### **Accepted Manuscript**

Amino acid-level signal-to-noise analysis of incidentally identified variants in genes associated with long QT syndrome during pediatric whole exome sequencing reflects background genetic noise

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PII: S1547-5271(18)30142-5

DOI: 10.1016/j.hrthm.2018.02.031

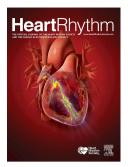
Reference: HRTHM 7500

To appear in: Heart Rhythm

Received Date: 4 January 2018

Please cite this article as: Landstrom AP, Fernandez E, Rosenfeld JA, Yang Y, Dailey-Schwartz AL, Miyake CY, Allen HD, Penny DJ, Kim JJ, Amino acid-level signal-to-noise analysis of incidentally identified variants in genes associated with long QT syndrome during pediatric whole exome sequencing reflects background genetic noise, *Heart Rhythm* (2018), doi: 10.1016/j.hrthm.2018.02.031.

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#### ACCEPTED MANUSCRIPT

# Amino acid-level signal-to-noise analysis of incidentally identified variants in genes associated with long QT syndrome during pediatric whole exome sequencing reflects background genetic noise

Running title: LQTS-associated variants in WES testing

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