

Case Study: The Role of Thyroid Function Tests in Diagnosing Allan-herndon-dudley Syndrome Revisited: A Novel Iran-based Mutation



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ABSTRACT

Introduction: Allan-Herndon-Dudley Syndrome (AHDS) is a rare X-linked recessive intellectual disability condition with neuromuscular involvements. Altered thyroid function tests are major milestones in AHDS diagnosis. However, due to phenotypic variations in the levels of thyroid hormones in AHDS patients, we believe that the disorder is often underdiagnosed. Here, we reported a 3.5-year-old boy with an AHDS diagnosis and healthy thyroid hormones.

Methods: Whole-Exome sequencing followed by data analysis was performed on the patient's sample. The mutation was confirmed by Sanger sequencing in the patient and his mother.

Results: We reported a 3.5-year-old boy with AHDS diagnosis and a novel synonymous missense mutation (c. 1026G>A) in the SLC16A2 gene manifesting normal levels of T3, T4, and TSH. The mutation causes no change in amino acid sequence; however, it affects splicing through alteration of an exonic splicing enhancer. To the best of our knowledge, there are only 3 similar reports in the literature reporting AHDS diagnosis and normal levels of thyroid hormones.

Conclusion: The altered levels of thyroid hormones are notable but not necessary markers for diagnosing AHDS. The candidate diagnosis of AHDS should be considered in patients with X-linked recessive intellectual disability syndrome with neuromuscular involvements irrespective of levels of thyroid hormones; otherwise, it could lead to the under-diagnosis of the disorder.

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Highlights

- A novel mutation (c. 1026G>A) in the SLC16A2 gene causes AHDS.
- The mutation causes no change in amino acid sequence and it only affects splicing.
- Patients with this mutation manifest normal levels of T3, T4, and TSH.

Plain Language Summary

Allan-Herndon-Dudley Syndrome (AHDS) is a rare intellectual disability condition with neuromuscular involvement. Altered thyroid function tests are significant milestones in AHDS diagnosis. However, due to phenotypic variations in the levels of thyroid hormones in AHDS patients, we believe that the disorder is often underdiagnosed. Here, we reported a 3.5-year-old boy with an AHDS diagnosis and healthy thyroid hormones. The DNA sequencing analysis showed that the mutation causes no immediate change in amino acid sequence; however, it affects protein splicing. To the best of our knowledge, there are only three similar reports in the literature reporting AHDS diagnosis and normal levels of thyroid hormones. We conclude that the altered levels of thyroid hormones are notable but not necessary markers for diagnosing AHDS. The candidate diagnosis of AHDS should be considered in patients with X-linked recessive intellectual disability syndrome with neuromuscular involvements irrespective of levels of thyroid hormones; otherwise, it could lead to the under-diagnosis of the disorder.

1. Introduction

Allan-Herndon-Dudley Syndrome (AHDS) is an X-linked recessive intellectual disability condition, characterized by muscular hypoplasia, hypotonia, spastic paraplegia (Dumitrescu, Liao, Best, Brockmann, & Refetoff, 2004). AHDS is a rare disease with an overall prevalence of less than one in a million. The disease manifestations appear in the neonatal period or early infancy.

AHDS is caused by mutations in the SLC16A2 gene (Xq13.2). This gene, expressed in the brain, encodes for Monocarboxylate Transporter 8 (MCT8) and is a specific transporter of thyroid hormone T3 into nerve cells (Schwartz et al., 2005). Gene mutations altering the structure and function of the SLC16A2 protein effectively disrupt T3 transport into nerve cells; thus, it disrupts normal brain development, resulting in intellectual disability and problems with movement. Moreover, T3 is not taken up by nerve cells; therefore, excess amounts of this hormone continue to circulate in the bloodstream and is a hallmark of AHDS diagnosis.

Diagnosis is based on clinical manifestations, delayed myelination of the brain on MRI radiological examination, and the presence of altered thyroid function tests. Affected males have abnormally high T3 levels, low to normal T4 levels, and normal to slightly elevated TSH levels. Diagnosis is confirmed by molecular genetic test-

ing revealing mutations in the SLC16A2 gene. As characteristics of X-linked recessive inheritance, the female can pass on the mutated gene; however, it usually does not experience signs and symptoms of the disorder. The carriers of SLC16A2 mutations have normal thyroid function tests and healthy intelligence and do not experience movement problems. In this study, we reported a 3.5-year-old boy with AHDS diagnosis and a novel synonymous missense mutation in the SLC16A2 gene manifesting normal levels of T3, T4, and TSH.

