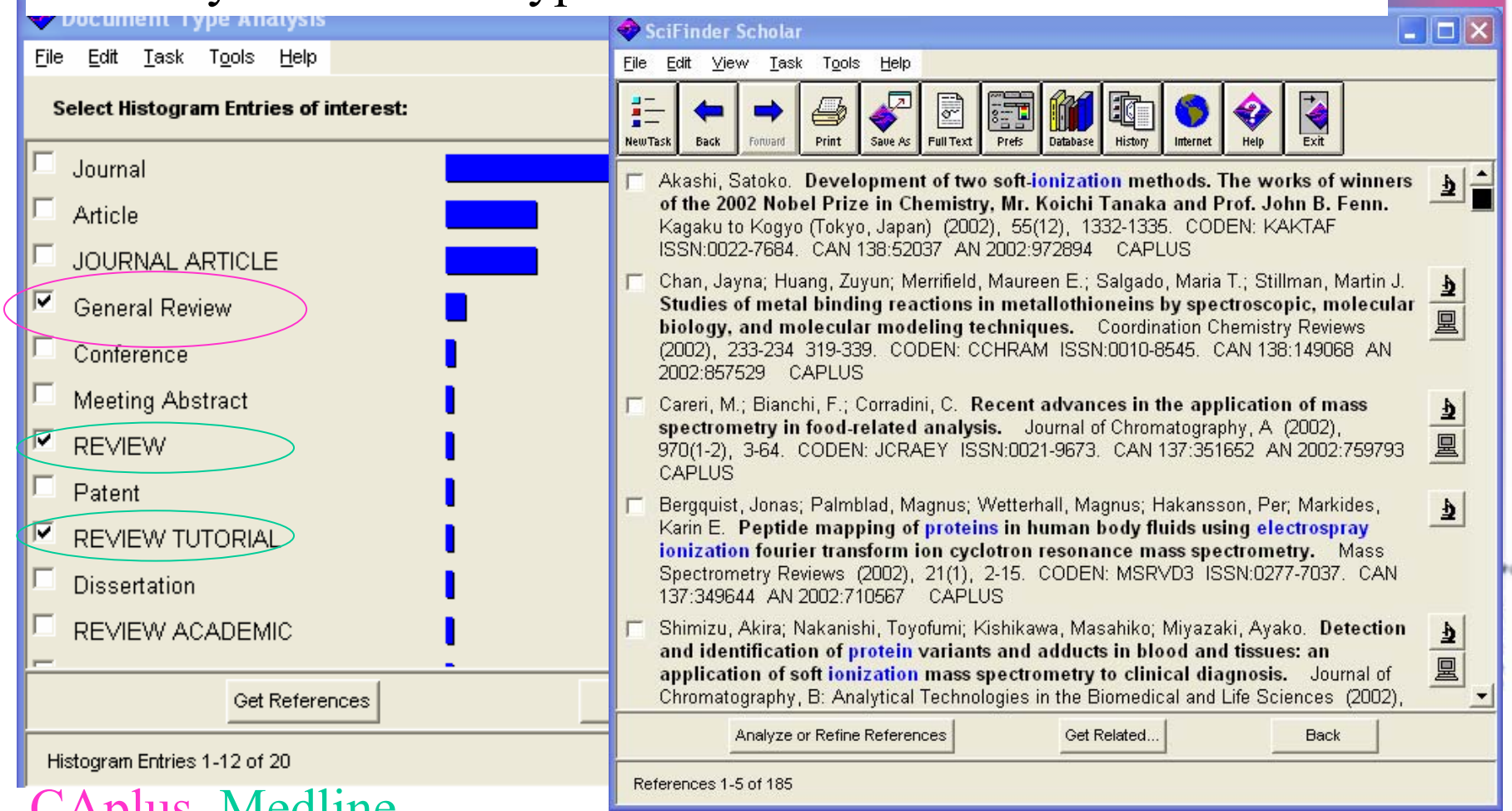

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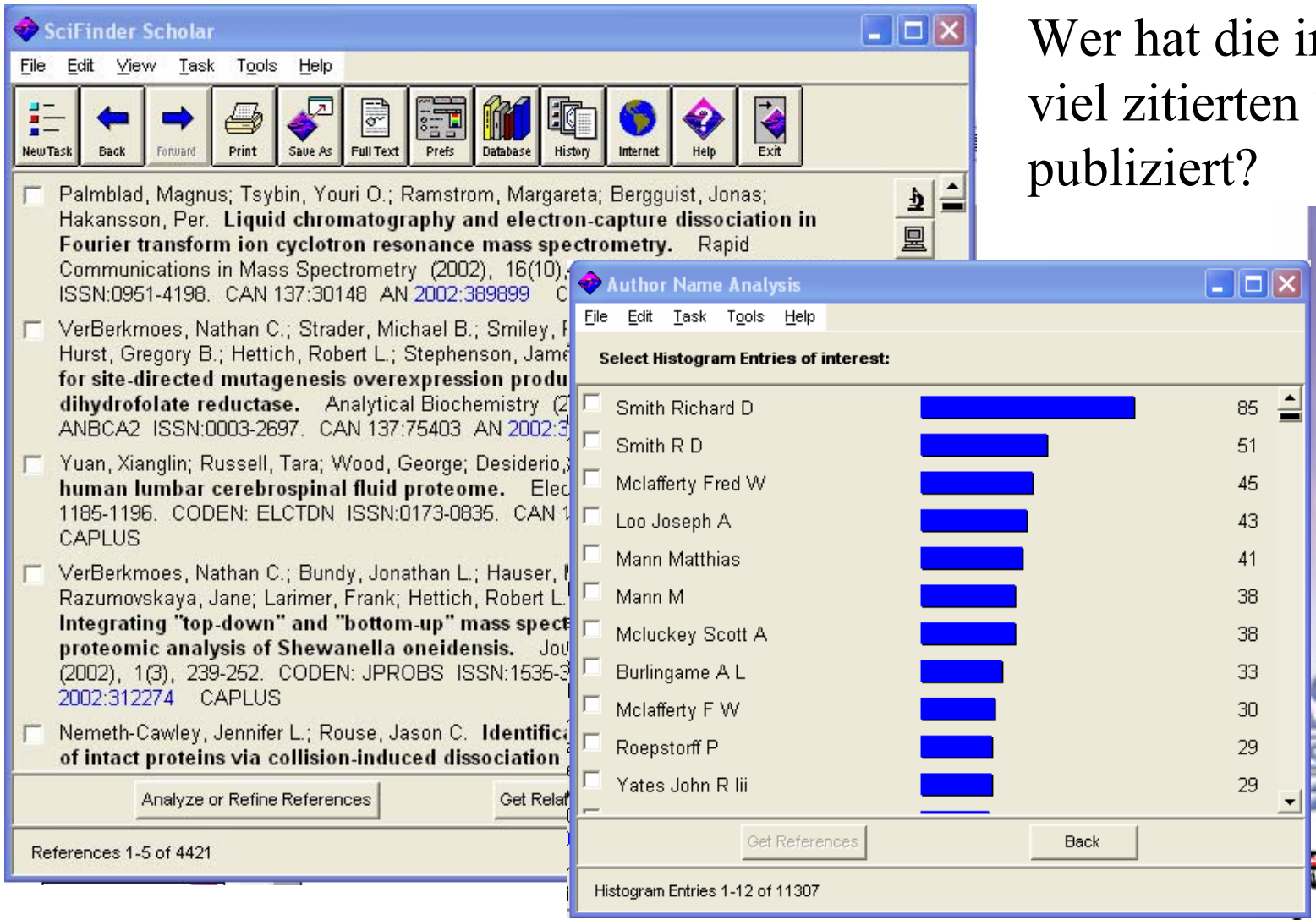

