CLINICAL VIGNETTE



Recurrent status epilepticus and severe bifrontal hypometabolism in PGAP1-related neurodevelopmental disorder

Samia Benabess¹ | Kenneth A. Myers^{2,3,4} ©

Correspondence

Kenneth A. Myers, Montreal Children's Hospital, McGill University Health Centre Glen Site, 1001 Boulevard Décarie, Montreal, PQ H4A 3J1, Canada.

Email: kenneth.myers@mcgill.ca

Funding information

Fonds de Recherche du Québec - Santé, Grant/Award Number: 282228 and 295639

KEVWORDS

congenital diaphragmatic hernia, focal epilepsy, PGAP1, status epilepticus

Glycophosphatidylinositol (GPI) plays an anchoring role, linking cell membranes to proteins. GPI biosynthesis involves >20 proteins, including phosphatidylinositol glycans (PIGs) and post-GPI attachment to proteins (PGAPs).¹ Pathogenic variants in genes encoding PIGs and PGAPs are associated with global developmental impairment and congenital malformations. PGAP1 encodes an enzyme involved in GPI biosynthesis through the catalysis of GPI inositol deacylation.² Eight patients with PGAP1-related disorders have been described from five families, all with biallelic apparent loss-of-function variants.³⁻⁷ The clinical phenotype involves severe to profound developmental impairment, with spastic quadriparesis, feeding problems, microcephaly, cerebral visual impairment, dyskinesia, and brain atrophy variably reported. Seizures were only reported in 2/8.^{4,5} We present two brothers with PGAP1related disorder, including the proband with recurrent status epilepticus and severe bifrontal positron emission tomography (PET) hypometabolism.

A Pakistani male was born at term via caesarean section following a pregnancy complicated by gestational diabetes, preeclampsia, polyhydramnios, and antenatal diagnosis of congenital diaphragmatic hernia (CDH; Figure 1D). CDH repair was done at 72h of life and was uncomplicated.

At age 6 months, recurrent status epilepticus developed with seizures involving unresponsiveness, writhing, and unusual eye movements, lasting >60 min and requiring emergency medication to stop. In adolescence, seizure semiology changed to involve tachycardia, unilateral head and eye deviation, and eventually bilateral tonic-clonic convulsions. These events lasted ~2 min but occurred in clusters that almost always required emergency medications, such as diazepam, midazolam, or phenytoin, to stop. This pattern continued for many years despite the sequential addition of clonazepam, clobazam, lamotrigine, and levetiracetam. The family used rectal diazepam 20 mg as initial status epilepticus treatment before calling an ambulance. Seizure severity improved in late adolescence on lamotrigine 200 mg bid, levetiracetam 1500 mg bid, clobazam 10 mg morning, 35 mg evening, and clonazepam 1.5 mg morning, 2.5 mg evening. When last seen at age 19 years, he had ~1 seizure/month but no longer had clusters or required rescue medication.

This is an open access article under the terms of the Creative Commons Attribution-NonCommercial-NoDerivs License, which permits use and distribution in any medium, provided the original work is properly cited, the use is non-commercial and no modifications or adaptations are made. © 2025 The Author(s). Epileptic Disorders published by Wiley Periodicals LLC on behalf of International League Against Epilepsy.

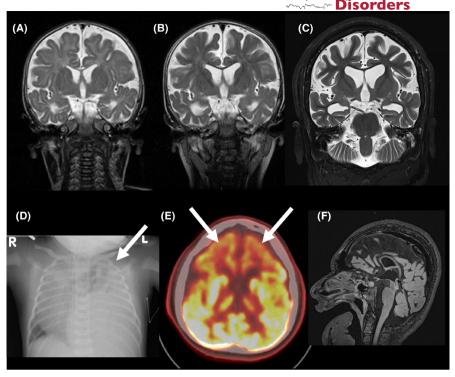
¹Faculty of Medicine and Health Sciences, McGill University, Montreal, Quebec, Canada

²Research Institute of the McGill University Health Centre, Montreal, Quebec, Canada

³Division of Neurology, Department of Pediatrics, Montreal Children's Hospital, McGill University Health Centre, Montreal, Quebec, Canada

⁴Department of Neurology and Neurosurgery, McGill University Health Centre, Montreal, Quebec, Canada

FIGURE 1 Brain atrophy in the proband is demonstrated with coronal T2 brain MRI showing progressive brain volume loss in images taken at (A) 5 months, (B) 18 months, and (C) 15 years of age. In (D), a chest x-ray taken at ~24h of life shows a left congenital diaphragmatic hernia. Gas-containing bowel loops can be seen in the left hemithorax (arrow). The left lung is not seen, and the right lung is compressed by the left diaphragmatic hernia. In (E), brain PET at age 15 years shows severe hypometabolism in the frontal regions (arrows). The cerebellum was relatively unaffected by atrophy, as shown in the sagittal FLAIR sequence at age 15 years (F).