2. Case Presentation

Medical examination and genetic counseling: A 3.5-year-old boy was referred to our clinic for evaluating floppiness. He was also under the observation of a pediatric neurologist because of cerebral palsy. He was born from unrelated parents (Figure 1); delivery occurred in 38 weeks, with a birth weight of 2.9 kg and length of 50cm. His progressive weakness was discovered a few months after birth. On our examination, he manifested a failure to thrive; his weight was 11 kg (< the 5th percentile of corresponding age) and his length was 87cm (<the 5th percentile of corresponding age). The head circumference was 47.5 cm (equal to the 50th percentile). He manifested excessive drooling, obvious head lag, and global developmental delay implying serious neurological damage. He could follow objects and pay attention to sounds but was unable to walk, speak, and feed independently. Other neuromuscular findings involved generalized hypotonia,

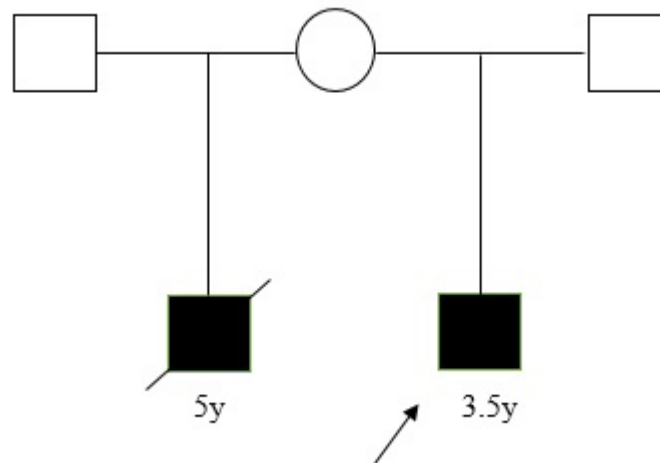


Figure 1. The pedigree of the family

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Proband is shown with a black arrowhead The patients with clinical AHDS manifestations, including the proband, are shown in black.

the hypotonia of limbs, and increased deep tendon reflexes. Genitalia was prepubertal and testes were descended and palpable. Laboratory tests revealed normal metabolic and lipid profile, normal liver function tests, and normal levels of free triiodothyronine (T3) (3.87 pg/mL within the normal range of 3.7-4.4 pg/mL), free thyroxine (T4) (1.23 ng/dL within the normal range of 1.1-1.4 ng/dL) and TSH (2.34 mIU/L within the normal range of 1.75-3.5 mIU/L). Auditory Brainstem Response (ABR) test data revealed no abnormal latency time; however, brain MRI manifested remarkable delayed myelination of the brain white matter. The mother disclosed that from another marriage, she had another son with similar phenotypes who had died at the age of 5 years.

Whole exome sequencing: This familial occurrence suggested an X-linked recessive disorder; however, because AHDS was not suspected before analysis, exome sequencing was performed. After obtaining our Institutional Ethical Committee's approval and receiving written informed consent from the patient's family, the patient's blood sample was obtained and genomic DNA was extracted. Clinical exome sequencing was performed using the Illumina HiSeq4000 sequencing panel.

3. Results

A novel single nucleotide alteration in exon 3 (c.1026G>A) was considered as the mutation related to

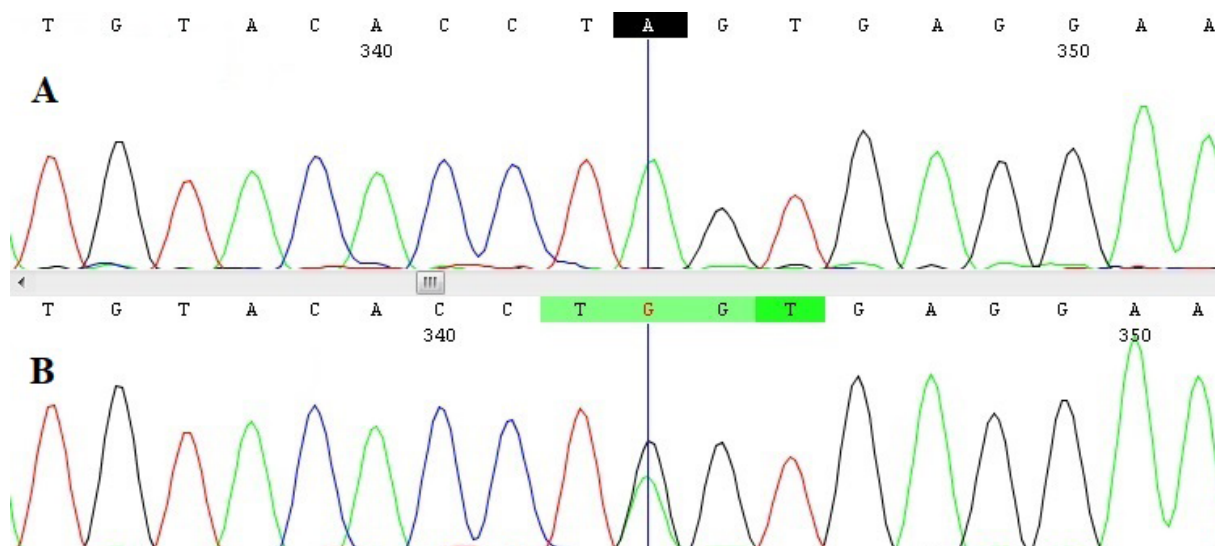


Figure 2. Chromatogram of the c. 1026G>A (p. L342L) mutation

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A: Sanger confirmation of c. 1026G>A mutation in the proband and B: The heterozygous mother.

