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DIDASE A IN PATIENTS DISTAGE RENAL DISEASE

Sperandeo¹, R Tuzzi¹, G Sebastio¹, G Andria¹ phrology, Federico II nia

netabolism resulting from the manifestations of FD include: and cardiac and brain impairments, from 13 independent of suspected FD. Plasma activity was deficient, genomic DNA from these sis, when available. We have from 6 independent families: and for Fabry disease patients aring plasma a-galactosidase of them a-galactosidase A

MPTOMATIC FABRY

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its in Fabry carriers.

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are described as symptomatic beyte α-galactosidase levels ate increased the abnormal die or asymptomatic women to females. By contrast, skin

sed to confirm the diagnosis in female probands, is the

IDENTIFICATION AND CHARACTERIZATION OF ALPHA-GALACTOSIDASE A MUTATIONS IN CHINESE PATIENTS WITH FABRY DISEASE

MUTATIONS IN CHINESE PATIENTS WITH FABRY DISEASE M-Y Liu¹, C-H Chen³, S-J Wu⁴, Y-T Hsieh⁵, T-T Liu², K-J Hsiao^{1,2,4}

Institute of Genetics and ²Genome Research Center, Natl. Yang-Ming University, Taipei; ³Institute of Human Genetics, Tzu-Chi Collage of Medicine, Hualien; ⁴Dept. of Med. Research and Education, Taipei Veterans General Hospital; ⁵Dept. of Obstetrics and Gynaecology, Taichung Veterans General Hospital; Taiwan, R.O.C.

Fabry disease (MIM 301500) is an X-linked inborn error of sphingolipid catabolism caused by alphagalactosidase A (gene symbol: GLA, E.C. 3.2.1.22) deficiency. Four mutations, namely 274G > T (D92Y), 802_805 del (L286X), 781G > A (G261S), and a gross gene deletion, in the GLA gene were identified from four unrelated Chinese patients with Fabry disease in this study. The gross gene deletion, designated S65/sX5, was about 2.6 Kb in length and identified to comprise partial intron 1 extending to partial intron 2, which resulted in exon 2 skipping of the GLA transcripts. The G261S alteration and gross gene deletion were novel mutations in the GLA gene, while the D92Y and 802_805 del were reported previously. The G261S and D92Y mutants were constructed and expressed in a mammlian system to characterize the functional relevance of these two mutations. The relative activity of D92Y was below 1% of normal GLA construct and its kinetic properties could not be determined. The K_m and V_{max} value of G261S mutant were determined to be 2.4 ± 0.02 mM (mean \pm SE. n=3, normal: 2.5 ± 0.14 mM) and 28 ± 8.8 nmol/min/mg (normal: 37.7 ± 14.5 nmol/min/mg), respectively. The relative activity of G261S was about $70.2\pm16.9\%$ of normal construct, which correlated with the atypical clinical phenotype presentation of the patient. These results suggest that G261S and D92Y are disease causing mutations.

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THERAPY FOR FABRY DISEASE: COMPARISON OF AGALSIDASE ALPHA AND BETA ENZYME PREPARATIONS

GE Linthorst. D Blom. D Speijer. CEM Hollak. JMFG Aerts
Department of Internal Medicine and Biochemistry. Academic Medical Center. Amsterdam.
Meibergdreet 9, 1105. AZ Amsterdam. Netherlands

Fabry disease is an X-linked disorder caused by the deficiency of the lysosomal enzyme α -galactosidase A (α -Gal). Two α -galactosidase A enzyme preparations have been approved for enzyme supplementation therapy for Fabry disease in the EU: agalsidase alpha (ReplagalTM, TKT Inc) and agalsidase beta (FabrazymeTM. Genzyme Corp) that are produced by different methodologies, gene activation in human fibroblasts and cDNA transduction in CHO cells, respectively. Previously it has been reported that editing of α -Gal A mRNA may results in amino acid changes. Detailed biochemical analysis of both products was therefore performed. Enzyme activity per mg protein was similar (6.6 ± 0.6 and 7.0 ± 0.5 mmol mg hr. respectively). MS analysis gave no indications for amino acid changes due to RNA-editing in both preparations. Extensive analysis of RNA of normal controls did not reveal the existence of the reported editing process. Kinetic studies on the uptake in Fabry fibroblasts showed that both products were similarly taken up and functionally corrected glycosphingolipid storage. In addition, patients receiving either α -Gal preparation generated cross-reactive antibodies. These results form the basis for a clinical study which aims to compare the efficacy and safety of agalsidase alpha and agalsidase beta in a dose of 0.2 mg kg in adults with Fabry disease.

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