Past medical history was notable for sialorrhea, constipation, scoliosis, right orchidopexy, and dysphagia requiring gastrostomy tube insertion. He had obstructive sleep apnea requiring tonsillectomy/adenoidectomy but still required bilevel positive airway pressure treatment. He had cortical visual impairment and profound global developmental impairment with spastic quadriparesis. He never walked, could not hold objects, was nonverbal, and did not show a clear capacity to understand words or gestures.

His parents were first cousins from Pakistan and had two other sons, one healthy with normal intelligence and a second with a phenotype very similar to the proband, including profound intellectual disability (ID), epilepsy, and microcephaly (but not CDH).

On examination, the proband had deep-set eyes and a high-arched palate. Appendicular tone was increased, and deep tendon reflexes were diffusely brisk. Brain MRI showed progressive volume loss (Figure 1A-C). Brain PET showed severe bifrontal hypometabolism (Figure 1E). EEG typically showed mildly slow background with multifocal spikes and sharp waves. During prolonged video EEG at age 15 years, focal seizures were recorded, originating independently from either hemisphere. A clinical gene panel (Blueprint Genetics) identified a novel homozygous intragenic *PGAP1* deletion, c.(1861+1 1862+2) (1952+1_1953-1)del, estimated to cover the region chr2:197712564-197712864, affecting exon 21; however, exact breakpoints could not be determined. Both parents were heterozygous for the deletion, and the proband's brother with profound ID was also homozygous. The

deletion is classified as pathogenic by ACMG criteria due to absence from control databases (PM2), predicted null variant with loss of function effect (PVS1), and familial segregation (PP1).⁸

This report clarifies the epilepsy phenotype that may arise with PGAP1 pathogenic variants and demonstrates that severe frontal hypometabolism can occur. While this finding indicates severe bilateral frontal lobe dysfunction, the underlying cause is unclear. The findings may also extend the phenotypic spectrum for PGAP1-related disorders to include CDH. The latter is unlikely to be coincidental, given that CDH is rare (1 in 3000 live births).^{9,10} CDH involves incomplete diaphragm development with consequent herniation of abdominal viscera into the chest cavity. It requires urgent medical intervention at birth, as the consequent respiratory distress is life-threatening. Both genetic and environmental factors are believed to play a role in CDH. In addition to copy number variants, CDH has been associated with at least 16 genes, 11 including some in the GPI-anchoring pathway including PIGA, PIGW, PIGL, PIGV, and PIGN. 12-17

AUTHOR CONTRIBUTIONS

Samia Benabess: Data collection, writing – original draft. Kenneth A. Myers: Conceptualization, data collection, preparation of figures, writing – review and editing.

ACKNOWLEDGMENTS

We thank the patient and his family. This study was supported by funding from Fonds de Recherche du Québec – Santé.



FUNDING INFORMATION

This study was supported by funding from the Fonds de Recherche du Québec – Santé (282228, 295639).

CONFLICT OF INTEREST STATEMENT

SB has no relevant disclosures. KAM is a site principal investigator for studies sponsored by Ultragenyx and LivaNova, and is a member of advisory boards for Jazz Pharmaceuticals and AS²Bio.

DATA AVAILABILITY STATEMENT

The data that support the findings of this study are available from the corresponding author upon reasonable request.

PATIENT CONSENT STATEMENT

Written consent for publication was obtained from the patients' parents.