the patient's condition. The mutation does not cause any changes in the amino acid sequence (p. L342L). This variant is not reported in the 1000 genomes database and ExAC browser. Using Mutation Taster, the variant was predicted as damaging. Using HSF (Human Splicing Finder), the mutation was predicted to alter a wild-type splicing donor site, most probably affecting splicing through the alteration of an Exonic Splicing Enhancer (ESE). This mutation was confirmed in the patient by Sanger sequencing (Figure 2A). This mutation was verified in the patient's mother (Figure 2B). Therefore, the diagnosis of AHDS was confirmed in the family.

4. Discussion

The AHDS is a rare X-linked recessive intellectual disability condition. The diagnosis is primarily based on clinical findings and altered thyroid hormone levels. Although the increased T3 levels are mentioned as an obligate finding in some studies (Friesema, Visser, Visser, & Endocrinology, 2010; Langley, Trau, Bean, Narravula, & Schrier Vergano, 2015), to the best of our knowledge, 3 studies are supporting the normal levels of thyroid hormones in AHDS patients (Boccone, Dessi, Meloni, & Loudianos, 2013; Shimojima et al., 2016; Tsurusaki et al., 2011). In this study, we reported a 3.5-year-old boy manifesting AHDS and normal thyroid function tests with a missense mutation in the SLC16A2 gene affecting splicing. In contrast to our finding, other studies reported different non-synonymous variants, one in exon 3 similar to our finding.

The reason why some patients with AHDS manifest normal thyroid hormones remains unclear. A previous study indicated a single mutation in the SLC16A2 gene in two males in a family caused phenotypic variability, as follows: the infant male showed increased T3 levels and the adult male presented normal T3 levels (Boccone et al., 2013). Boccone et al. discussed that this variable expressivity could probably be attributed to the age of the patients. From a genetic point of view, variable expressivity refers to the range of signs and symptoms that can occur in different individuals with the same genetic condition, i.e., the same pathogenic mutation in the SLC16A2 gene. Variable expressivity is probably caused by a combination of genetic (interactions with other loci), age, environmental, and lifestyle factors, most of which remained unidentified. Variable expressivity of a genetic disease makes it challenging to diagnose.

Clinical considerations: The normal thyroid hormones profile in some AHDS patients implies clinicians should consider AHDS as a candidate diagnosis in patients with

X-linked recessive intellectual disability syndrome with neuromuscular involvements irrespective of the levels of thyroid hormones. Screening for mutations in the SLC16A2 gene in these cases obviates the need for expensive Whole Exome Sequencing (WES). This is particularly important in developing countries where WES is costly due to limited resources. Furthermore, this screening is essential from the view of genetic counseling. Since there is no treatment available for this disease and the disease compromises the quality of life by affecting the ability of the patient to live independently; genetic counseling to prevent the birth of an affected child is of significance. Affected families should be informed that prenatal or preimplantation diagnosis of a male with AHDS is possible if the mutation in his mother is identified. In our case, the ultimate diagnosis of AHDS in one infant uncovered his previously undiagnosed affected brother who had expired of the disease. If the diagnosis of AHDS was considered in the first brother, it could have led to prenatal or preimplantation diagnosis to prevent the birth of a second affected child.

5. Conclusion

In brief, altered levels of thyroid hormones are a notable but not necessary marker for the diagnosis of AHDS. Due to phenotypic variations in the levels of thyroid hormones in AHDS patients, the disorder is often under-diagnosed imposing economic and mental burden. Therefore, the candidate diagnosis of AHDS should be considered in patients with X-linked recessive intellectual disability syndrome with neuromuscular involvements irrespective of levels of thyroid hormones.

Ethical Considerations

Compliance with ethical guidelines

All ethical principles are considered in this article. The participants were informed of the purpose of the research and its implementation stages. They were also assured about the confidentiality of their information. They were free to leave the study whenever they wished, and if desired, the research results would be available to them. Written consent has been obtained from the subjects. Principles of the Helsinki Convention were also observed.

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Authors' contributions

All authors equally contributed to preparing this article.

Conflict of interest

The authors declared no conflict of interest.

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