ARTICLE

Prevalence of lysosomal storage diseases in Portugal

Rui Pinto^{1,2}, Carla Caseiro¹, Manuela Lemos¹, Lurdes Lopes¹, Augusta Fontes¹, Helena Ribeiro¹, Eugénia Pinto¹, Elisabete Silva¹, Sónia Rocha¹, Ana Marcão², Isaura Ribeiro^{1,2}, Lúcia Lacerda^{1,2}, Gil Ribeiro^{1,2}, Olga Amaral^{1,2}, MC Sá Miranda^{*,1,2}

¹Instituto de Genética Médica Jacinto de Magalhães, Porto, Portugal; ²Instituto de Biologia Molecular & Celular (IBMC), Portugal

Lysosomal storage diseases (LSDs) are a group of inherited metabolic disorders individually considered as rare, and few data on its prevalence has been reported in the literature. The overall birth prevalence of the 29 different LSDs studied in the Portuguese population was calculated to be 25/100 000 live births, twice the prevalence previously described in Australia and in The Netherlands. The comparison of the prevalence profile of the LSDs presenting a prevalence higher than 0.5/100 000 in the Portuguese, Dutch and Australian populations showed, in the Portuguese, the existence of a higher prevalence of GM2 gangliosidosis (B variant), mucopolysaccharidoses (II and III), Niemann-Pick type C and metachromatic leukodystrophy (MLD), and a lower prevalence of Pompe and Fabry. The highest prevalence value for a single LSD is the one of GM2 gangliosidosis (B variant), corresponding to 3/100 000, a value which is significantly higher than the prevalence of the most frequent LSD in Dutch, Pompe disease (2/100 000) and Australians, Gaucher's disease (GD) (1.8/100 000). It is worth noting that the highest prevalence of GM2 gangliosidosis found in the Portuguese is mainly due to the existence of a unique subtype, the rare juvenile B1 variant.

European Journal of Human Genetics (2004) 12, 87–92. doi:10.1038/sj.ejhg.5201044
Published online 17 December 2003

Keywords: inherited metabolic disorders; lysosomal storage diseases; genetic epidemiology; birth prevalence; Portuguese population

Introduction

Lysosomal storage disorders (LSDs) are a heterogeneous group of inherited metabolic diseases characterized by the impairment of the intralysosomal catabolic pathways (for a recent review, see the corresponding chapters in Scriver *et al*¹).

With the exception of particular diseases in specific ethnic groups, few data on the frequency of LSDs can be found in the literature¹ and, as far as we know, prevalence data on LSDs considered as a group have only been reported in The Netherlands² and in Australia.³ Besides contributing to the knowledge of the genetic character-

istics of each population, prevalence data are important to delineate prevention and therapeutic strategies.

Between 1982 and 2001, 353 patients with an LSD were identified in this laboratory by studying 29 different LSDs in about 4700 cases suspected of an Inherited metabolic disease. On the basis of these data, we report for the first time, the relative frequency and the birth prevalence of LSDs in a Southern European population.

Patients and methods

Geographical origin of the cases studied

The Enzymology Unit of Instituto de Genética Médica Jacinto de Magalhães, located in Porto, is the only Portuguese Laboratory providing post- and prenatal diagnosis of LSDs and receiving samples from Health Services located throughout the country. Probably due to its geographical location, in the North of Portugal, a higher

*Correspondence: Dr MC Sá Miranda, IBMC, Rua do Campo Alegre 823, 4150-180 Porto, Portugal. Tel.: 351 22 607 4970; Fax: 351 22 609 2404
E-mail: mcsamir@ibmc.up.pt

Received 16 August 2002; revised 28 April 2003; accepted 13 May 2003

number of requests are received from Health Services located in the North and Central regions of the country than from other regions. Since there is only one other Portuguese laboratory, located in Lisbon, offering postnatal diagnosis for just a few LSDs, during the last 10 years a time tendency for a more uniform coverage of the country by our lab has been observed (data not shown). The relative frequency was determined considering two groups of patients, according to their possible origin, from the North and from other regions. As the population from the Northern part of the country has been studied more extensively, we decided to use only these data in the birth prevalence calculation.