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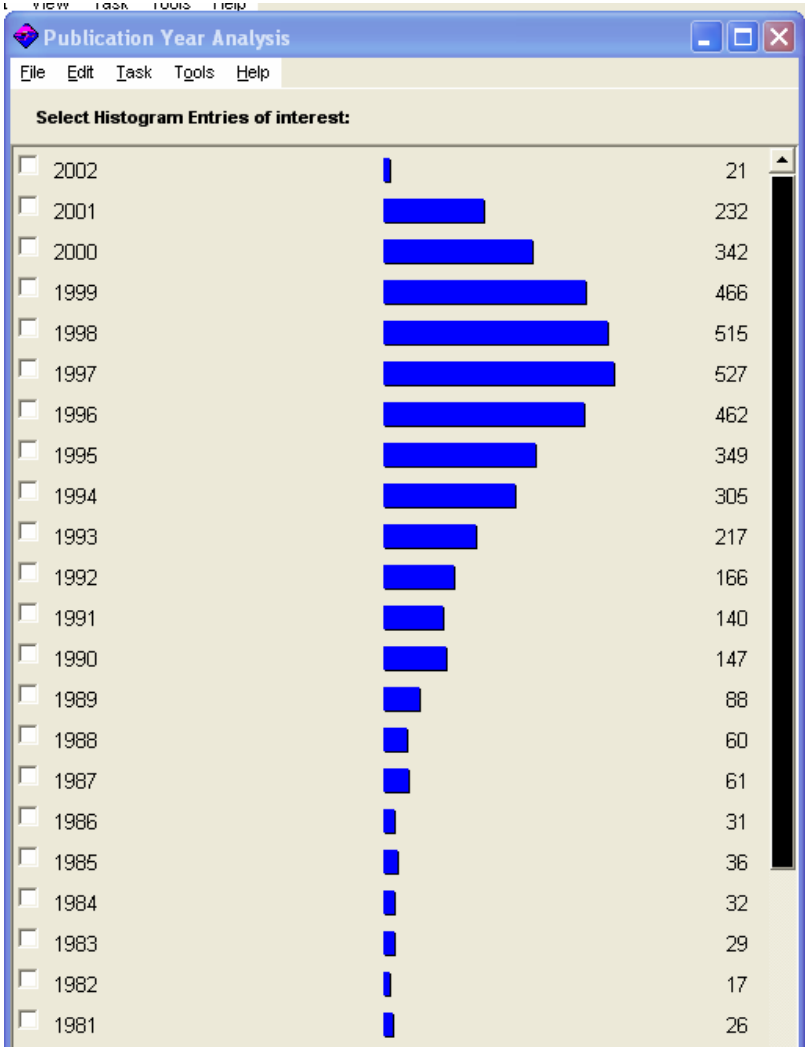

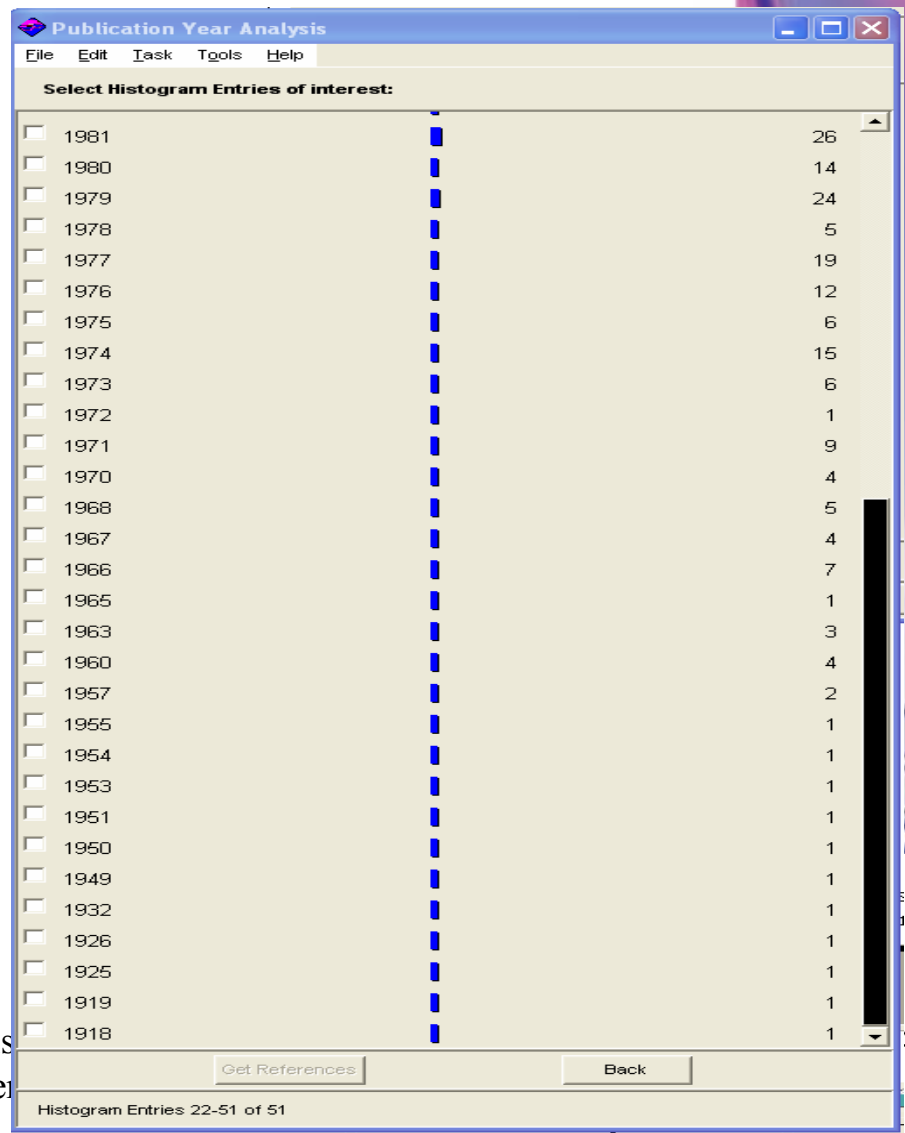

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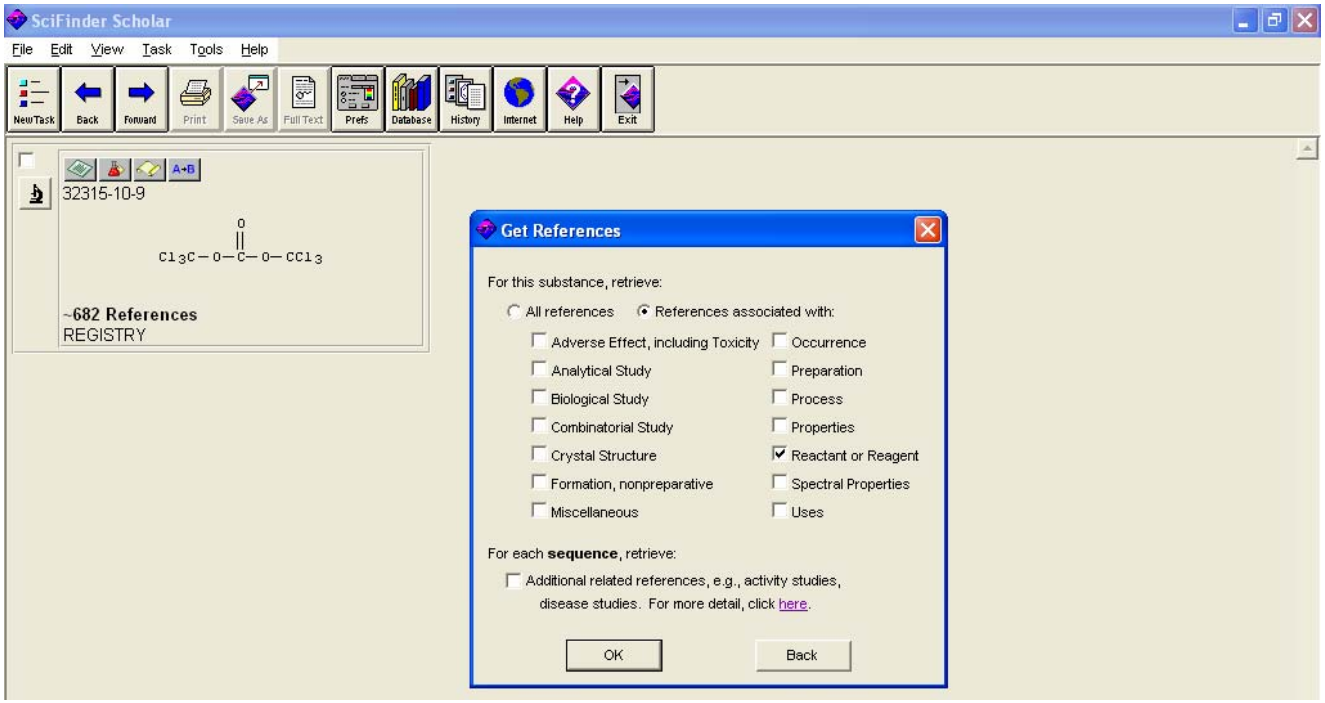

