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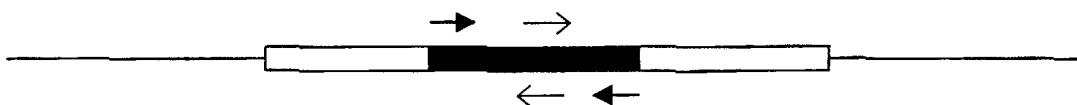
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(54) Title: METHOD OF IDENTIFYING A NUCLEIC ACID SEQUENCE



(57) Abstract: The invention describes a method of utilizing sequence from a partial nucleic acid fragment to identify flanking sequences that are contiguous to that partial nucleic acid fragment. The invention also includes optimized parameters for high throughput assay to isolate and characterize clones containing upstream and downstream flanking sequences from any piece of DNA to be used as a starting point. A kit containing components used in the method of the invention is provided.

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INTERNATIONAL SEARCH REPORT

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A. CLASSIFICATION OF SUBJECT MATTER

IPC 7 C12Q1/68

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 7 C12Q

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

EPO-Internal, WPI Data, BIOSIS

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category °	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 96 38591 A (INCYTE PHARMA INC) 5 December 1996 (1996-12-05)	1-3, 6, 7, 11-16, 18-20, 29-32, 39
Y	Abstract page 10, line 4 -page 13, line 11; figures 1-6; example 1 * whole document *	1-39
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☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

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Y	WO 93 12257 A (HYBRITECH INC) 24 June 1993 (1993-06-24) page 9, line 22; figure 1	1-39

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Int. Patent Application No

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Information on patent family members

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