# 藥學科技(二)報告

# B303098072 藥三A班 楊上嬅

# --- SciFinder

(1) <u>搜尋題目</u>: Plavix (保栓通) 抗血栓藥

(2) <u>使用成果:</u>

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在下方加入搜尋條件,縮小資料範圍,並增加精確度。

Publication Year(s) 🚸	2008-2011 Examples: 1997, 1995-	1999, 1995-, -1995		
Document Type(s) 🚸	Biography	Dissertation	Patent	
	Book	Editorial	Preprint	
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## 我加入的條件 年分:2008~2011 語言:English、Chinese 公司名稱:Bristol-Myers Squibb

#### 搜尋中的頁面如下:



#### 結果顯示有幾筆相關資料

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#### 點開第一篇文章,全部的內容就顯示出來了。 右邊有一剛開始附加的搜尋條件。

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L. Targeted quantitative bioanalysis in paceurate mass spectrometry: an evaluat power and extraction window, with com By: Xia, Yuan-Qing; Lau, Jim; Olah, Timothy; Jemal, Mohan here is a growing interest in exploring the use of Iq, cl bioanal. labs. as an alternative to the current practice ave investigated the theor, and practical aspects of LC.	blasma using liquid chromatography/high-resolution ion of global selectivity as a function of mass resolving parison of centroid and profile modes med romatog. coupled with full-scan high resoln. accurate mass spectrometry (LC of using LC coupled with tandem mass spectrometry (LC/MS/MS). Therefor IRMS as it relates to the quantitation of drugs in plasma which is the most item set to the second set of the most spectrometry (LC/MS/MS). Therefor item set of the set of the quantitation of drugs in plasma which is the most	Rapid Communications in Mass Spectrometry Volume 25 Issue 19 Pages 2863-2878 Journal 2011 C/HRMS) CoDEN: RCMSEF ISSN: 0951-4198
commonly used matrix in pharmacokinetics studies. In nterferences from endogenous plasma components by moder chromatog. conditions typically used for LC/MS/M esolving power in both profile and centroid modes. Fre he [M + H]+ ions of 153 model drugs using different m n each ELC. Fewer endogenous peaks are detected usi power can be considered adequate for the selective det world missing data points in the analyte ELC peak, the M 'ull width at half max. (FWHM) of the profile mass peak development under the specified chromatog. and sampl resolving power, is smaller for the profile mode when cc	order to assess the overall selectivity of HRM5, we evaluated the potential analyzing acetonitrile-pptd. blank human plasma ext. using an LC/HRM5 syst 5 bioanal, with the acquisition of total ion chromatograms (TICs) using 10 k m each TIC, we generated extd. ion chromatograms (EICs) of the exact mas ass extn. windows (MEWs) and detd. the no. of plasma endogenous peaks of plingher resolving power, narrower MEW, and centroid mode. A 20 k resolv of using in plasma. To achieve desired analyte EIC selectivity and simulta IV used should not be too wide or too narrow and should be a small fractic It is recommended that the optimum MEW be established during method prepn. conditions. In general, the optimum MEW, typically ≤ ±20 pm for moared with the centroid mode. Copyright (2 011 John Wiley & Sons, Ltd.	DOI: 10.1002/rcm.5178 tem and 20 k Getected Ving aneously on of the r 20 k
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在這裡可以詳細的閱讀文章,並再依目標做搜尋條件的修改。

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老師上課操作的時候有示範<專利地圖>給我們看,但我們學校的版本沒有這個功能。

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## <u>(3) 學習心得:</u>

Thomson Innovation 這個系統非常多功能,也很容易上手。他的介面簡單、清楚, 不會太難懂。

我覺得布林邏輯運算元的功能很方便,利用簡單的數學觀念,輕鬆篩選自己要的 資料,使搜尋結果更精確。更重要的是可以減少查資料的時間。

除了布林邏輯外,搜尋時也可以加入\*(開放式又切截)、?(後面加一個字), 等 符號幫助查尋。

這個專利檢索平台包含的範圍非常廣闊,方便發明人及時搜尋專利資訊,跟 Scifinder 一樣,避免不必要的時間浪費。