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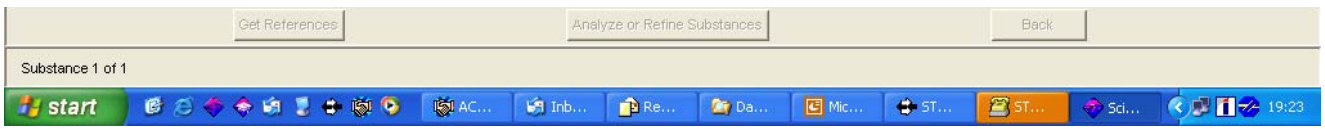
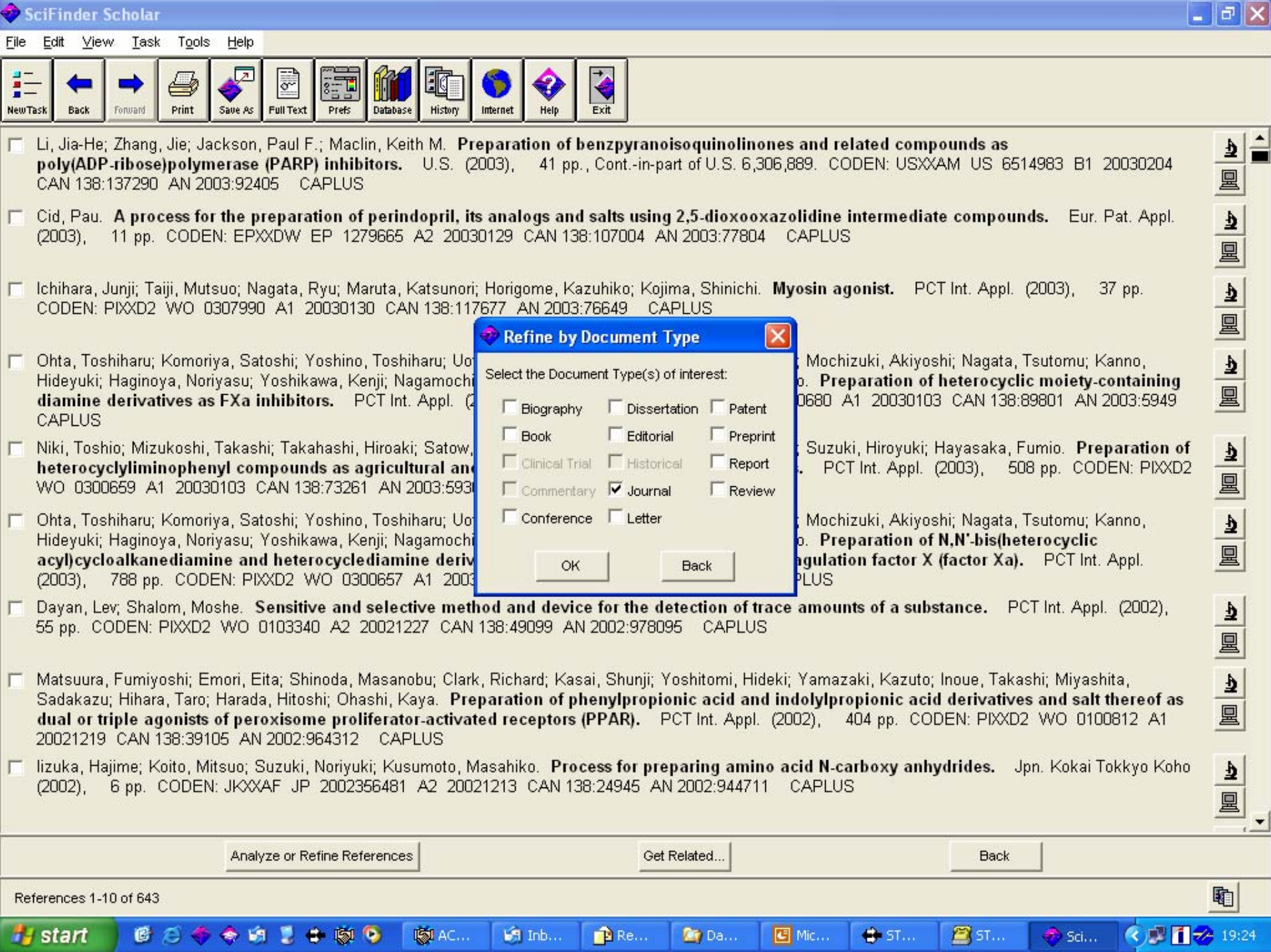
als Phosgen - Ersatzstoff für meine Synthesen von Heterozyklen

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Ohta, Toshiharu; Komoriya, Satoshi; Yoshino, Toshiharu; Uo Mochizuki, Akiyoshi; Nagata, Tsutomu; Kanno, Hideyuki; Haginoya, Noriyasu; Yoshikawa, Kenji; Nagamochi o. Preparation of N,N'-bis(heterocyclic acyl)cycloalkanediamine and heterocyclediamine deriv gulation factor X (factor Xa). PCT Int. Appl. (2003), 788 pp. CODEN: PIXXD2 WO 0300657 A1 2003 LUS
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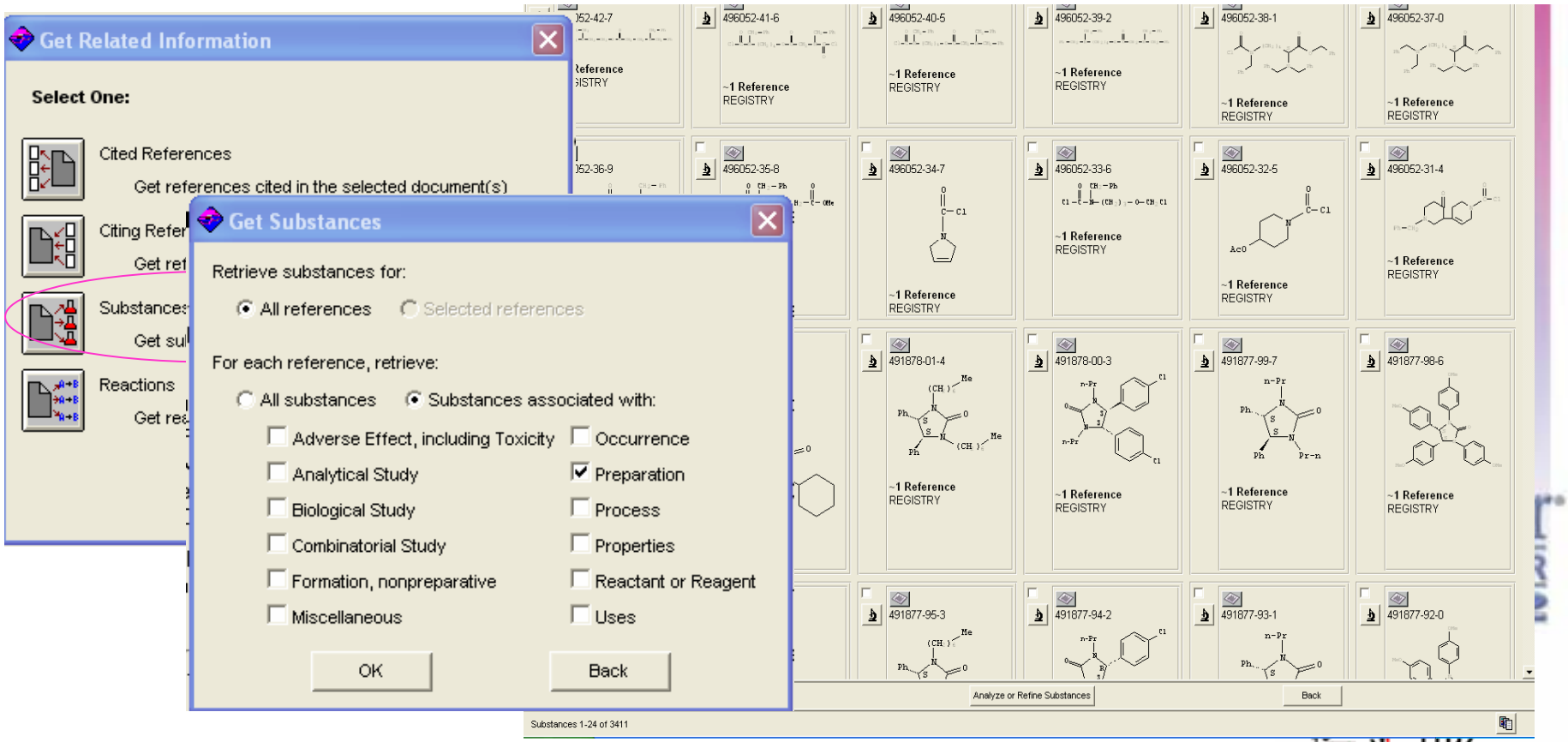

