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De novo mutation of *NAXE (APOAIBP)*-related early-onset progressive

encephalopathy with brain edema and/or leukoencephalopathy-1: A case report and

literature review

Le Ding, Ting-Ting Huang, Guohuan Ying, Shang-Yu Wang, Hai-Feng Xu, Hao Qian,

Faiza Rahman, Xiaopeng Lu, Hu Guo, Guo Zheng, Gang Zhang

Abstract

BACKGROUND

Early-onset progressive encephalopathy with brain edema and/or

leukoencephalopathy-1 (PEBEL1) is a rare autosomal recessive severe neurometabolic

disease. The aim of this study was to investigate the clinical characteristics and genetic

pathogenicity of PEBEL1 caused by rare NAXE (or APOA1BP)-related defects.

CASE SUMMARY

The patient was a girl aged 2 years and 10 mo. She was hospitalized due to walking

disorder for > 40 d. The clinical manifestations were ataxia, motor function regression,

hypotonia, and eyelid ptosis. Within 1 mo of hospitalization, she developed sigh

breathing, respiratory failure, cerebellar edema and brain hernia, and finally she died.

Changes were found in cranial imaging, including cerebellar edema accompanied by

symmetrical myelopathy. Through whole exome sequencing, we detected

NAXE compound heterozygous variation (NM 144772.3) c.733A>C (p. Lys245Gln,

dbSNP: rs770023429) and novel variation c.370G>T (p.Gly124Cys) in the germline gene.

The clinical features and core phenotypes of this case were consistent with 18

previously reported cases of PEBEL1.

CONCLUSION

This is the first case of NAXE-related PEBEL1 with severe clinical phenotype in Mainland China. The p.Gly124Cys mutation discovered in this case has enriched the pathogenic variation spectrum of *NAXE*.

INTRODUCTION

Early-onset progressive encephalopathy with brain edema and/or leukoencephalopathy 1 (PEBEL1, OMIM: #617186) is a rare autosomal recessive severe neurometabolic disease^[1]. At present, more than 10 cases have been reported worldwide since 2016^[1-3], but only one case has been reported in China. In this study, the clinical data of a child with PEBEL1 treated in the Department of Neurology, Children's Hospital Affiliated to Nanjing Medical University in July 2021 were retrospectively analyzed, and the related literature was reviewed to improve the understanding, diagnosis, and treatment of the disease.

CASE PRESENTATION

Chief complaints

The patient was a girl aged 2 years and 10 mo. She was hospitalized because of walking disorder for > 40 d.

History of present illness

Forty days ago, she presented with unstable walking and was suspected of having synovitis and stayed in bed at home. She had an episode of fever 30 d prior to that, and the fever spike was 38°C. Blood examination in the outpatient clinic showed leukocytosis, and the fever subsided after oral administration of cephalosporin for 1 wk. During this period, the patient did not leave bed or walk again. Two weeks ago, she developed binocular movement disorder, slow eyeball pursuit, and slight drooping

eyelids. Gradually, the patient had wobbling in sitting, and resisted sitting, and cried when urinating, so she was admitted to hospital.

History of past illness

There was a sudden suspicious episode of choking during a meal 3 mo ago, with gaze, unconsciousness, and clenched teeth for 6–7 min, which was relieved after patting on the back and vomiting. There was no abnormality in computed tomography (CT) examination of the head and chest in the local hospital. She was scratched by a cat 2 mo ago and was injected with rabies vaccine three times in 2 wk. Fifty days ago, she suddenly developed generalized weakness and bowed her head during playing, and then her limbs became stiffer, which lasted for about 1 min, followed by vomiting and incontinence. The follow-up was as usual, and the parents made an appointment for a video electroencephalogram examination, which was not done because the patient was immobilized at home.

Personal and family history

There was no special personal history, and the milestone of intellectual and motor development was normal since childhood. Her parents were in good health and denied any consanguinity. The 11-year-old brother also was in good health.

Physical examination

There was no special personal history, and the milestone of intellectual and motor development was normal since childhood. Her parents were in good health and denied any consanguinity. The 11-year-old brother also was in good health.