Analytical methods

Current methodology, extensively reviewed, was used in the determination of enzyme activities^{4,5} and in the analysis of substrates.^{6,7} Standard molecular biology techniques, such as PCR, SSCP, RFLP and sequencing, were used in mutation analysis. Genotype analysis was performed to exclude pseudodeficiency alleles in MLD,⁸ to identify polymorphisms responsible for low enzymatic activity in Krabbe's disease⁹ and as diagnostic criteria in NCL.^{3,10} Genotype analysis was also carried out in the four most frequent sphingolipidoses.¹¹

Patients

In general, the enzymatic activity was determined in a blood sample, and subsequently confirmed in cultured skin fibroblasts. Urinary excretion of substrates was considered in mucopolysaccharidoses (MPSs), oligosaccharidoses and metachromatic leukodystrophy. Patient's birthplace, age and clinical phenotype were collected from the laboratory records.

Prevalence estimation

Prevalence was calculated for each disease by dividing the total number of diagnosed cases (post- and prenatal diagnosis) by the total number of live births that occurred between the years of birth of the older and the younger patients (birth period). In relation to GM2 gangliosidosis B variant and MLD, diseases presenting three different clinical phenotypes (infantile, juvenile and adult or chronic), the prevalence of each phenotype was determined and the overall prevalence was calculated by adding the prevalence of each phenotype. The noneuronopathic form of GD (type 1) includes a very wide spectrum of clinical phenotypes, the age of onset ranging between 3 and 80 years. In order to overcome the underestimation due to the very large birth period, two subgroups were considered according to the age of clinical manifestations, before and after 15 years age. For diseases in which only one patient was diagnosed, the prevalence was calculated using the number of live births during the ascertainment period (1982–2001). Data on the number of live births in

Portugal were collected at the Instituto Nacional de Estatística, Porto, Portugal.

Results

The relative frequency of LSDs in Portuguese population from the North and from other regions is compared in Table 1. Three main groups were considered:

MPSs (MLP), sphingolipidoses and oligosaccharidoses; mucopolipidoses, Pompe and neuronal ceroid lipofuscinoses were considered apart. The relative frequency of each LSD was calculated: in relation to the total number of diagnosed patients and in relation to the number of patients of each main group considered. As a group, MPSs and sphingolipidoses present similar frequencies in populations of both geographical regions. Exceptions to the relatively uniform distribution of MPSs along the country are MPS I and II. The first one (Hurler's disease) represents 13% in the North and 35% in the rest of the country; in contrast, Hunter's disease represents 34 and 8% of the MPS in the same respective regions. The relatively high frequency of MPS III (Sanfilippo) among MPSs is mainly due to MPS IIIB.

Regarding the sphingolipidoses, the most striking difference between both samples is the higher frequency of MLD in the North (17 vs 2%). Mucopolipidoses (II and III), generally considered as rare, account for 11 patients out of 353. Neuronal ceroid lipofuscinoses (NCL), whose study begun only in 1999, account for 7% of total LSDs in the Northern and 5% in the non-Northern samples.

Birth prevalence was calculated using the more extensively studied Northern Portuguese population data. Table 2 compares the results obtained with the birth prevalence previously described in the Netherlands² and in Australia.³

Genotype analysis data were used to investigate if the high prevalence of the more frequent sphingolipidoses was due to a high frequency of particular mutations. As can be observed in Table 3, one single mutation represents 58, 53 and 64% of the mutated alleles in unrelated patients with Tay-Sachs disease, Gaucher's disease and MLD, respectively.

