

Title (en)

METHOD FOR DETERMINING THE MAXIMUM MASS PEAK IN MASS SPECTROMETRY

Title (de)

VERFAHREN ZUR BESTIMMUNG DES MAXIMUMS DES MASSENPEAKS IN DER MASSENSPEKTROMETRIE

Title (fr)

PROCÉDÉ DE DÉTERMINATION DU MAXIMUM DU PIC DE MASSE DANS LA SPECTROMÉTRIE DE MASSE

Publication

EP 2820667 B1 20200422 (DE)

Application

EP 13708748 A 20130228

Priority

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- EP 2013054055 W 20130228

Abstract (en)

[origin: WO2013127933A2] A fast method for determining molecular mass using mass spectrometry has the following steps: specifying a first adjusting value (M1) of the mass spectrometer, recording the associated signal amplitude (A1), specifying a second adjusting value (M2) which is different to the first, measuring the associated second signal amplitude (A2), specifying a third adjusting value (M3) which is different to the first (M1) and the second (M2) adjusting value, measuring the associated third signal amplitude (A3), determining a quadratic function containing the measured amplitude values as y-values and the specified adjusting values as x-values, determining the maximum of the quadratic function, wherein the searched adjusting value is determined from the x-value of the maximum.

IPC 8 full level

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CPC (source: EP RU US)

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Citation (examination)

DOMINIC M. DESIDERIO ET AL: "High resolution mass spectral photoplate data acquired and reduced with a real time remote time-shared digital computer", ANALYTICAL CHEMISTRY, vol. 40, no. 14, 1 December 1968 (1968-12-01), US, pages 2090 - 2096, XP055388103, ISSN: 0003-2700, DOI: 10.1021/ac50158a027

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