Laboratory examinations

After admission, blood routine tests were done. Biochemical studies, blood ammonia, blood lactic acid, erythrocyte sedimentation rate, procalcitonin, six items of coagulation, complete set of autoantibodies, cellular immunity, humoral immunity, four items of

infectious diseases, and seven items of thyroid function tests were normal. Screening of hereditary metabolic diseases in hematuria was not abnormal. Four items of tumor: nonspecific enolase was 40.33 ng/mL (0–16.3 ng/mL), a-fetoprotein, carcinoembryonic antigen and carbohydrate antigen 19-9 were normal. Cerebrospinal fluid biochemistry showed: protein 0.27 g/L and glucose 4.82 mmol/L. Routine cerebrospinal fluid analysis showed: nucleated cell count 3×10⁶/L; cerebrospinal fluid-MP, EB, cytomegalovirus and herpes simplex virus DNA was negative; cerebrospinal fluid and blood autoimmune encephalitis antibody, central demyelination antibody, and peripheral nerve disease spectrum antibody were negative; and cerebrospinal fluid pathogen macro gene was negative.

Imaging examinations

Plain CT of the chest and abdomen showed no abnormalities, with normal abdominal contents and obvious bladder filling. No abnormality was found in skull magnetic resonance imaging (MRI), MR angiography and MR venography on day 2 after admission, and spinal MRI showed T2 high signal in the spinal cord at the T7–10 Level (poor coordination, heavy artifact) (Figure 3). Electroencepholography showed background moderation, poor rhythmicity and responsiveness, with spike waves and spike slow waves in bilateral anterior and middle temporal regions during sleep, which were not synchronized between left and right.

Further diagnostic work-up

With the approval of the Medical Ethics Committee of the hospital (approval number: 202111116-1) and the informed consent of the guardians of the child, 2 mL peripheral blood from the patient and her parents were taken for whole exome sequencing (Beijing Zhiyin Dongfang). The whole-exome library was constructed using xGen ®Exome Research Panel v1.0 (IDT, USA) capture probe, and the NovaSeq 6000 (Illumina, USA) series sequencer was used for high-throughput sequencing. The compound heterozygous mutations c.733A > C (p.Lys245Gln, dbSNP: rs770023429) and c.370G > T

(p.Gly124Cys) in NAXE (APOA1BP) inherited from her parents were identified and verified by Sanger sequencing (Figure 1). According to the American Society of Medical Genetics 2015 [3], Lys245Gln (reported in the PS1+PM1, gnomAD East Asia MAF:0.0019, PEBEL1 case [4]) was likely pathogenic and Gly124Cys (PM1+PM2+PP3, no MAF record) was annotated as unknown pathogenicity variation. The variant amino acid residues were analyzed using VarSite (https://www.ebi.ac.uk/thorntonsrv/databases/VarSite); a variant analysis tool provided by European Institute of Bioinformatics, and the two variants were both highly evolutionarily conservative (Figure 2). The conservatism of Lys245Gln in 174 homologous sequences was 0.5, which was lower than that of Gly124Cys in 190 homologous sequences (0.1). Using SWISS-MODEL and Swiss-pdb Viewer software to predict the pathogenicity of the mutation sites, the protein structure, the residues of the mutation site, and the functional sites nearby, it is suggested that the possible pathogenicity, c.733A>C site is shown in Figure 6, c.370G>T mutation site is shown in Figure 7.

FINAL DIAGNOSIS

The patient was diagnosed with PEBEL1.

TREATMENT

After admission, a detailed examination was done and the patient was treated with acyclovir, dexamethasone (0.5 mg/kg), gammaglobulin (2 g/kg), and other anti-inflammatory drugs. Urinary retention occurred on the day 3 and indwelling catheterization was performed. On day 5, the patient developed confusion, irregular respiratory rhythm, and poor response, and was transferred to the pediatric intensive care unit. This was followed by a decrease in heart rate and blood oxygen saturation, and she received cardiopulmonary resuscitation and mechanical ventilation. Nutritional support with levocarnitine and coenzyme Q10 was given, and simultaneous high-dose methylprednisolone (20 mg/kg) was given for 3 d, and plasma exchange was performed for 2.5 h on day 10. Bilateral pupil dilation, the disappearance of light reflex

and deep coma appeared on day 12. Cranial CT suggested diffuse cerebellar swelling with hydrocephalus and ventricular dilatation, and further improvement of cranial CT angiography enhancement suggested an unclear display of straight sinus and no significant parenchymal enhancing mass shadow was seen (Figure 4). On day 14, after multidisciplinary team discussion, intraventricular drilling and drainage, and intracranial pressure probe implantation were performed. About 100 mL cerebrospinal fluid was drained after surgery, and the intracranial pressure was maintained at 20-74 mmHg. On day 17, whole exosome sequencing detected NAXE compound heterozygous mutation, and we added 100 mg nicotinamide intravenous drip. The patient gradually deteriorated and developed a slow heart rate, hypotension, central diabetes insipidus and electrolyte disturbance, and was treated with symptomatic support such as volume expansion, plasma, albumin, electrolyte supplement, blood transfusion, and maintenance of blood pressure using dopamine, norepinephrine, epinephrine, and posterior pituitary hormone. On day 26, cerebral MRI showed cerebellar swelling, possible herniation of the cerebellar curtain notch, and multiple abnormal signals in the brain parenchyma and cervical spinal cord (Figure 5).