Discussion

Overall, the birth prevalence of LSD determined in the Northern Portuguese population (25 per 100 000 live births) is about the double of that previously reported in the Netherlands² and in Australia;³ it must be noticed that the observed discrepancies cannot be explained by the slight differences in the calculation methods. In fact, the incidence rates of the different LSDs in the Portuguese (data not shown) are similar to the birth prevalence values presented. When only MPSs are considered, birth prevalence (4.8 per 100 000) is very close to the values previously reported.^{2,3} In contrast, the sphingolipidoses

Table 1 Relative frequency of LSD in Portugal

Disease	Studied since	Postnatal		Post- and prenatal					
		No. of cases	Age at diagnosis	North			Other regions		
				N	% Group	% Total	N	% Group	% Total
MPS IH+IS (Hurler/Scheie)	1982	20		8	12.9	3.6	14	35.0	9.2
MPS IH (Hurler)		17	0–19	6	9.7	2.7	13	32.5	8.5
MPS IS (Scheie)		3	11–26	2	3.2	0.9	1	2.5	0.7
MPS II (Hunter)	1982 ^a	22	1–25	21	33.9	9.5	3	7.5	2.0
MPS III (Sanfilippo)	1982	23		14	22.6	6.3	10	25.0	6.5
MPS IIIA (Sanfilippo A)	^a	3	3–18	0	0.0	0.0	3	7.5	2.0
MPS IIIB (Sanfilippo B)		17	1–27	13	21.0	5.9	5	12.5	3.3
MPS IIIC (Sanfilippo C)	^a	3	5–8	1	1.6	0.5	2	5.0	1.3
MPS IVA (Morquio A)	1982	10	0–16	6	9.7	2.7	5	12.5	3.3
MPS VI (Maroteaux-Lamy)	1982	15	1–22	10	16.1	4.5	6	15.0	3.9
MPS VII (Sly)	1982	1	1–1	0	0.0	0.0	1	2.5	0.7
Multiple sulfatase deficiency	1982	4	1–10	3	4.8	1.4	1	2.5	0.7
Total mucopolysaccharidoses		91		62	100.0	27.9	40	100.0	26.1
GM1-gangliosidosis	1982	9	0–2	4	3.3	1.8	6	6.5	3.9
GM2-gangliosidosis (α -subunit)	1982	46		32	26.4	14.4	15	16.3	9.8
GM2-gangliosidosis (α -subunit) infantile		7	1–2	3	2.5	1.4	4	4.3	2.6
GM2-gangliosidosis (α -subunit) juvenile		30	2–14	23	19.0	10.4	8	8.7	5.2
GM2-gangliosidosis (α -subunit) chronic		9	10–44	6	5.0	2.7	3	3.3	2.0
GM2-gangliosidosis (β -subunit)	1982	3	1–3	3	2.5	1.4	1	1.1	0.7
Krabbe	1988 ^b	11	0–4	9	7.4	4.1	3	3.3	2.0
MLD	1982	21		20	16.5	9.0	2	2.2	1.3
MLD late infantile		11	0–8	11	9.1	5.0	1	1.1	0.7
MLD juvenile		3	4–13	2	1.7	0.9	1	1.1	0.7
MLD adult		7	17–29	7	5.8	3.2	0	0.0	0.0
Fabry	1982	4	32–51	1	0.8	0.5	3	3.3	2.0
Gaucher	1982	84		36	29.8	16.2	48	52.2	31.4
Gaucher type 1 early		21	3–39	8	6.6	3.6	13	14.1	8.5
Gaucher type 1 late		59	16–84	25	20.7	11.3	34	37.0	22.2
Gaucher neurologic (type 2 and 3)		4	0–5	3	2.5	1.4	1	1.1	0.7
Niemman-Pick A+B	1983	11		7	5.8	3.2	4	4.3	2.6
Niemman-Pick type A		4	0–1	4	3.3	1.8	0	0.0	0.0
Niemman-Pick type B		7	2–53	3	2.5	1.4	4	4.3	2.6
Niemman-Pick type C	1985 ^c	18	1–27	9	7.4	4.1	10	10.9	6.5
Total sphingolipidoses		207		121	100.0	54.5	92	100.0	60.1
α -Mannosidosis	1982	3	8–22	1	16.7	0.5	2	22.2	1.3
β -Mannosidosis	1989	1	4–4	1	16.7	0.5	0	0.0	0.0
Sialidosis	1985	2	35–41	0	0.0	0.0	2	22.2	1.3
Fucosidosis	1982	1	5–5	0	0.0	0.0	1	11.1	0.7
Galactosialidosis	1982	2	0–2	2	33.3	0.9	3	33.3	2.0
Aspartylglucosaminuria	1990	3	9–10	2	33.3	0.9	1	11.1	0.7
Total oligosaccharidoses		12		6	100.0	2.7	9	100.0	5.9
MLP II/III	1982	11		10		4.5	2		1.3
MLP II (I cell disease)		7	0–1	7		3.2	1		0.7
MLP III		4	3–11	3		1.4	1		0.7
Pompe	1982	10	0–54	7		3.2	3		2.0
Neuronal ceroid lipofuscinoses		22		16		7.2	7		4.6
NCL1 (Santavuori)	1999	2	6–27	2		0.9	0		0.0
NCL2 (Jansky-Bielschowsky)	1999	3	5–38	2		0.9	1		0.7
NCL3 (Spielmeyer-Vogt)	1999	10	11–28	7		3.2	4		2.6
NCL6	2001	7	10–15	5		2.3	2		1.3
Total LSD		353		222		100.0	153		100.0

