

Title (en)
METHODS AND COMPOSITIONS FOR EVALUATING GENETIC MARKERS

Title (de)
VERFAHREN UND ZUSAMMENSETZUNG ZUR BEWERTUNG GENETISCHER MARKER

Title (fr)
PROCÉDÉS ET COMPOSITIONS POUR L'ÉVALUATION DE MARQUEURS GÉNÉTIQUES

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Application
EP 14762322 A 20140314

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Abstract (en)
[origin: WO2014143994A2] Aspects of the invention relates to methods and compositions that are useful to reduce bias and increase the reproducibility of multiplex analysis of genetic loci. In some configurations, predetermined preparative steps and/or nucleic acid sequence analysis techniques are used in multiplex analyses for a plurality of genetic loci in a plurality of samples.

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Citation (search report)
• [E] WO 2014074246 A1 20140515 - GOOD START GENETICS INC [US]
• [Y] US 2012165202 A1 20120628 - PORRECA GREGORY [US], et al
• [Y] ISOSOMPPI JUHA ET AL: "Disease-causing mutations in the CLRN1 gene alter normal CLRN1 protein trafficking to the plasma membrane.", MOLECULAR VISION 2009, vol. 15, 2009, pages 1806 - 1818, XP055310082, ISSN: 1090-0535
• [IP] MARK A. UMBARGER ET AL: "Next-generation carrier screening", GENETICS IN MEDICINE, vol. 16, no. 2, 13 June 2013 (2013-06-13), pages 132 - 140, XP055186040, ISSN: 1098-3600, DOI: 10.1038/gim.2013.83
• [IP] STEPHANIE HALLAM ET AL: "Validation for Clinical Use of, and Initial Clinical Experience with, a Novel Approach to Population-Based Carrier Screening using High-Throughput, Next-Generation DNA Sequencing", JOURNAL OF MOLECULAR DIAGNOSTICS, THE, vol. 16, no. 2, 27 December 2013 (2013-12-27), US, pages 180 - 189, XP055309745, ISSN: 1525-1578, DOI: 10.1016/j.jmoldx.2013.10.006
• See references of WO 2014143994A2

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