

Title (en)

MULTIPORE DETERMINATION OF FRACTIONAL ABUNDANCE OF POLYNUCLEOTIDE SEQUENCES IN A SAMPLE

Title (de)

MULTIPORENBESTIMMUNG DER FRAKTIONIERTEN ABUNDANZ VON POLYNUKLEOTIDSEQUENZEN IN EINER PROBE

Title (fr)

DÉTERMINATION PAR MULTIPORES DE L'ABONDANCE FRACTIONNAIRE DE SÉQUENCES POLYNUCLÉOTIDIQUES DANS UN ÉCHANTILLON

Publication

**EP 3959331 A1 20220302 (EN)**

Application

**EP 19925663 A 20190422**

Priority

US 2019028518 W 20190422

Abstract (en)

[origin: WO2020219011A1] Disclosed herein are methods and compositions for determining an improved estimate of the fractional abundance of target analytes (e.g., specific polynucleotide sequences) in a sample using a nanopore sensor having one or more nanopores-, e.g., by correcting errors inherent to identifying and correlating electrical signals to amounts of a target analyte or reference analyte in a sample.

IPC 8 full level

**C12Q 1/68** (2018.01); **G01N 27/447** (2006.01); **G01N 27/49** (2006.01)

CPC (source: EP KR)

**C12Q 1/68** (2013.01 - EP KR); **G01N 27/447** (2013.01 - KR); **G01N 27/49** (2013.01 - KR); **G01N 33/48721** (2013.01 - KR); **C12Q 2565/631** (2013.01 - KR); **G01N 33/48721** (2013.01 - EP)

Citation (search report)

See references of WO 2020219011A1

Designated contracting state (EPC)

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Designated extension state (EPC)

BA ME

DOCDB simple family (publication)

**WO 2020219011 A1 20201029**; CN 113966403 A 20220121; EP 3959331 A1 20220302; JP 2022530016 A 20220627; KR 20220011639 A 20220128

DOCDB simple family (application)

**US 2019028518 W 20190422**; CN 201980097297 A 20190422; EP 19925663 A 20190422; JP 2021562858 A 20190422; KR 20217038017 A 20190422