ORCID

Kenneth A. Myers https://orcid.org/0000-0001-7831-4593

REFERENCES

- 1. Wu T, Yin F, Guang S, He F, Yang L, Peng J. The glycosylphosphatidylinositol biosynthesis pathway in human diseases. Orphanet J Rare Dis. 2020;15(1):129.
- Tanaka S, Maeda Y, Tashima Y, Kinoshita T. Inositol deacylation of glycosylphosphatidylinositol-anchored proteins is mediated by mammalian PGAP1 and yeast Bst1p. J Biol Chem. 2004;279(14):14256–63.
- 3. Bosch DG, Boonstra FN, Kinoshita T, Jhangiani S, de Ligt J, Cremers FP, et al. Cerebral visual impairment and intellectual disability caused by PGAP1 variants. Eur J Hum Genet. 2015;23(12):1689–93.
- 4. Kettwig M, Elpeleg O, Wegener E, Dreha-Kulaczewski S, Henneke M, Gärtner J, et al. Compound heterozygous variants in PGAP1 causing severe psychomotor retardation, brain atrophy, recurrent apneas and delayed myelination: a case report and literature review. BMC Neurol. 2016;16:74.
- 5. Murakami Y, Tawamie H, Maeda Y, Büttner C, Buchert R, Radwan F, et al. Null mutation in PGAP1 impairing Gpi-anchor maturation in patients with intellectual disability and encephalopathy. PLoS Genet. 2014;10(5):e1004320.
- Granzow M, Paramasivam N, Hinderhofer K, Fischer C, Chotewutmontri S, Kaufmann L, et al. Loss of function of PGAP1 as a cause of severe encephalopathy identified by whole exome sequencing: lessons of the bioinformatics pipeline. Mol Cell Probes. 2015;29(5):323–9.

- 7. Williams C, Jiang YH, Shashi V, Crimian R, Schoch K, Harper A, et al. Additional evidence that PGAP1 loss of function causes autosomal recessive global developmental delay and encephalopathy. Clin Genet. 2015;88(6):597–9.
- 8. Richards S, Aziz N, Bale S, Bick D, das S, Gastier-Foster J, et al. Standards and guidelines for the interpretation of sequence variants: a joint consensus recommendation of the American College of Medical Genetics and Genomics and the Association for Molecular Pathology. Genet Med. 2015;17(5):405–24.
- Burgos CM, Frenckner B. Addressing the hidden mortality in CDH: a population-based study. J Pediatr Surg. 2017;52(4):522-5.
- McGivern MR, Best KE, Rankin J, Wellesley D, Greenlees R, Addor MC, et al. Epidemiology of congenital diaphragmatic hernia in Europe: a register-based study. Arch Dis Child Fetal Neonatal Ed. 2015;100(2):F137–F144.
- 11. Yu L, Hernan RR, Wynn J, Chung WK. The influence of genetics in congenital diaphragmatic hernia. Semin Perinatol. 2020;44(1):151169.
- Crenshaw MM, Thompson L, Pique DG, Micke K, Saenz M, Baker PR 2nd. Congenital diaphragmatic hernia in siblings with PIGA-related congenital disorder of glycosylation. Am J Med Genet A. 2023;191(12):2860-7.
- 13. Bayat A, Knaus A, Pendziwiat M, Afenjar A, Barakat TS, Bosch F, et al. Lessons learned from 40 novel PIGA patients and a review of the literature. Epilepsia. 2020;61(6):1142–55.
- Brady PD, Moerman P, De Catte L, Deprest J, Devriendt K, Vermeesch JR. Exome sequencing identifies a recessive PIGN splice site mutation as a cause of syndromic congenital diaphragmatic hernia. Eur J Med Genet. 2014;57(9):487–93.
- Meier N, Bruder E, Lapaire O, Hoesli I, Kang A, Hench J, et al. Exome sequencing of fetal anomaly syndromes: novel phenotype-genotype discoveries. Eur J Hum Genet. 2019;27(5):730-7.
- Reynolds KK, Juusola J, Rice GM, Giampietro PF. Prenatal presentation of Mabry syndrome with congenital diaphragmatic hernia and phenotypic overlap with Fryns syndrome. Am J Med Genet A. 2017;173(10):2776–81.
- Winter-Paquette LM, Al Suwaidi HH, Sajjad Y, Bricker L. Congenital diaphragmatic hernia and early lethality in PIGL-related disorder. Eur J Med Genet. 2022;65(5):104501.

How to cite this article: Benabess S, Myers KA. Recurrent status epilepticus and severe bifrontal hypometabolism in PGAP1-related neurodevelopmental disorder. Epileptic Disord. 2025;27:304–306. https://doi.org/10.1002/epd2.20336