OUTCOME AND FOLLOW-UP

After 30 d, the child became brain dead and her parents abandoned treatment.

DISCUSSION

PEBEL1 is a rare fatal encephalopathy caused by a double allele mutation of *NAXE* (*APOA1BP*) on chromosome 1q22. In 2016, Spiegel [2] reported for the first time that five children from Israel, who were near relatives were affected. The age at onset was 6–12 mo, with loss of motor function after infection, bedridden at the age of 2 years, mechanically ventilated, and finally in a vegetative state. Four patients died between 1 and 3 years of age, and one child was supported by a ventilator until 5.5 years of age, and MRI revealed deep white matter lesions. Kremer *et al* [1] summarized the clinical features of five patients who presented with infantile/early childhood onset, usually

caused by fever with rapidly progressive deterioration of neurological function. The patients showed muscular hypotonia, motor development regression, cognitive loss, ataxia, nystagmus, seizures, quadriplegia, and respiratory failure, which eventually led to a vegetative state and brain death. Brain and spinal cord imaging may show white matter abnormalities, cerebral atrophy, cerebellar edema, and myelopathy. It has also been reported [1] that a large area of subacute bullous dermatosis occurred within a few weeks after the onset of neurological symptoms.

The keywords NAXE, APOAIBP, NAD(P)HX epimerase and AIBP were searched in PubMed, ClinVar, Wanfang, and CNKI, and papers with clear clinical data were selected. As of October 2021, 17 reported pathogenic NAXE variants in 12 cases had been collected (Table 1). These included 11 missense mutations (Glu85Asp, Ala94Asp, Gly121Arg, Arg129Pro, Gly189Ser, Ile214Val, Ile214Ser, Asp218Asn, Asp218Val, Lys245Gln and Gly253Ser, 64.7%, 11/17), two splice site mutations (c. 516+1G > A and 665-1G > A, 11.8%, 2/17), two nonsense mutations (Tyr59Ter Gln66Ter,11.8%,2/17), one frameshift mutation (Ala248fs, 5.9%, 1/17), and one deletion/insertion mutation (c.804_807delinsA/Lys270del, 5.9%, 1/17) [1, 2, 4-7]. The Lys245Gln detected in the present case was the reported homozygous pathogenic variant [4]. The clinical characteristics of the present case and 18 previous cases from 12 families are summarized as follows [1-8]. The age of onset of PEBEL1, except for a German family with two girls who developed the disease at the age of 20 and 22 years, respectively [4], was < 3 years in 89% of the patients (17/19). Most cases were induced by febrile infection (58%, 11/19). The main manifestations were rapidly progressive or recurrent respiratory insufficiency (79%, 15/19), motor cognitive regression (79%, 15/19), decreased muscle strength and muscle tone (79%, 15/19), ataxia (42%, 8/19), nystagmus (32%, 6/19), strabismus (21%, 4/19), seizures (21%,4/19), bilateral blepharoptosis (11%, 2/19), dysarthria and dysphagia (21%, 4/19), and skin erythema rash (11%, 2/19). Eleven of the 19 patients had brain MRI changes, including brain edema (55%, 6/11), brain white matter abnormalities (27%, 3/11), brain atrophy (27%, 3/11), myelopathy (27%, 3/11), cortical and basal ganglia lesions (10%, 1/11), and brain

stem and intracranial hemorrhage (10%, 1/11). The cerebrospinal fluid lactic acid test was elevated in seven of the 19 cases, (71%, 5/7), and 12 of 19 cases died of end-stage coma within 3 years of age (63%, 12/19).

