

Adult-onset KMT2B-related dystonia

Edoardo Monfrini^{1,2}, A. Ciolfi³, F. Cavallieri^{4,5}, M. Ferilli³, P. Soliveri^{6,7}, L. Pedace⁸, R. Erro⁹, F. Del Sorbo^{6,7}, F. Valzania⁴, V. Fioravanti⁴, G. Cossu¹⁰, M. Pellegrini¹¹, L. Salviati¹², F. Invernizzi¹³, V. Oppo¹⁰, D. Murgia¹⁰, B. Giometto¹¹, M. Picillo⁹, B. Garavaglia¹³, F. Morgante^{14,15}, M. Tartaglia³, M. Carecchio^{16,17}, A.B. Di Fonzo²

¹Dino Ferrari Center, Neuroscience Section, Department of Pathophysiology and Transplantation, University of Milan, Milan, Italy

²Foundation IRCCS Ca' Granda Ospedale Maggiore Policlinico, Neurology Unit, Milan, Italy

³Genetics and Rare Diseases Research Division, Ospedale Pediatrico Bambino Gesù, IRCCS, Rome, Italy

⁴Neurology Unit, Neuromotor & Rehabilitation Department, Azienda USL-IRCCS di Reggio Emilia, Reggio Emilia, Italy

⁵Clinical and Experimental Medicine PhD Program, University of Modena and Reggio Emilia, Reggio Emilia, Italy

⁶Parkinson Institute, ASST G. Pini-CTO, Milan, Italy

⁷Fondazione Grigioni per il Morbo di Parkinson, Milan, Italy

⁸Department of Onco-Hematology, Cell Therapy, Gene Therapy and Hemopoietic Transplant, Ospedale Pediatrico Bambino Gesù, IRCCS, Rome, Italy

⁹Department of Medicine, Surgery and Dentistry "Scuola Medica Salernitana", Neuroscience Section, University of Salerno, Baronissi (SA), Italy

¹⁰Department of Neuroscience, Brotzu Hospital, Cagliari, Italy

¹¹Neurology Unit, Trento Hospital, Azienda Provinciale per i Servizi Sanitari (APSS) di Trento, Trento, Italy

¹²Clinical Genetics Unit, Department of Woman and Child Health, University of Padua, Padua, Italy

¹³Medical Genetics and Neurogenetics Unit, Fondazione IRCCS Istituto Neurologico C. Besta, Milan, Italy

¹⁴Neurosciences Research Centre, Molecular and Clinical Sciences Research Institute, St George's, University of London, London, UK

¹⁵Department of Experimental and Clinical Medicine, University of Messina, Messina, Italy

¹⁶Parkinson Disease and Movement Disorders Unit, Department of Neuroscience, University of Padua, Padua, Italy

¹⁷Study Center for Neurodegeneration (CESNE), University of Padua, Padua, Italy

Introduction: KMT2B-related dystonia (DYT-KMT2B, also known as DYT28) is an autosomal dominant neurological disorder characterized by varying combinations of generalized dystonia, psychomotor developmental delay, mild-to-moderate intellectual disability, and short stature. Disease onset occurs typically before ten years of age [1-3].

We report the clinical and genetic findings of a series of subjects affected by adult-onset dystonia, hearing loss, or intellectual disability carrying rare heterozygous KMT2B variants.

Twelve cases from five unrelated families carrying four rare KMT2B missense variants predicted to impact protein function are described. Seven affected subjects presented with adult-onset focal or segmental dystonia, three developed isolated progressive hearing loss, and one displayed intellectual disability and short stature. Genome-wide DNA methylation profiling allowed to discriminate these adult-onset dystonia cases from controls and early-onset DYT-KMT2B patients [4].

These findings document the relevance of KMT2B variants as a potential genetic determinant of adult-onset dystonia and prompt to further characterize KMT2B carriers investigating non-dystonic features.

References:

- [1] Carecchio M, Invernizzi F, González-Latapi P, et al. Frequency and phenotypic spectrum of KMT2B dystonia in childhood: A single-center cohort study. *Movement Disorders*. 2019;34(10):1516-1527. doi:10.1002/mds.27771.
- [2] Zech M, Boesch S, Maier EM, et al. Haploinsufficiency of KMT2B, Encoding the Lysine-Specific Histone Methyltransferase 2B, Results in Early-Onset Generalized Dystonia. *Am J Hum Genet*. 2016;99(6):1377-1387. doi:10.1016/j.ajhg.2016.10.010.
- [3] Meyer E, Carss KJ, Rankin J, et al. Mutations in the histone methyltransferase gene KMT2B cause complex early-onset dystonia. *Nat Genet*. 2017;49(2):223-237. doi:10.1038/ng.3740.
- [4] Ciolfi A, Foroutan A, Capuano A, et al. Childhood-onset dystonia-causing KMT2B variants result in a distinctive genomic hypermethylation profile. *Clinical epigenetics*. 2021;13(1):157. doi:10.1186/s13148-021-01145-y.