Some diagnoses have been confirmed by

^aO. van Diggelen and WJ Kleijer (Rotterdam).^bMT Vanier (Lyon) and D Wenger (Philadelphia).^cMT Vanier (Lyon). MPS = Mucopolysaccharidoses; p/MLP = Mucopolipidoses; NCL = Neuronal Ceroid Lipofuscinoses.

Table 2 Birth Prevalence of LSD

Disease	No. of cases	Birth period	Post and prenatal		
			Northern Portugal	Birth prevalence*	
			NP	NL (2)	AU (3)
MPS IH+IS (Hurler/Scheie)	8		1.33	1.19	1.14
MPS IH (Hurler)	6	1984–1995	1.05		
MPS IS (Scheie)	2	1976–1987	0.28		
MPS II (Hunter)	21	1967–2000	1.09	0.67	0.74
MPS III (Sanfilippo)	14		0.84	1.89 ^a	
MPS IIIA (Sanfilippo A)	0		0.00	1.16	0.88
MPS IIIB (Sanfilippo B)	13	1966–1994	0.72	0.42	0.47
MPS IIIC (Sanfilippo C)	1	1982–2001	0.12	0.21	0.07
MPS IVA (Morquio A)	6	1979–1998	0.60	0.22	0.59
MPS VI (Maroteaux-Lamy)	10	1962–1999	0.42	0.15	0.43
MPS VII (Sly)	0		0.00	0.24	0.05
Multiple sulfatase deficiency	3	1978–1988	0.48	0.05	0.07
Total mucopolysaccharidoses	62		4.8	4.5^b	
GM1-gangliosidosis	4	1985–1998	0.62	0.41	0.26
GM2-gangliosidosis (α -subunit)	32		3.13	0.41	0.50
GM2-gangliosidosis (α -subunit) infantile	3	1985–1990	1.02		
GM2-gangliosidosis (α -subunit) juvenile	23	1974–1997	1.77		
GM2-gangliosidosis (α -subunit) chronic	6	1959–1980	0.34		
GM2-gangliosidosis (β -subunit)	3	1985–1988	1.49	0.34	0.26
Krabbe	9	1984–1999	1.21	1.35	0.71
MLD	20		1.85	1.42 ^c	1.09
MLD late infantile	11	1980–2001	1.12	0.52	
MLD juvenile	2	1974–1984	0.29	0.51	
MLD adult	7	1954–1971	0.45	0.24	
Fabry	1	1982–2001	0.12	0.21	0.86
Gaucher	36		1.35	1.16	1.75
Gaucher type 1 early	8	1953–1987	0.30	0.26	
Gaucher type 1 late	25	1917–1981	0.50	0.64	
Gaucher neurologic (type 2 and 3)	3	1977–1985	0.55	0.26	
Niemman-Pick A+B	7		0.60	0.53	0.40
Niemman-Pick type A	4	1983–2000	0.50		
Niemman-Pick type B	3	1952–1991	0.10		
Niemman-Pick type C	9	1979–1985	2.20	0.35	0.47
Total sphingolipidoses	121		12.6	6.2	
α -Mannosidosis	1	1982–2001	0.12	0.09	0.10
β -Mannosidosis	1	1982–2001	0.12	0.13	
Sialidosis	0		0.00	0.05	0.02
Fucosidosis	0		0.00	0.05	
Galactosialidosis	2	1994–2001	0.77	0.04	
Aspartylglucosaminuria	2	1982–1983	1.72	0.13	0.05
Total oligosaccharidoses	6		2.7		
MLP II/III	10		2.70		0.31
MLP II (I cell disease)	7	1982–2000	0.81	0.16	
MLP III	3	1984–1986	1.89	0.08	
Pompe	7	1940–1999	0.17	2.0	0.69
Neuronal ceroid lipofuscinoses	16		2.14		
NCL1 (Santavuori)	2	1974–1994	0.17		
NCL2 (Jansky-Bielschowsky)	2	1950–1987	0.07		
NCL3 (Spielmeyer-Vogt)	7	1973–1999	0.48		
NCL6	5	1984–1990	1.43		
Total LSD	222		25	14^d	12.99

