## **Short Report**



Eur J Hum Genet 1997;5:336-337

Received: November 13, 1996 Revision received: June 30, 1997 Accepted: July 7, 1997

Angela F. Brady<sup>a</sup>
C. Ruth Jamieson<sup>a</sup>
Ineke van der Burgt<sup>b</sup>
Andrew Crosby<sup>a</sup>
Margo van Reen<sup>b</sup>
Hannie Kremer<sup>b</sup>
Edwin Mariman<sup>b</sup>
Michael A. Patton<sup>a</sup>
Steve Jeffery<sup>a</sup>

## Further Delineation of the Critical Region for Noonan Syndrome on the Long Arm of Chromosome 12

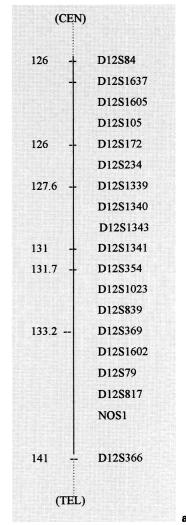
- <sup>a</sup> Medical Genetics Unit, St. George's Hospital Medical School, London, UK;
- Department of Human Genetics, University Hospital Nijmegen, The Netherlands

**Key Words** 

Noonan syndrome  $\cdot$  Chromosome  $12q \cdot$  Mapping studies

Noonan syndrome (NS) is an autosomal dominant disorder characterised by typical facies, short stature and congenital heart disease [1], with an estimated incidence between 1 in 1,000 and 1 in 2,500 [2]. We have recently localised a gene for NS to 12q22-qter in a large 3-generation Dutch family [3, 4]. Recombination events placed the gene in a 14-cM interval between the markers D12S84 and D12S366.

In the present study, we analysed the 3-generation family with newly isolated CA-repeat markers derived from the interval between D12S84 and D12S366 (fig. 1a) [5]. Genotyping was performed as previously described [3]. At the proximal border of the critical region, a recombination was detected between D12S105 and D12S1637 (fig. 1b, III.3). D12S1605 was not informative. In the distal part of the critical region, a recombination was found between D12S79 and NOS1 (fig. 1b, III.4). Thus, the interval which contains the gene mutated in this family to produce NS is now defined by markers D12S105 and NOS1. As NOS1 has not been positioned genetically, it is impossible to determine precisely by what amount the physical and genetic intervals have been reduced by these new recombinations. Existing physical maps indicate that



**Fig. 1. a** Genetic map of the relevant region of chromosome 12q. Distances are given in centimorgans [5, 6]. CEN = Centromere; TEL = telomere. **b** Pedigree and haplotypes of the Dutch family. The markers used for haplotype construction are shown. The cross-hatched bars represent haplotypes, which co-segregate with the disorder, the black bars represent areas of uninformativeness and the white bars represent the unaffected haplotype.

NOS1 is positioned less than 1.7 Mb distally of D12S79, the most distal marker in the NS interval to be assigned a genetic distance [5]. D12S79 is 6 cM distal to D12S105, so the NS interval can be summarized by: D12S105-(6 cM)-D12S79-(1.7 Mb)-NOS1, which is likely to represent a substantial reduction in the interval.

## **Acknowledgements**

We are most grateful for the financial support from the British Heart Foundation, the Birth Defects Foundation and the Dutch Heart foundation (grant No. R96025).

## References

- Sharland M, Burch M, McKenna WM, Patton MA: A clinical study of Noonan syndrome. Arch Dis Child 1992;67:178–183.
- 2 Allanson JE: Noonan syndrome. J Med Genet 1985;24:9–13.
- 3 Jamieson CR, van der Burgt I, Brady AF, van Reen M, Elsawi MM, Hol F, Jeffery S, Patton MA, Mariman E: Mapping a gene for Noonan syndrome to the long arm of chromosome 12. Nature Genet 1994;8:357–360.
- 4 van der Burgt I, Berends E, Lommen E, Beersum S, Hanel B, Mariman E: Clinical and molecular studies in a large Dutch family with Noonan syndrome. Am J Med Genet 1994;53: 1\_8
- 5 Krauter K, Montgomery K, Yoon S-J, LeBlanc-Straceski J, Renault B, Marondel I, Herdman V, Cupelli L, Banks A, Lieman J, Menninger J, Bray-Ward P, Nadkarni P, Weissenbach J, Le Paslier D, Rigault P, Chumakov I, Cohen D,
- Miller P, Ward D, Kucherlapati R: A secondgeneration YAC contig map of human chromosome 12. Nature 1995;337(suppl):321–333.
- 6 Dib C, Faure S, Fizames C, Samson D, Drouot N, Vignal A, Millasseau P, Marc S, Lathrop M, Gyapay G, Morissette J, Weissenbach J: A comprehensive genetic map of the human genome based on 5,264 microsatellites. Nature 1996;380:152-154.