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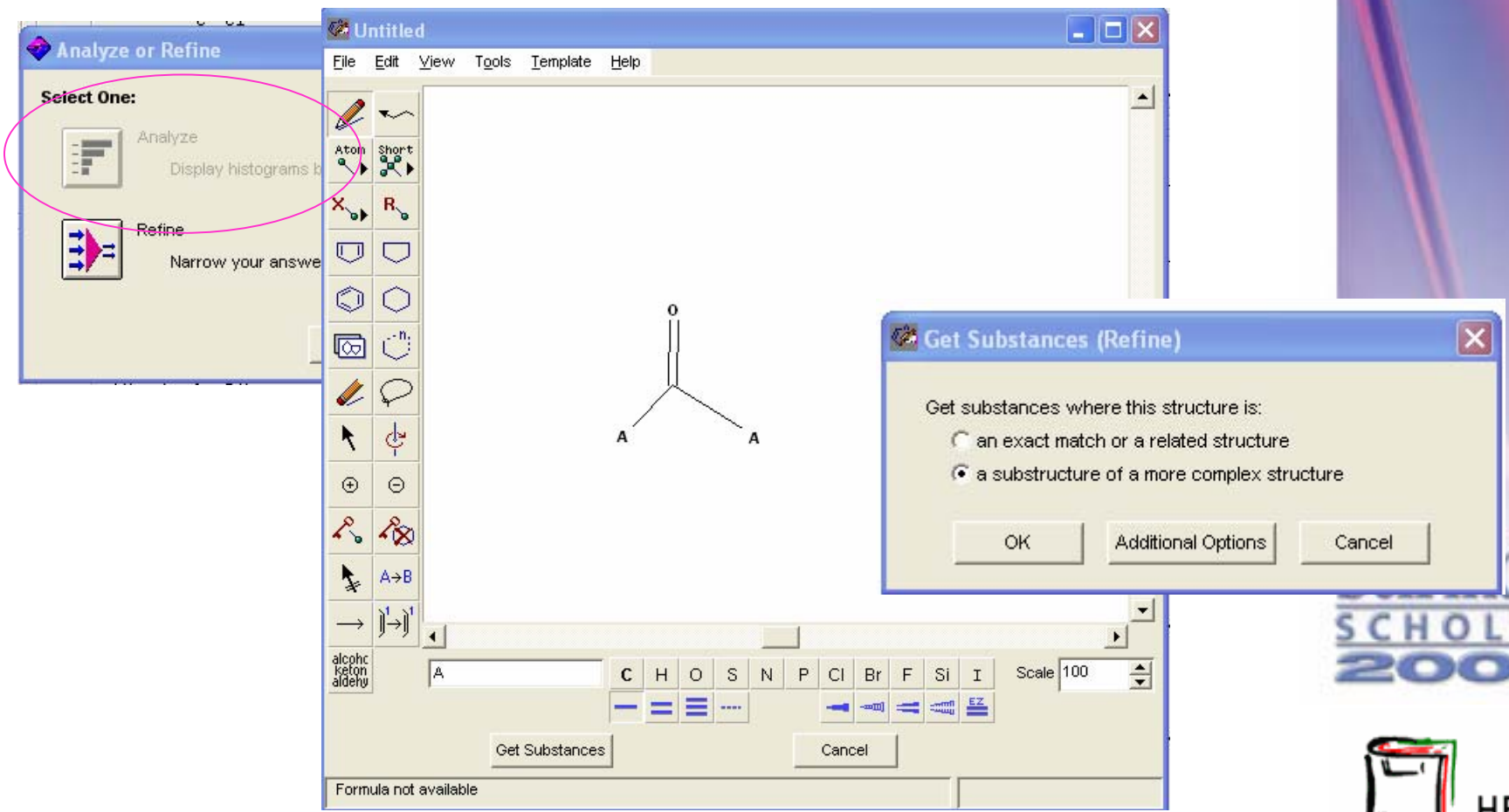

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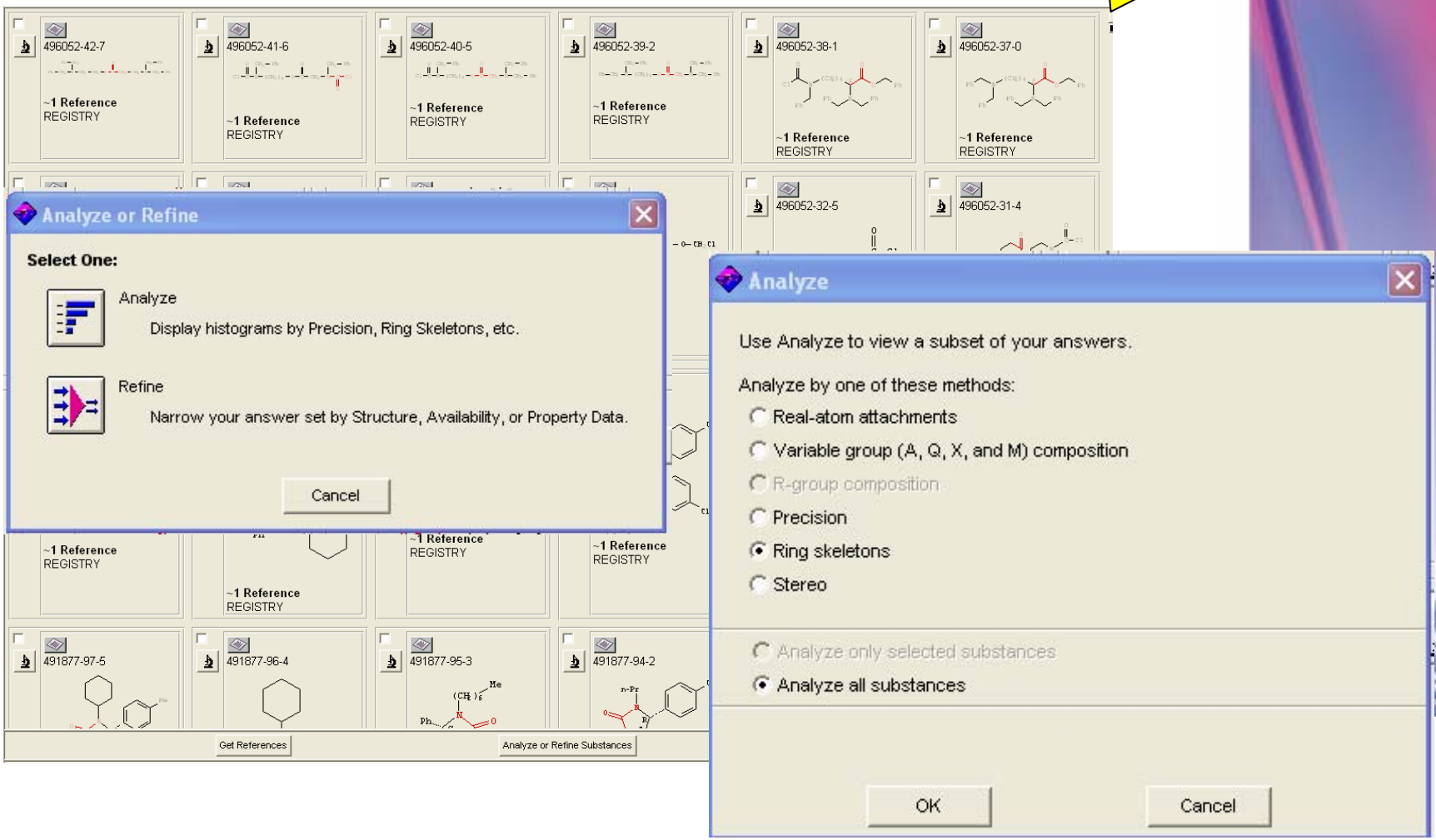

