

## Many testing roads can lead to KCNQ2 diagnosis.

Amal Yussuf<sup>1</sup>, Zoe Powis<sup>2</sup>, Kirsten Blanco<sup>1</sup>, Jing Wang<sup>2</sup>

<sup>1</sup>Clinical Research, <sup>2</sup>Clinical Genomics, Ambry Genetics

### Objective:

To evaluate different genetic testing approaches for *KCNQ2* related disorders.

### Background:

Mutations in the *KCNQ2* gene are associated with a broad phenotypic spectrum, from Benign Familial Neonatal Epilepsy to Neonatal Epileptic Encephalopathy. A diagnosis of *KCNQ2*-related epilepsy can be a confusing process. Genetic testing is key to confirm clinical diagnosis and guide prompt therapeutic treatment.

### Design/Methods:

We reviewed clinical indications for individuals with pathogenic mutations in *KCNQ2* detected by NGS multi-gene panel testing (MGPT) and diagnostic exome sequencing (DES).

### Results:

*KCNQ2* mutations were identified in a total of 13 patients, age range from 3 months to 23 years old. Six (3.75%) patients were tested by MGPT and seven (0.63%) by DES. Of the detected alterations, 5 were truncation and 7 were missense mutations. There was 1 whole gene deletion detected by MGPT. Interestingly, *KCNQ2* alterations in all 7 patients tested by DES were confirmed to be *de novo*. All patients have history of seizures. Only 3/6 patients who had MGPT presented other phenotypic features: cardiac or cutaneous disorders and migraine. None of these cases demonstrated developmental delay (DD). 57.1% of EEG and MRI results were normal. Patients tested by DES displayed a wider variety of additional symptoms such as vision disorder, hearing impairment, dysmorphic features, gastrointestinal disorder, and spasticity. 6/7 patients had DD. 86% of the patients had at least one metabolic screening and 40% of EEG and MRI results were abnormal.

### Conclusions:

*KCNQ2* positive patients tested by DES had a more complex medical history than those tested by MGPT. Patients with a more complex phenotype may benefit DES testing to avoid a long diagnostic journey, while others may benefit from direct MGPT testing. Although ~10% of pathogenic *KCNQ2* variants have been reported to be *de novo*, our findings suggest the *de novo* rate is higher than previously reported.