^aIncludes MPS IIID.^bIncludes MPS IIID and MPS IVB.^cIncludes some unspecified types.^dIncludes Schindler, and ISSD does not include NCL.

NP = Northern Portugal; NL = Netherlands; AU = Australia. *Birth prevalence expressed as number of cases per 100 000 live births.

Table 3 Frequency of mutated alleles in unrelated Portuguese patients with the four most frequent sphingolipidoses: MLD, GM2-Gangliosidosis, Gaucher and Niemann-Pick type C diseases

Gene	Mutation	MIM Id	Nb alleles	%
ARSA	ivs2+1	250100.0033	18	60.0
	I179S	250100.0008	2	6.7
	Other		10	33.3
	Total		30	100.0
HEXA	R178H	606869.0006	44	59.5
	R137X	606869.0019	7	9.5
	Other		23	31.1
	Total		74	100.0
GBA	N370S	606463.0003	68	53.1
	L444P	606463.0001	18	14.1
	Other		42	32.8
	Total		128	100.0
NPC1	A1035V	257220.0017	5	14.7
	C177Y	257220.0016	3	8.8
	I1061T	257220.0010	2	5.9
	Other		24	70.6
	Total		34	100.0

birth prevalence (12.6 per 100 000 live births) in the Northern Portuguese population is the double of the birth prevalence reported in the Dutch (6.2 per 100 000 live births) and Australian (6.3 per 100 000) populations.

The most frequent LSD in the populations of the three countries is Gaucher's disease, its birth prevalence in the Portuguese population (1.4 per 100 000 live births) is similar to the one reported in the Netherlands and in Australia, 1.2 and 1.8 per 100 000 live births, respectively.

Concerning birth prevalence values higher than 0.5 per 100 000 live births in the three populations, the Portuguese present higher prevalence values of GM2 gangliosidosis, mucopolidoses II/III, NPC and MLD, and lower prevalence values of Pompe and Fabry diseases. The highest value for a single LSD is observed with GM2 gangliosidosis, B variant in the Portuguese (3.1 per 100 000 live births), which is considerably higher than the highest prevalence of a single LSD in The Netherlands, Pompe disease (2 per 100 000) and Gaucher's disease (1.8 per 100 000) in Australia.