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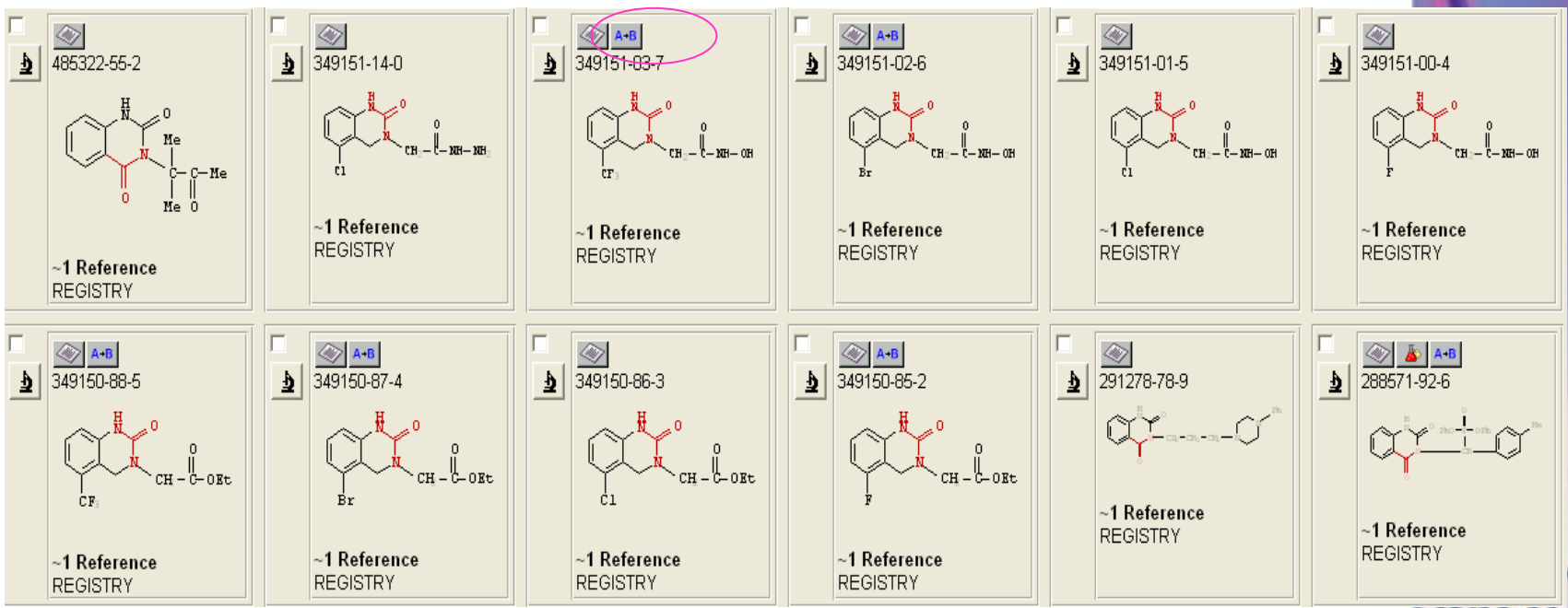

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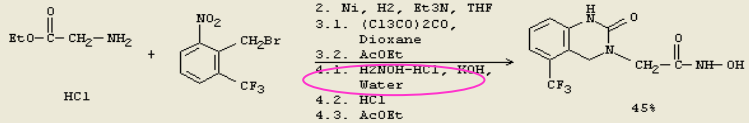

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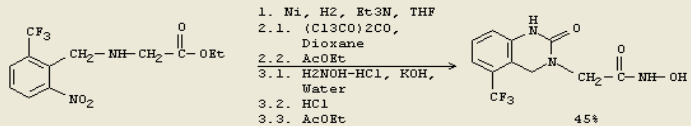

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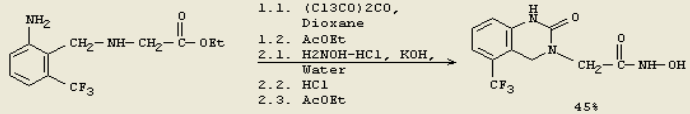

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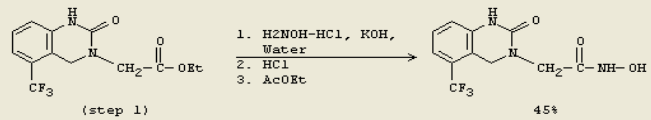

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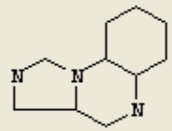

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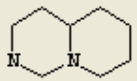

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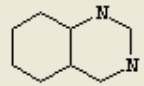

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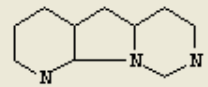

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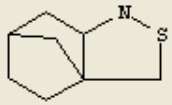

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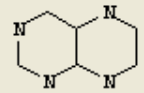

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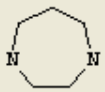

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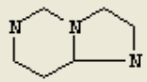

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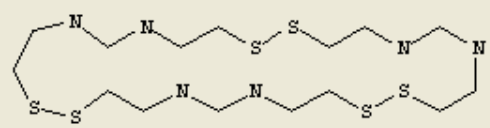

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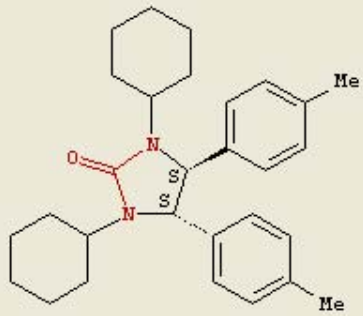

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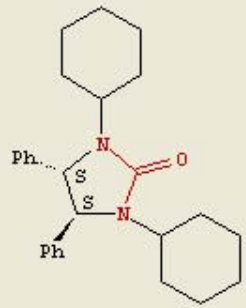

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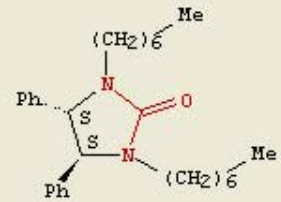

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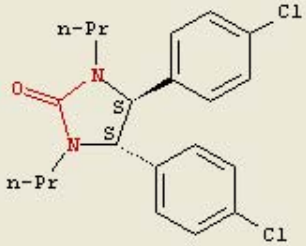

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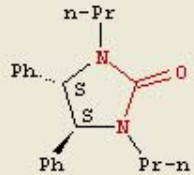

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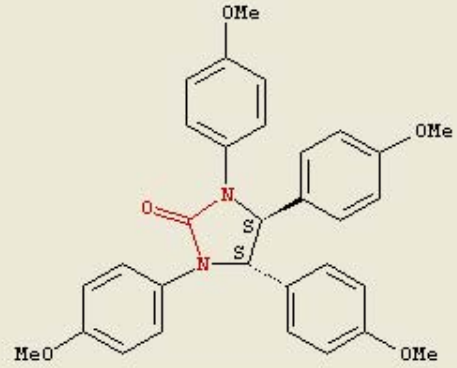

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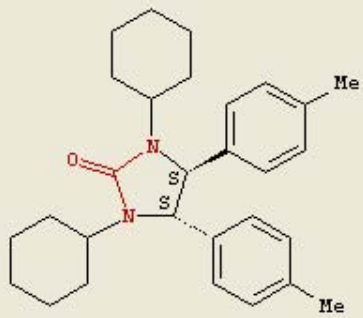

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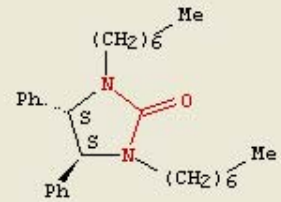

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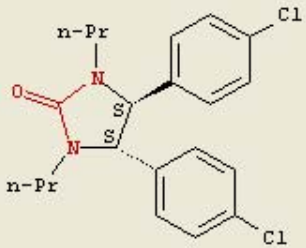

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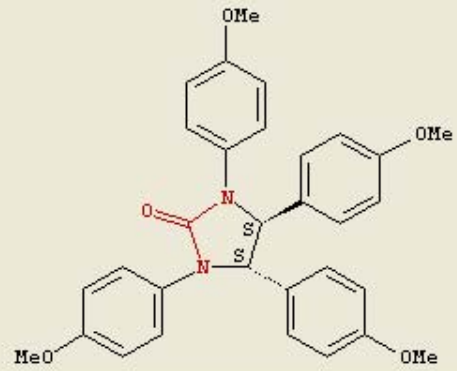

