

TARGETED SEQUENCING OF HEXA GENE SHOWS MISSENSE SUBSTITUTION (P.ARG499HIS) IN A LARGE PAKISTANI FAMILY WITH TAY-SACHS DISEASE

F.K. BAZAI¹, M.U.I. HASSAN¹, H. TAYYAB², S. NAUDHANI³, S. SIRAJ⁴, M. TARIQ⁵, S.A. SHAH⁶, J. AHMAD⁷, S. DAUD^{3*}

¹ Quetta Institute of Medical Sciences, Quetta, Pakistan

² Department of Biochemistry and Biomedical Sciences, McMaster University, Ontario, Canada

³ Department of Biotechnology, BUIEMS, Quetta, Pakistan

⁴ Institute of Basic Medical Sciences, Khyber Medical University, Peshawar, Pakistan

⁵ Division of Science and Technology, University of Education, Lahore, Pakistan

⁶ Institute of Biotechnology and Microbiology, Bacha Khan University, Charsadda, Pakistan

⁷ Montreal Neurological Institute, McGill University, Montreal, Canada

E-mail: fariya_khan98@outlook.com, mehmoorkhan.mengal@gmail.com, tayyabh@mcmaster.ca, saranoudhani@yahoo.com, samisiraj.ibms@kmu.edu.pk, muhammad.tariq@ue.edu.pk, : sajjadbiotec@gmail.com, jamil.ahmad@mcgill.ca, shakeela.daud@buitms.edu.pk

*Corresponding Author: Shakeela Daud, PhD, e-mail: shakeela.daud@buitms.edu.pk

Tay-Sachs disease or GM2 gangliosidosis, is caused by a deficiency of beta-hexosaminidase A (HEXA), resulting in lysosomal accumulation of GM2 ganglioside. However, deficiencies or reduced activities of HEXA and HEXB result in Sandhoff disease. The patients manifest with the macular cherry-red spots due to lipid-laden ganglion cells, hypotonia, low muscle tone, intractable seizures, developmental arrest, blindness, and neurological deterioration. The aim of this study was to identify the TSD-causing variant in a large Pakistani family showing typical symptoms of Tay-Sachs disease. Here, we studied a large Pakistani family with six TSD patients for the identification of the pathogenic variant by targeted DNA sequencing. As a result, we identified a missense substitution (p.Arg499His) in exon 13 of HEXA that was completely cosegregated among affected and normal individuals. In conclusion, we identified a missense substitution (p.Arg499His) in HEXA gene in a large consanguineous Pakistani family and further enriched the mutational spectrum of HEXA through Pakistani patients for the early diagnosis of the disease.

Key words: HEXA, HEXB, TSD, GM2 gangliosidosis, β -N-acetylhexosaminidase A.

ЦІЛЬОВЕ СЕКВЕНУВАННЯ ГЕНУ HEXA ДЕМОНСТРУЄ МІСЕНС-ЗАМІНУ (P.ARG499HIS) У ВЕЛИКІЙ ПАКИСТАНСЬКІЙ РОДИНІ З ХВОРОБОЮ ТЕЯ-САКСА

Хвороба Тея-Сакса або GM2 гангліоліпідоз зумовлена дефіцитом бета-гексозамінідази А (HEXA), що призводить до лізосомного накопичення GM2 гангліозиду. Однак, недостатність або знижена активність HEXA і HEXB призводять до хвороби Сандхоффа. У пацієнтів з'являються такі симптоми, як вишнево-червоні плями на очних яблуках через накопичення ліпідів у гангліонарних клітинах, гіпотонія, слабкість м'язів, некеровані випадки, затримка розвитку, сліпота і погіршення неврологічних показників. Мета цього дослідження полягала в ідентифікації варіанту, який спричинив хворобу Тея-Сакса у великій пакистанській родині, яка демонструвала типові симптоми цієї хвороби і налічувала шість пацієнтів із хворобою Тея-Сакса. Ідентифікацію патогенного варіанту проводили за допомогою цільового секвенування ДНК. У результаті дослідження ми ідентифікували місенс заміну (p.Arg499His) в екзоні 13 HEXA, яка була повністю cosegregованою серед хворих та здорових осіб у великій єдинокровній пакистанській родині. Отже, ми ідентифікували місенс заміну (p.Arg499His) у гені HEXA і збагатили спектр мутацій HEXA серед пакистанських пацієнтів для ранньої діагностики захворювання.

Ключові слова: HEXA, HEXB, TSD, GM2 гангліоліпідоз, β -N-ацетилгексозамінідаза А.

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