Translational Medicine @ UniSa - ISSN 2239-9747

2019, Special Issue 1(5): 5

## OC.1- NOVEL CHD2 AND KIAA2022 MUTATIONS ASSOCIATED WITH EYELID MYOCLONIA WITH ABSENCES

<u>A. Coppola</u><sup>1</sup>, S. Krithika<sup>2,3</sup>, M. Iacomino<sup>6</sup>, S. Balestrini<sup>2,3</sup>, L. Hernandez-Hernandez<sup>2,3</sup>, S. Meletti<sup>4,5</sup>, G. Gobbi<sup>7</sup>, E. Ferlazzo<sup>8,9</sup>, L. Giordano<sup>10</sup>, S. Casellato<sup>11</sup>, V. Sofia<sup>12</sup>, P. Striano<sup>13</sup>, L. Bilo<sup>1</sup>, V. Nigro<sup>14,15</sup>, A. Torella<sup>14,15</sup>, F. Musacchia<sup>14</sup>, F. Zara<sup>6</sup>, and S.M. Sisodiya<sup>2,3</sup>

<sup>1</sup>Department of Neuroscience, Reproductive and Odontostomatological Sciences, Epilepsy Centre, Federico II University, Naples, Italy; <sup>2</sup>Department of Clinical and Experimental Epilepsy, UCL Institute of Neurology; <sup>3</sup>Chalfont Centre for Epilepsy, Chalfont St Peter, UK Department of Biomedical, Metabolic, and Neural Science; <sup>4</sup>University of Modena and Reggio Emilia, Modena, Italy; <sup>5</sup>Neurology Unit, OCSAE Hospital, AOU Modena, Italy; <sup>6</sup>Laboratory of Neurogenetics, Department of Neurosciences, Institute "G. Gaslini", Genoa, Italy; <sup>7</sup>IRCCS, Institute of Neurological Sciences of Bologna, Child Neurology Unit, via Altura 3, Bologna, Italy; <sup>8</sup>Department of Medical and Surgical Sciences, Magna Græcia University of Catanzaro, Viale Europa, Germaneto, Catanzaro, Italy; <sup>9</sup>Regional Epilepsy Center, Great Metropolitan Hospital, Via Melacrino, Reggio Calabria, Italy; <sup>10</sup>Child Neuropsychiatry Unit- Civile Hospital, Brescia, Italy; <sup>11</sup>Pediatric Neuropsychiatric Division Unit, AOU Sassari, Italy; <sup>12</sup>Department of Medical and Surgical Sciences and Advanced Technologies "G.F. Ingrassia", University of Catania, Catania, Italy; <sup>13</sup>Pediatric Neurology and Muscular Diseases Unit, Department of Neurosciences, Rehabilitation, Ophtalmology, Genetics, Maternal and Child Health, University of Genoa, "G. Gaslini" Institute, Genova, Italy; <sup>14</sup>Telethon Institute of Genetics and Medicine, Pozzuoli, Naples, Italy; <sup>15</sup>Dipartimento di Biochimica, Biofisica e Patologia Generale, Università degli Studi della Campania "Luigi Vanvitelli", Naples, Italy

**Purpose**: Eyelid myoclonia with absences (EMA) is a generalized genetic epilepsy syndrome characterised by the hallmark of rapid blinking of the eyelids and an upward deviation of the eyeballs. Understanding the genetic architecture of EMA is a challenging task as eyelid myoclonia is also seen in other epilepsy syndromes.

Method: The present study is based on a cohort of individuals with either only EMA, showing eyelid myoclonia with eye closure EEG bursts and/or photosensitivity (EMA), or with EMA associated with clinically evaluated intellectual disability or psychiatric conditions (EMA+). Whole-exome sequencing (WES) was performed on probands and their parents, when available. WES data was analysed, trios were analysed for de novo and/or unique variants and the singletons for unique variants. Sanger sequencing was used to confirm variants.

**Results**: We sequenced 112 individuals, including 19 trios, 40 singletons and 2 multiplex families. We identified 2 novel de novo CHD2 mutations in 2 EMA+ trios: a missense variant [c.4598 T>G] and a splice donor variant [c.3455+2 T>G]. We also identified a de novo frameshift deletion affecting KIAA2022 [c.2171del] in another EMA+ trio. All these variants were unique, with CADD score>20. Two singletons showed a previously reported mutation in the CHD2 gene: a frameshift variant [c.3734dupA] and a splice region variant [c.2577+7T>C]. The former variant was identified in an EMA+ case, and the latter in an EMA case. Overall, 10.5% of trios showed a de novo CHD2 variant and 5% of singletons harboured mutation in this gene.

**Conclusion**: CHD2 gene plays a key role in cerebral cortical development. Mutations in this genehave been reported for a broad spectrum of neurodevelopmental disorders, including EMA. KIAA2022 encodes a neurite extension and migration factor. Mutations in this gene have never been associated with EMA. Our study adds growing evidence implicating novel CHD2 and KIAA2022 mutations in EMA.