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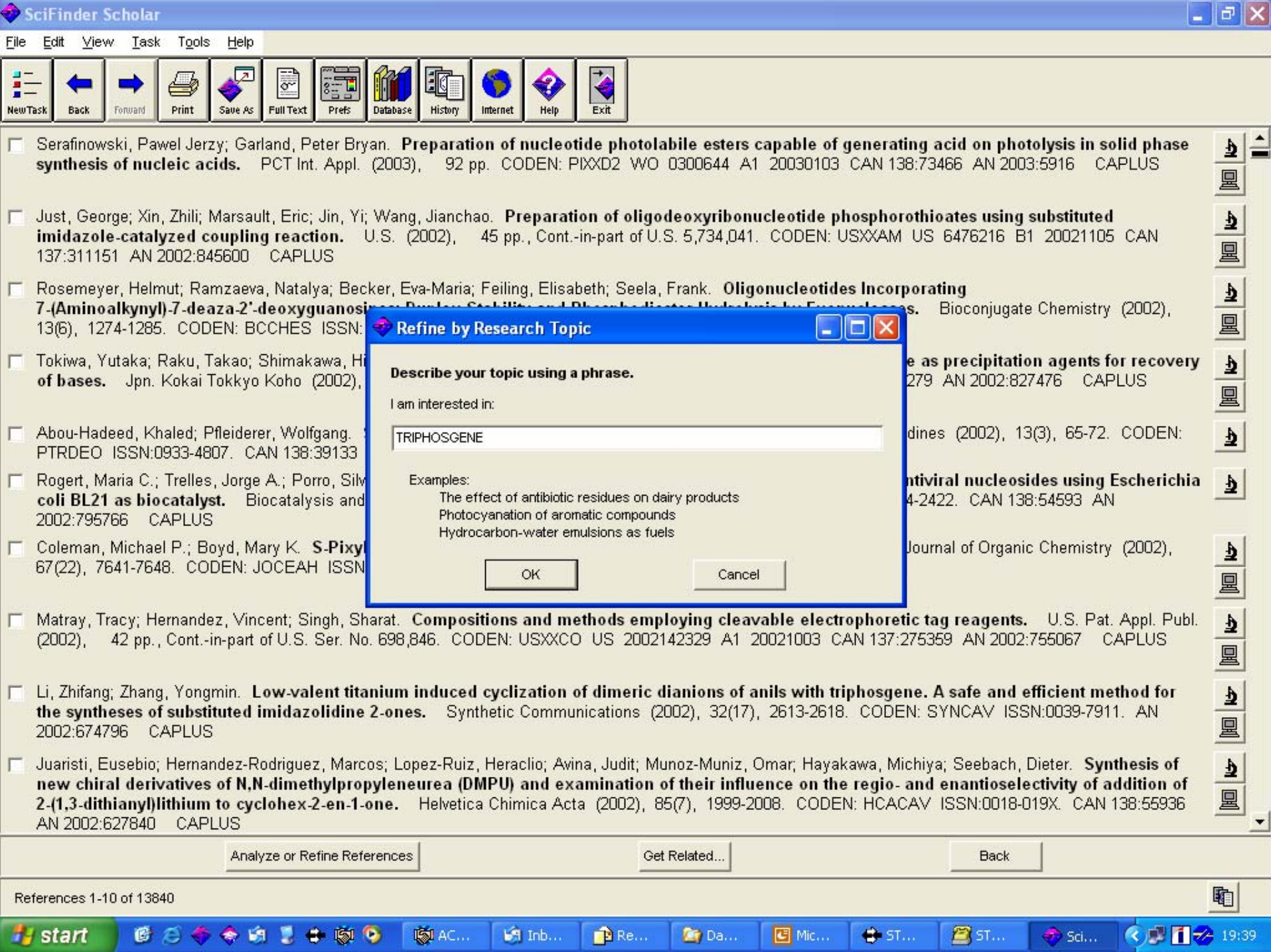
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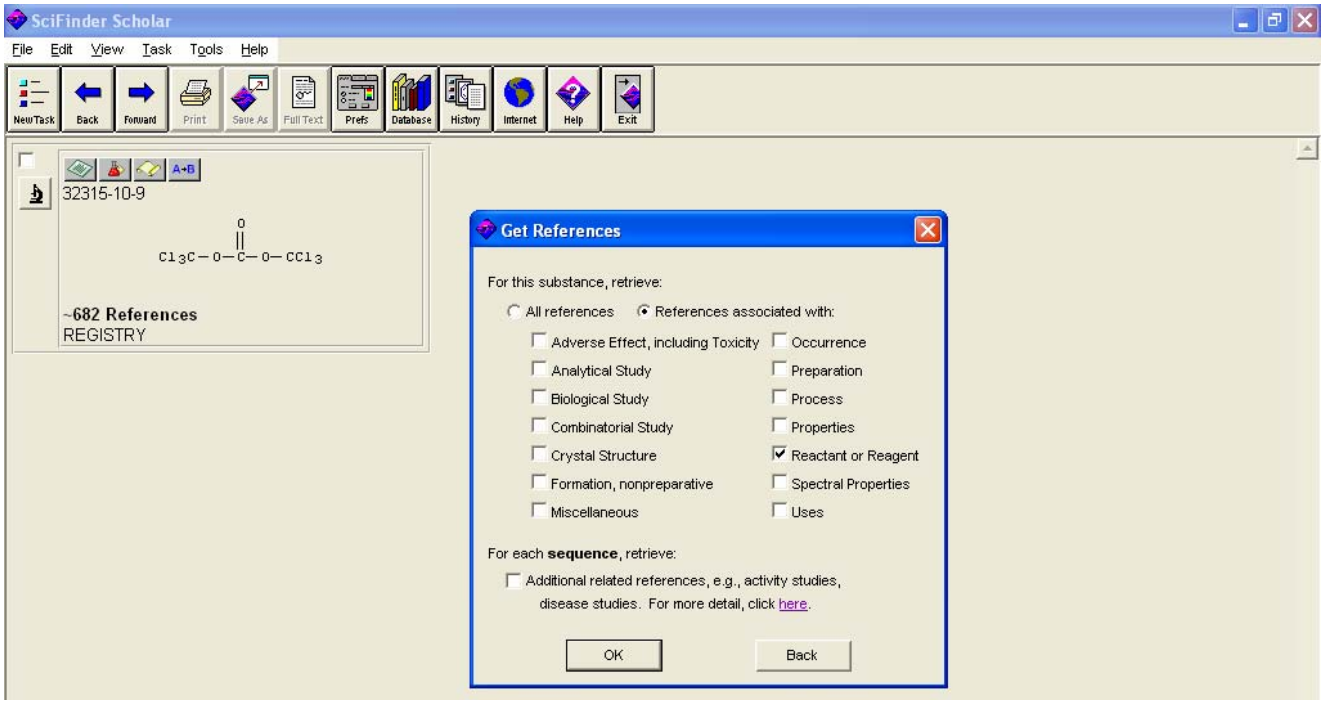

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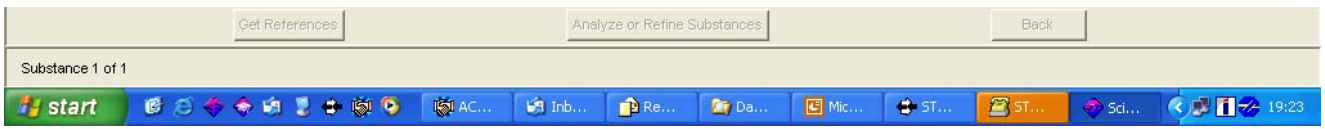

