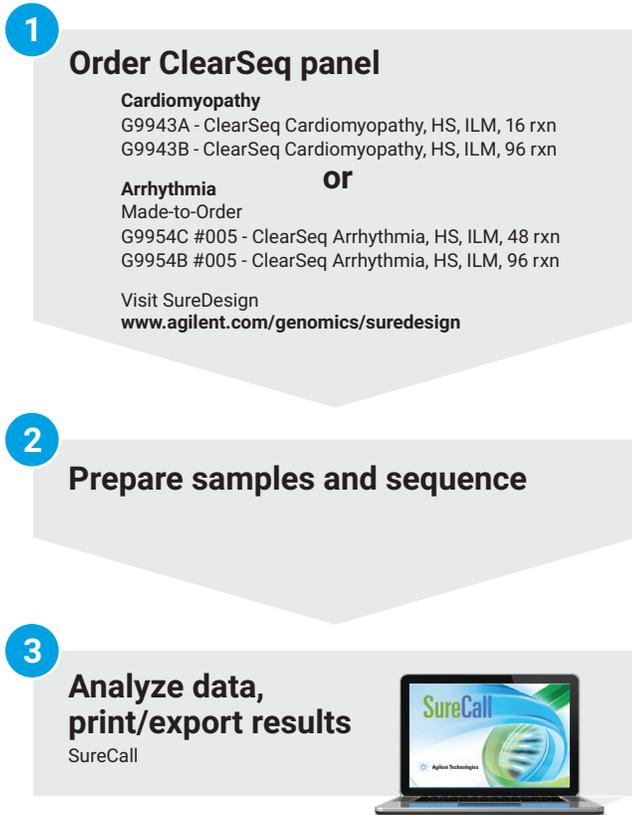


ClearSeq Arrhythmia

Targeting genomic regions known to be associated with four inherited arrhythmia related heart disorders, ClearSeq Arrhythmia (made-to-order) is a next generation sequencing target enrichment panel. It is a result of a thorough review of publications for arrhythmia and drew on information in GeneReviews in order to incorporate 21 genes known to correlate with Long QT syndrome, Short QT syndrome, Brugada syndrome, and Catecholaminergic polymorphic ventricular tachycardia. The genes associated with different types of arrhythmia are overlapping in some cases, as shown in the table below, and using ClearSeq Arrhythmia, a comprehensive arrhythmia profile can be assembled for clinical research samples.

KCNQ1	KCNH2	KCNJ2	ANK2	KCNE1	KCNE2	CACNA1C	CAV3	SCN4B	AKAP9	SNTA1	SCN5A	GPD1L	CACNA1C	CACNB2	SCN1B	KCNE3	SCN3B	RYR2	KCNJ2	CASQ2
Short QT syndrome			Long QT syndrome								Brugada syndrome						Catecholaminergic polymorphic ventricular tachycardia			

From Sample to Result in Less than 2 Days



Premium Target Enrichment Performance

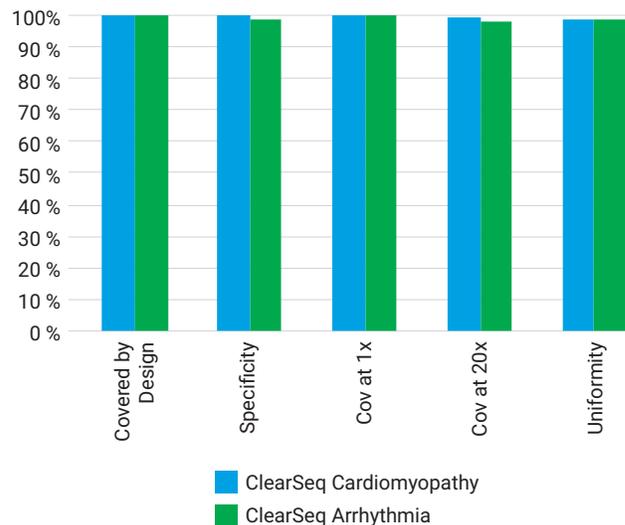


Figure 1. Results from Illumina MiSeq runs; Uniformity calculated as percent bases covered at $\geq 10\%$ of mean coverage.

Request more information or buy online:

www.agilent.com/genomics/ngs

For Research Use Only. Not for use in diagnostic procedures.

This information is subject to change without notice.

PR7000-0350
 © Agilent Technologies, Inc. 2017
 Published in the USA, September 27, 2017
 5991-2525EN