

Title (en)
METHODS TO IDENTIFY POLYNUCLEOTIDE AND POLYPEPTIDE SEQUENCES WHICH MAY BE ASSOCIATED WITH PHYSIOLOGICAL AND MEDICAL CONDITIONS

Title (de)
VERFAHREN ZUR IDENTIFIZIERUNG VON POLYNUKLEOTID- UND POLYPEPTIDSEQUENZEN WELCHE MIT PHYSIOLOGISCHEN UND MEDIZINISCHEN ZUSTÄNDEN ASSOZIIERT SIND

Title (fr)
TECHNIQUES D'IDENTIFICATION DE SEQUENCES DE POLYNUCLEOTIDES ET DE POLYPEPTIDES POUVANT ETRE ASSOCIEES A DES ETATS PHYSIOLOGIQUES ET MEDICAUX

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Abstract (en)
[origin: WO9939006A2] The present invention provides methods for identifying polynucleotide and polypeptide sequences in human and/or non-human primates which may be associated with a physiological condition, such as disease (including susceptibility (human) or resistance (chimpanzee) to development of AIDS). The methods employ comparison of human and non-human primate sequences using statistical methods. Sequences thus identified may be useful as host therapeutic targets and/or in screening assays.

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Citation (examination)
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• HUGHES A.L.; NEI M.: "PATTERN OF NUCLEOTIDE SUBSTITUTION AT MAJOR HISTOCOMPATIBILITY COMPLEX CLASS I LOCI REVEALS OVERDOMINANT SELECTION", NATURE, vol. 335, 8 September 1988 (1988-09-08), NATURE PUBLISHING GROUP, LONDON, GB, pages 167 - 170, XP001237263
• HUGHES A.L.; NEI M.: "NUCLEOTIDE SUBSTITUTION AT MAJOR HISTOCOMPATIBILITY COMPLEX CLASS II LOCI: EVIDENCE FOR OVERDOMINANT SELECTION", PROC. NATL. ACAD. SCI. USA, vol. 86, February 1989 (1989-02-01), WASHINGTON, DC, US, pages 958 - 962, XP001237260

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