In the present case, the patient had normal development since childhood and had a history of rabies vaccination in the early stage of onset, and the nervous system symptoms were gradually aggravated after fever. It was possible to be misdiagnosed as an immune inflammatory disease; however, the condition still deteriorated rapidly after active immunotherapy, with respiratory failure, cerebellar edema, and cerebral herniation, which posed major challenges to clinicians. Carefully combing the medical history, we found that the patient had two paroxysmal events within 3 mo of PREBLE1 onset, which were suspected to be epileptic seizures. Physical examination on admission showed that the patient had droopy eyelids, eye movement disorder, and hypotonia of the extremities, and the pathological reflex was positive and the spinal cord lesions were symmetrical. Due to the rapid change of the disease, electromyographic examination was not conducted, and it was necessary to be alert to the possibility of hereditary metabolic diseases (mitochondria), so gene detection was performed as soon as possible. The clinical features, imaging findings, and disease progression of the child were consistent with the characteristics of PEBEL1.

The human nicotinamide nucleotide repair system consists of two chaperone enzymes: NAD(P)HX differential isomerase (NAXE, formerly known as APOA1BP, OMIM:608862), which converts R-NAD(P)HX to S-NAD(P)HX, and the NAD(P)HX dehydratase (NAXD, formerly CARKD, OMIM:615910), which converts SNAD(P)HX back to NAD(P)H [9] in an ATP-dependent manner. The presence of NAD(P)HX repair enzymes in all tissues and species, coupled with the core metabolism of their cofactors which play a protective role, suggests that the repair system is essential for life maintenance^[10].

The NAXE (APOAIBP) gene encodes NAD(P)HX differential isomerase or apolipoprotein AI binding protein (AIBP), located on the 1q22 chromosome. There are five transcripts, including six exons (https://www.uniprot.org/uniprot/Q8NCW5),

spanning 2.5 kb and composed of 288 amino acids. It contains mitochondrial transport peptide (residues 1-47), NAD(P)H hydrate isomerase (residues 48-288), yjefn terminal domain (residues 65-275), two NAD(P)HX region (residues 119-223, residues 189-195). NAXE protein exists in cerebrospinal fluid and urine, and is widely expressed in the kidneys, heart and liver [11]. In in vitro experiments in fibroblasts from PEBEL1 patients, NAXE deficiency caused a significant increase in circulating NADHX and two NADHX isoforms (S and R isoforms) under normal body temperature (37°C). The elevated levels of circulating NADHX were greater in NAXE-deficient cells after treatment with heat stress (40°C), and the elevation of S- and R-type NADHX isoforms was attenuated. In addition, the activity of intact mitochondria in the muscle tissue of two patients also decreased significantly [1]. These prove that NAXE deficiency leads to the accumulation of the toxic metabolites of the biochemical reaction cofactors NADH and NADPH, circulating NADHX, then affects the oxidative phosphorylation of mitochondria and leads to the deficiency/disorder of mitochondrial energy metabolism. Therefore, it can be predicted that the brain with high demand for energy supply produced by mitochondria is particularly vulnerable to the damage of NAD(P)HX repair, which can also explain the deterioration of neurological symptoms clinically in children after fever. Although NAXE is highly expressed in the human brain (https://gtexportal.org/home/gene/NAXE), the mechanism of its defect in the central nervous system is still unclear.

Vitamin B3 and coenzyme Q are specific treatments for elevated NAD+ levels, so they are also expected to benefit PEBEL1 patients. A 22-year-old female patient reported by TRINH [4] in Germany, in addition to receiving vitaminB3 treatment (niacin 40 mg bid to 2×40 mg bid), received anticonvulsant therapy (topiramate 50 mg bid, clomazone 20 mg/d, lamotrigine 300 mg/d, levetiracetam 1000 mg/d, piracetam 2400 mg/d, vitamin D3, esomeprazole, and laxative). The patient survived up to 29 years of age and had significant improvement in spasticity with some recovery in motor abilities. In our case, the patient had acute onset and progressed rapidly. Although nicotinamide (vitamin B3) and coenzyme Q were used on day 17 of onset, they still could not

reverse/delay brain edema, and the fatal pathological changes need to be proved by practice.

CONCLUSION

This is the first case of severe clinical phenotypic PEBEL1 reported in Mainland China, and a compound heterozygous variation of the *NAXE* gene was detected and a new variant Gly124Cys was identified, which expanded the pathogenicity variation spectrum of PEBEL1. Due to the rapid progress of PEBEL1 and mostly sporadic cases, its early presentation, often characterized by acute onset of fever-induced deterioration of central nervous function or even cerebral edema, is not easy to indicate hereditary diseases, which may be an important reason for the lack of awareness of the disease among clinicians. For patients suspected of PEBEL1, vitamin B3 and coenzyme Q therapy should be tried first, and gene detection should be carried out as soon as possible to make a definite diagnosis. For the family of this case, we suggest that the *NAXE* mutation test should be used as a necessary part of prenatal genetic screening.

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