

Comparison of two subtelomeric assays for the screening of chromosomal rearrangements: analysis of 383 patients, literature review and further recommendations

Por: [Maria, LS](#) (Santa Maria, Lorena)^[1]; [Faundes, V](#) (Faundes, Victor)^[1]; [Curotto, B](#) (Curotto, Bianca)^[1]; [Morales, P](#) (Morales, Paulina)^[1]; [Morales, K](#) (Morales, Karla)^[1,2]; [Aliaga, S](#) (Aliaga, Solange)^[1]; [Pugin, A](#) (Pugin, Angela)^[1]; [Alliende, MA](#) (Angelica Alliende, Maria)^[1]

JOURNAL OF APPLIED GENETICS

Volumen: 57

Número: 1

Páginas: 63-69

DOI: 10.1007/s13353-015-0295-4

Fecha de publicación: FEB 2016

[Ver información de revista](#)

Resumen

Intellectual disability (ID) and global development delay (GDD) are caused by genetic factors such as subtelomeric rearrangements (SR) in 25 % of patients. There are several assays currently available to detect SR, but subtelomeric fluorescence in situ hybridisation (Subt-FISH) and subtelomeric multiplex ligation-dependent probe amplification (Subt-MLPA) have been the most frequently used. However, the diagnostic yield of each technique has not been compared. We reviewed the results of SR screening over a ten-year period in Chilean patients with ID/GDD using Subt-FISH and/or Subt-MLPA, compared the diagnostic yield of both tools and reviewed the corresponding literature. A total of 383 cases were included in this study, of which 53.8 % were males. The overall diagnostic yield was 8.9 % between both methods, but Subt-MLPA showed a higher performance than Subt-FISH ($p = 0.002$). A total of 4,181 patients with ID/GDD have been studied worldwide with Subt-MLPA and other subtelomeric assays, and 244 (5.84 %) had a pathogenic SR. It is estimated that Subt-MLPA may detect 92.6 % of the total cases with SR. The capacity of detecting tandem duplication and other critical regions, as well as the use of two MLPA kits, may explain the higher performance of this tool over Subt-FISH. Therefore, we recommend the use of this subtelomeric method as a cost-effective way to study ID/GDD patients.

Palabras clave

Palabras clave de autor: [Intellectual disability](#); [Global development delay](#); [Subtelomeric rearrangements](#); [Fluorescence in situ hybridisation](#); [Multiplex ligation-dependent probe amplification](#)

KeyWords Plus: [DEPENDENT PROBE AMPLIFICATION](#); [UNEXPLAINED MENTAL-RETARDATION](#); [INTELLECTUAL DISABILITY](#); [DEVELOPMENTAL-DISABILITIES](#); [CONGENITAL-ANOMALIES](#); [MLPA](#); [FISH](#); [IDENTIFICATION](#); [INDIVIDUALS](#); [MECHANISMS](#)

Información del autor

Dirección para petición de copias: Faundes, V (autor para petición de copias)

- + Univ Chile, Lab Genet & Enfermedades Metab, INTA, Ave El Libano 5524, Santiago 13811, Chile.

Direcciones:

- + [1] Univ Chile, Lab Genet & Enfermedades Metab, INTA, Ave El Libano 5524, Santiago 13811, Chile
- + [2] Univ Desarrollo, Escuela Tecnol Med, Fac Med, Santiago, Chile

Direcciones de correo electrónico: vfaundes@inta.cl

Editorial

SPRINGER HEIDELBERG, TIERGARTENSTRASSE 17, D-69121 HEIDELBERG, GERMANY

Categorías / Clasificación

Áreas de investigación: Biotechnology & Applied Microbiology; Genetics & Heredity

Categorías de Web of Science: Biotechnology & Applied Microbiology; Genetics & Heredity

Información del documento

Tipo de documento: Review

Idioma: English

Número de acceso: [WOS:000369406200007](#)

ISSN: 1234-1983

eISSN: 2190-3883

Información de la revista

- **Impact Factor:** [Journal Citation Reports®](#)

Otra información

Número IDS: DC7ND

Referencias citadas en la Colección principal de Web of Science: **30**

Veces citado en la Colección principal de Web of Science: **0**