As happens with other genetic diseases, LSDs are genetically very heterogeneous.¹ However, in certain ethnic groups, a high incidence of specific mutations has been described. The high frequency of HEXA mutations associated with the classic infantile form of Tay-Sachs disease observed among the Ashkenazi Jews (1/4100) as compared with the one observed within the non-Jewish individuals (1/112 000)¹² is an example. Another example is Gaucher's disease N370S GBA mutation, presenting a frequency of 1/17.5 among the Ashkenazi Jews¹³ contrasting with a 1/210 frequency in a non-Ashkenazi population.¹⁴

Genotype analysis of the Portuguese patients presenting GM2 gangliosidosis showed that R178H HEXA mutation (DN allele) represents 58% of the mutated alleles; 44% of the patients are homozygous for this mutation, corresponding to a juvenile form of Tay-Sachs B1 variant.¹⁵ Except for two Spanish patients originated from Galiza¹⁶ no other patients homozygous for 'DN-allele' were, to our knowledge, described in the literature.

Similarly, in the case of Portuguese patients with MLD, a single ARSA mutation, IVS2 + 1GA (allele I), represents 60% of the alleles. The higher frequency of this mutation, in comparison with the 15–43% reported in other European patients,¹⁷ reflects the higher prevalence of the infantile form detected in the Portuguese patients.

In relation to Gaucher's disease, the N370S GBA mutation represents 53% of the mutated alleles, a frequency higher than the one reported in non-Ashkenazi GD patients.¹⁸ According to the gene frequency previously determined in the general Portuguese population¹⁴ about 200 homozygotes for N370S GBA mutation can be predicted in Portugal, contrasting with 21 homozygotes out of 84 Portuguese Gaucher's disease patients diagnosed. This fact suggests that the great majority of the N370S homozygotes are asymptomatic.

Surprisingly, the higher Niemann-Pick type C prevalence observed in the Portuguese population (2.2 per 100 000), as compared with the one estimated in other European (France, West Germany and UK) populations 1/150 000,¹⁹ cannot be explained by a frequent mutation. In fact, a wide genetic heterogeneity was observed, the I1061 T allele of the NPC1 gene, which represents 15% of the mutant alleles in British and French NPC patients,¹⁹ represents only 6% of the NPC1-mutated alleles in the Portuguese NPC patients.²⁰

In conclusion, as a group, LSDs can be considered as a very frequent inborn error of metabolism in the Portuguese Population, presenting a birth prevalence (1/4000 live births) higher than birth prevalence reported in the case of PKU (1/12 000 live births) and congenital hypothyroidism (1/6000 live births).²¹

Acknowledgements

We are grateful to all clinicians who referred cases for LSD study, to Doctors OP van Diggelen, WJ Kleijer (Rotterdam), MT Vanier (Lyon) and DA Wenger (Philadelphia) for invaluable help in difficult diagnoses, and to Mrs Clementina Maia for the search and collection of population data at Instituto Nacional de Estatística, Porto, Portugal.

References

- 1 Scriver SR, Beaudet AL, Sly WS, Valle D (eds): *The metabolic bases of inherited disease*. New-York: McGraw-Hill, 2001, Vol. 3.
- 2 Poorthuis BJ, Wevers RA, Kleijer WJ *et al*: The frequency of lysosomal storage diseases in The Netherlands. *Hum Genet* 1999; **105**: 151–156.
- 3 Meikle PJ, Hopwood JJ, Clague AE, Carey WF: Prevalence of lysosomal storage disorders. *JAMA* 1999; **281**: 249–254.