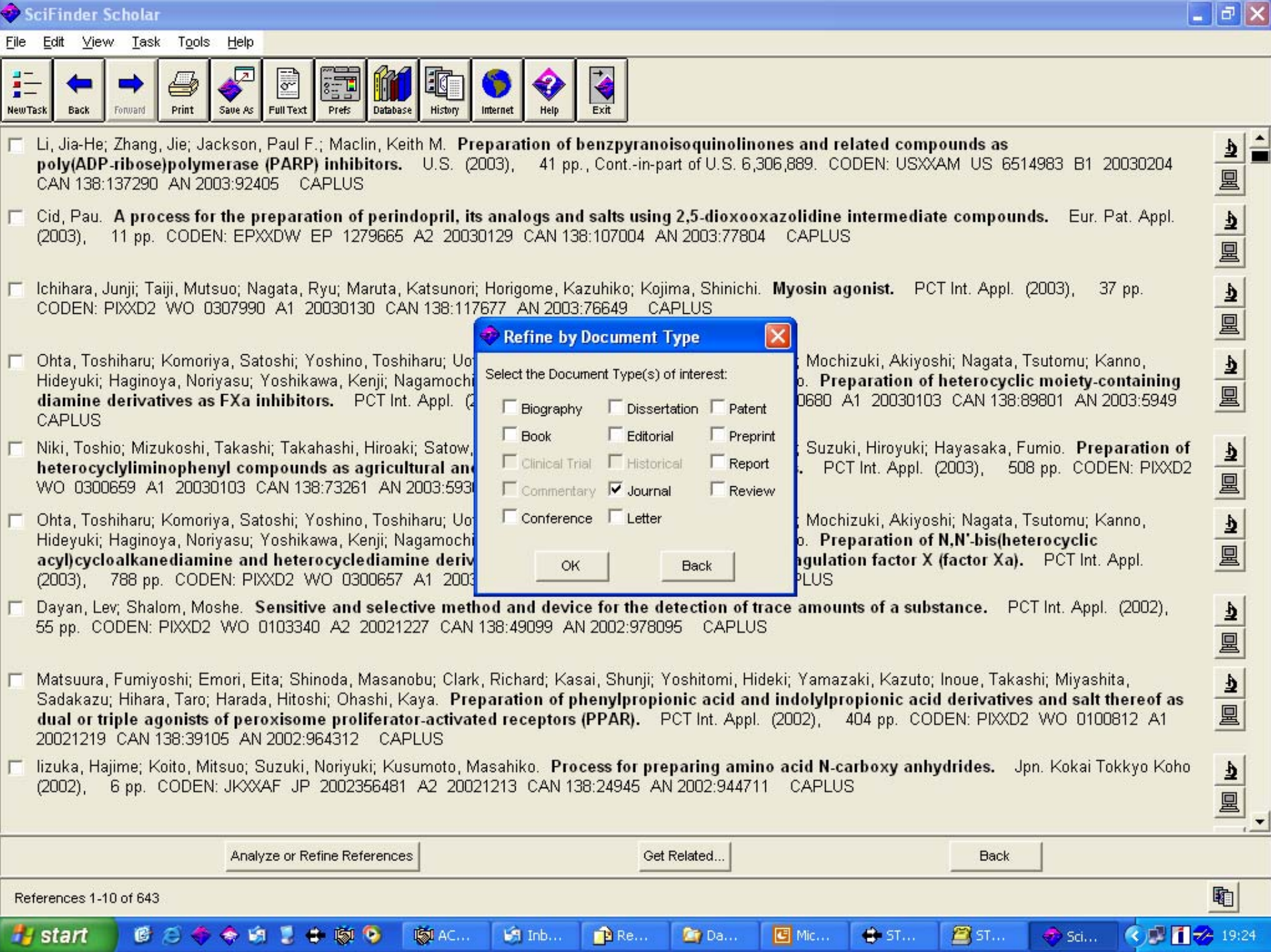
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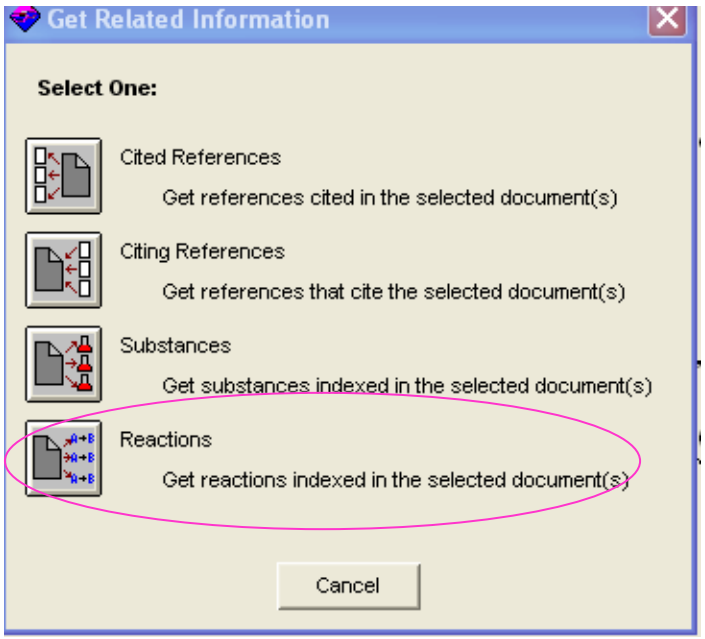

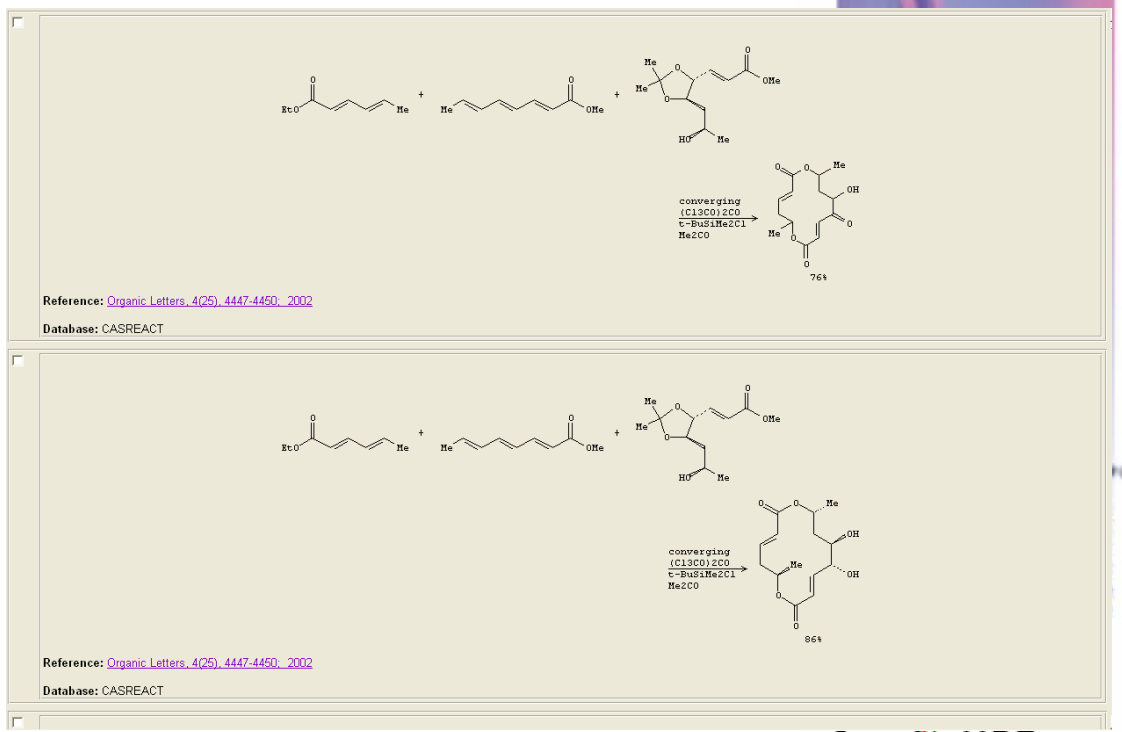

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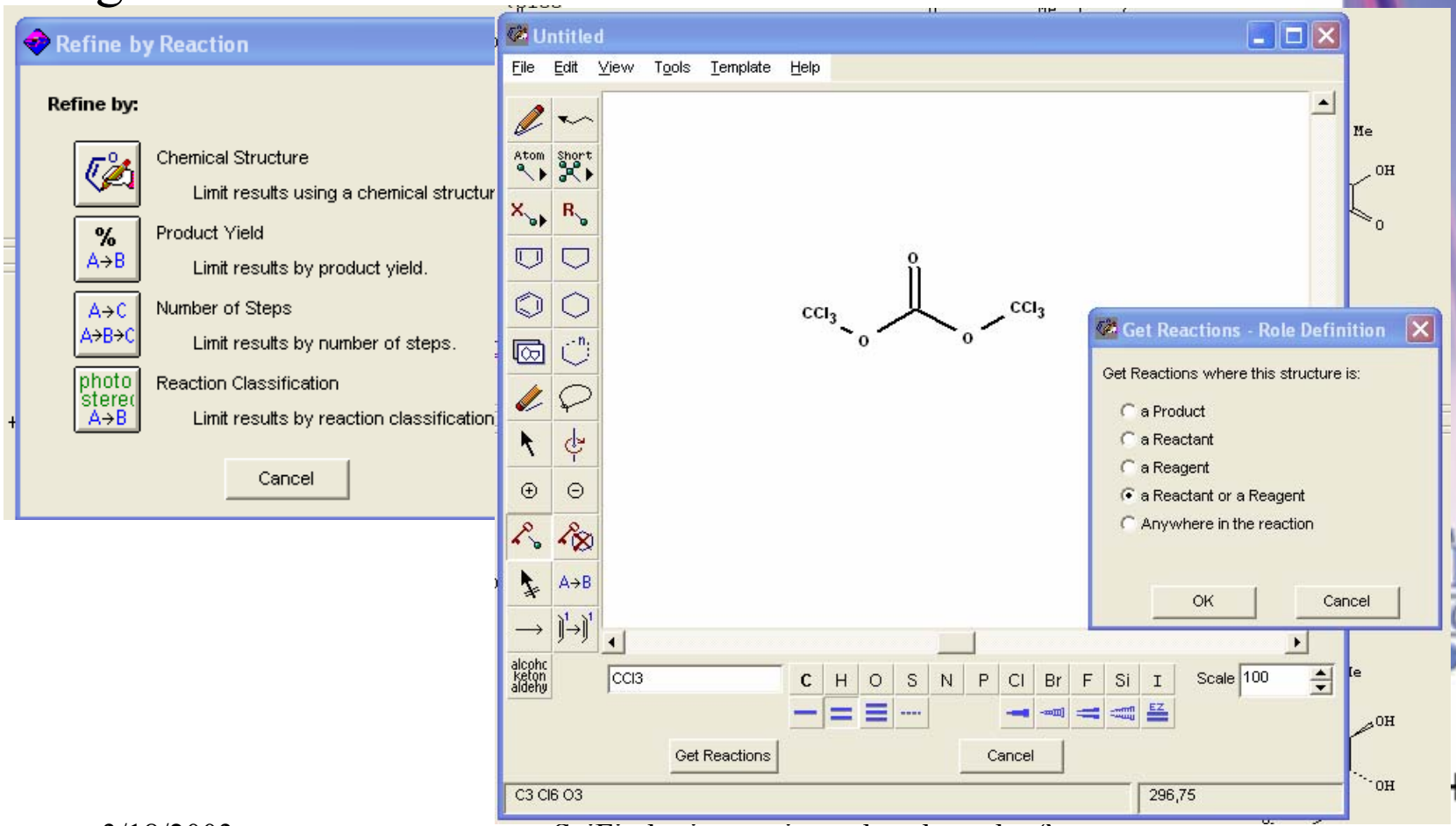