- 4 Wenger D, Williams C: Screening for lysosomal disorders; in Hommes FA (ed): *Techniques in diagnostic human biochemical genetics*. New York: Wiley-Liss, 1991, pp 587–617.
- 5 Galjaard H: *Genetic metabolic diseases. Early diagnosis and prenatal analysis*. Amsterdam: Elsevier/North Holland, 1980.
- 6 Hopwood JJ, Harrison JR: High resolution electrophoresis of urinary glycosaminoglycans: an improved screening test for the MPS. *Anal Biochem* 1982; **119**: 120–112.
- 7 Sewell AC: Urinary Oligosaccharides; in Hommes FA (ed): *Techniques in diagnostic human biochemical genetics: a laboratory manual*. New York: Wiley-Liss, 1991, pp 219–231.
- 8 Ricketts M, Zhang X, Manowitz P: A method for the rapid detection of arylsulfatase A pseudodeficiency mutations. *Hum Hered* 1995; **45**: 235–240.
- 9 Wenger D, Rafi M, Luzi P: Molecular genetics of Krabbe Disease (globoid cell leukodystrophy). Diagnostic and clinical implications. *Hum Mut* 1997; **10**: 268–279.
- 10 Zhong N, Wisniewski KE, Kaczmarek AL *et al*: Molecular screening of Batten disease: identification of a missense mutation (E295 K) in the CLN3 gene. *Hum Genet* 1998; **102**: 57–62.
- 11 Amaral O, Marcão A, Pinto E, Ribeiro MG, Sá Miranda MC: Análise do genótipo na prevenção das esfingolipidoses. *Arqui Med* 1997; **11**: 24–26.
- 12 Gravel RA, Kaback M, Proia R, Sandhoff K, Suzuki K: The GM2-gangliosidosis; in Scriver SR, Beaudet AL, Sly WS, Valle D (eds): *The metabolic basis of inherited disease*. New-York: McGraw-Hill, 2001, pp 3827–3876.
- 13 Horowitz M, Tzuri G, Eyal N *et al*: Prevalence of nine mutations among Jewish and non-Jewish Gaucher disease patients. *Am J Hum Genet* 1993; **53**: 921–930.
- 14 Lacerda L, Amaral O, Pinto R, Oliveira P, Aerts J, Sá Miranda MC: Gaucher Disease: N370S glucocerebrosidase gene frequency in the Portuguese population. *Clin Genet* 1994; **45**: 298–300.
- 15 Dos Santos MR, Tanaka A, Sá Miranda MC, Ribeiro MG, Maia M, Suzuki K: GM2-Gangliosidosis B1 variant: analysis of β -hexosaminidase α gene mutations in 11 patients from a defined region in Portugal. *Am J Hum Genet* 1991; **49**: 886–890.
- 16 Eiris J, Chabas A, Coll M J, Castro-Gago M: Late infantile and Juvenile form of GM2-gangliosidosis variant B1. *Rev Neurol* 1999; **29**: 435–438.
- 17 von Figura K, Gieselmann V, Jaeken J: Metachromatic leukodystrophy; in Scriver SR, Beaudet AL, Sly WS, Valle D (eds): *The metabolic basis of inherited disease*. New-York: McGraw-Hill, 2001, pp 3695–3724.
- 18 Koprivica V, Stone DL, Park JK *et al*: Analysis and classification of 304 mutant alleles in patients with type 1 and type 3 Gaucher disease. *Am J Hum Genet* 2000; **66**: 1777–1786.
- 19 Patterson M, Vanier MT, Suzuki K *et al*: Niemann-Pick disease type C: a lipid trafficking disorder; in Scriver SR, Beaudet AL, Sly WS, Valle D (eds): *The metabolic basis of inherited disease*. New-York: McGraw-Hill, 2001, pp 3611–3633.
- 20 Ribeiro I, Marcão A, Amaral O, Sá Miranda MC, Vanier MT, Millat G: Niemann-Pick type C disease: NPC1 mutations associated with severe and mild cellular cholesterol trafficking alterations. *Hum Genet* 2001; **109**: 24–32.
- 21 Osório, R Vilarinho L, Soares J, Almeida M, Carmona C, Martins E: Programa Nacional de Diagnóstico Precoce. *Arqui Med* 1999; **13**: 163–168.