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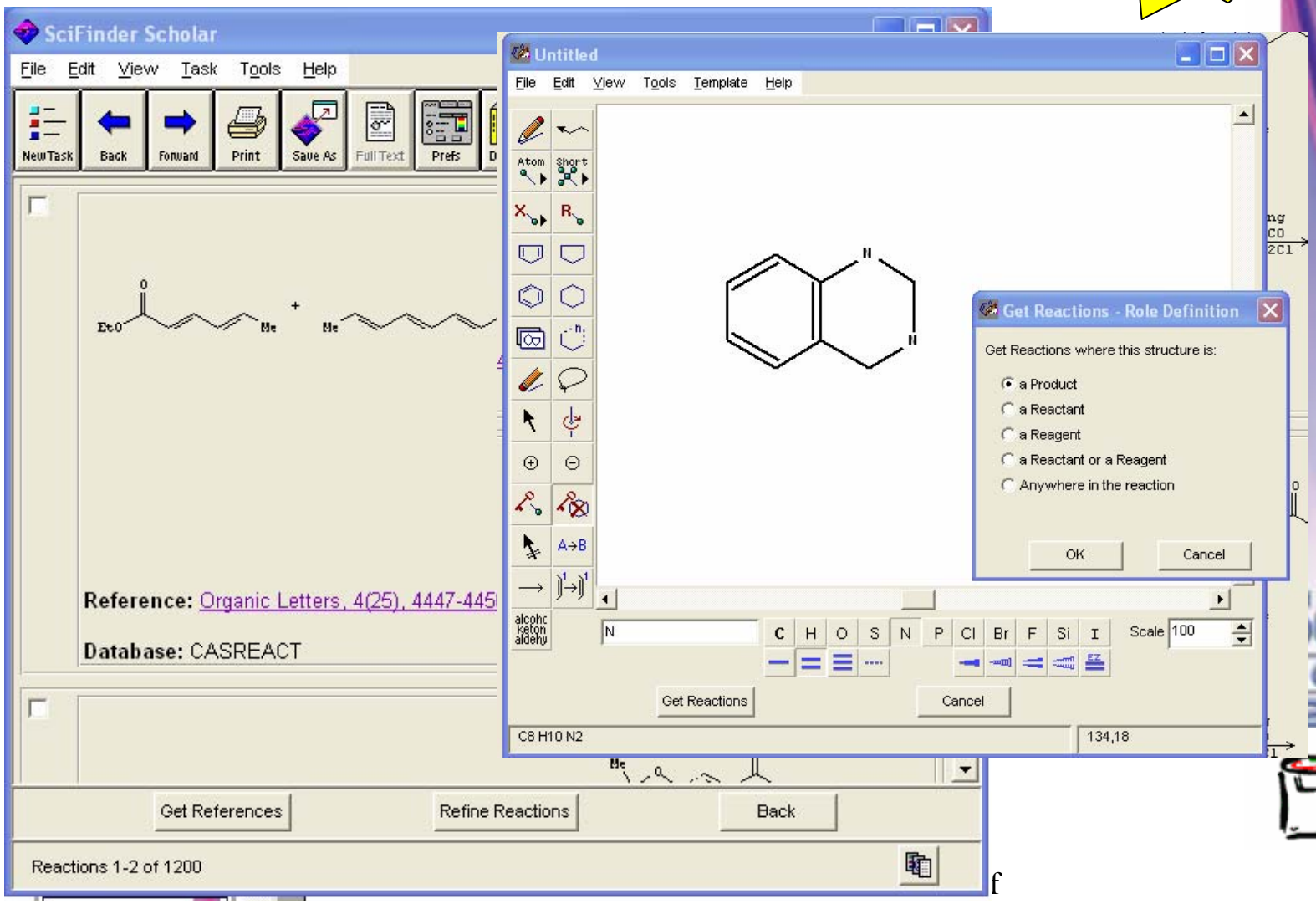

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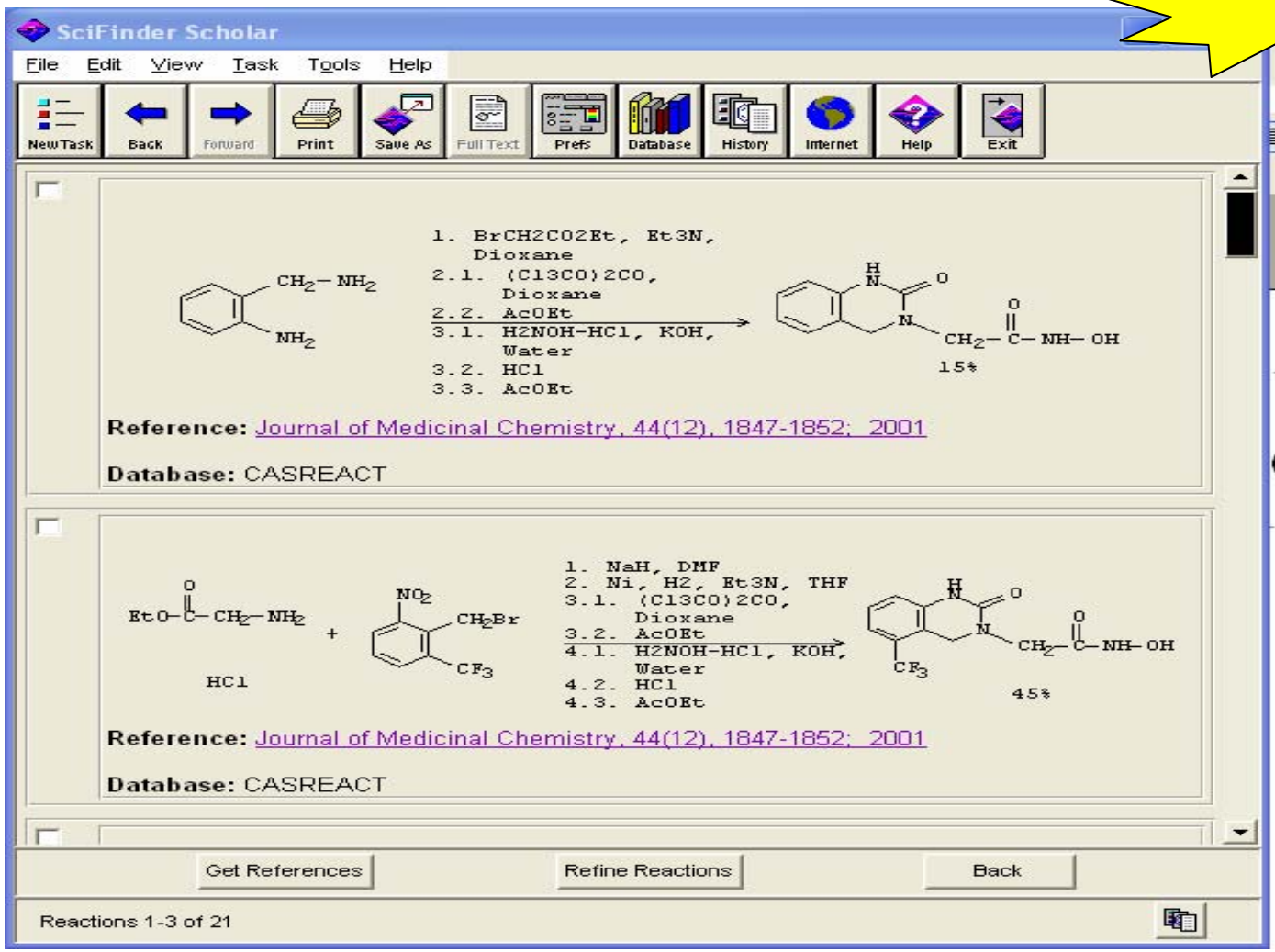

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