

Figure 3: Multiplexing Capabilities of the MiSeqDx Cystic Fibrosis 139-Variant Assay. A highly multiplexed method simultaneously sequences up to 48 samples in a single sequencing run.

Widely Adopted NGS Platform

Illumina sequencing by synthesis (SBS) technology is widely adopted in the sequencing community. Through massively parallel sequencing using a proprietary reversible terminator-based method, SBS enables detection of single bases as they are incorporated into growing DNA strands. A fluorescently labeled terminator is imaged as each dNTPs (dATP, dCTP, dGTP, or dTTP) are added and then cleaved to allow incorporation of the next base. Because all four reversible terminator-bound dNTPs are present during each sequencing cycle, natural competition minimizes incorporation bias. The result is base-by-base sequencing for highly accurate data even in difficult regions, such as homopolymers.

Easy Results Interpretation

Results from the MiSeqDx Cystic Fibrosis 139-Variant Assay are presented in an easy-to-read fashion that a board-certified molecular geneticist or equivalent can readily interpret. The report includes assay name, sample ID, dbSNP ID, and the call rate for each sample (Figure 4). Call rates must be $\geq 99\%$ to be considered valid.

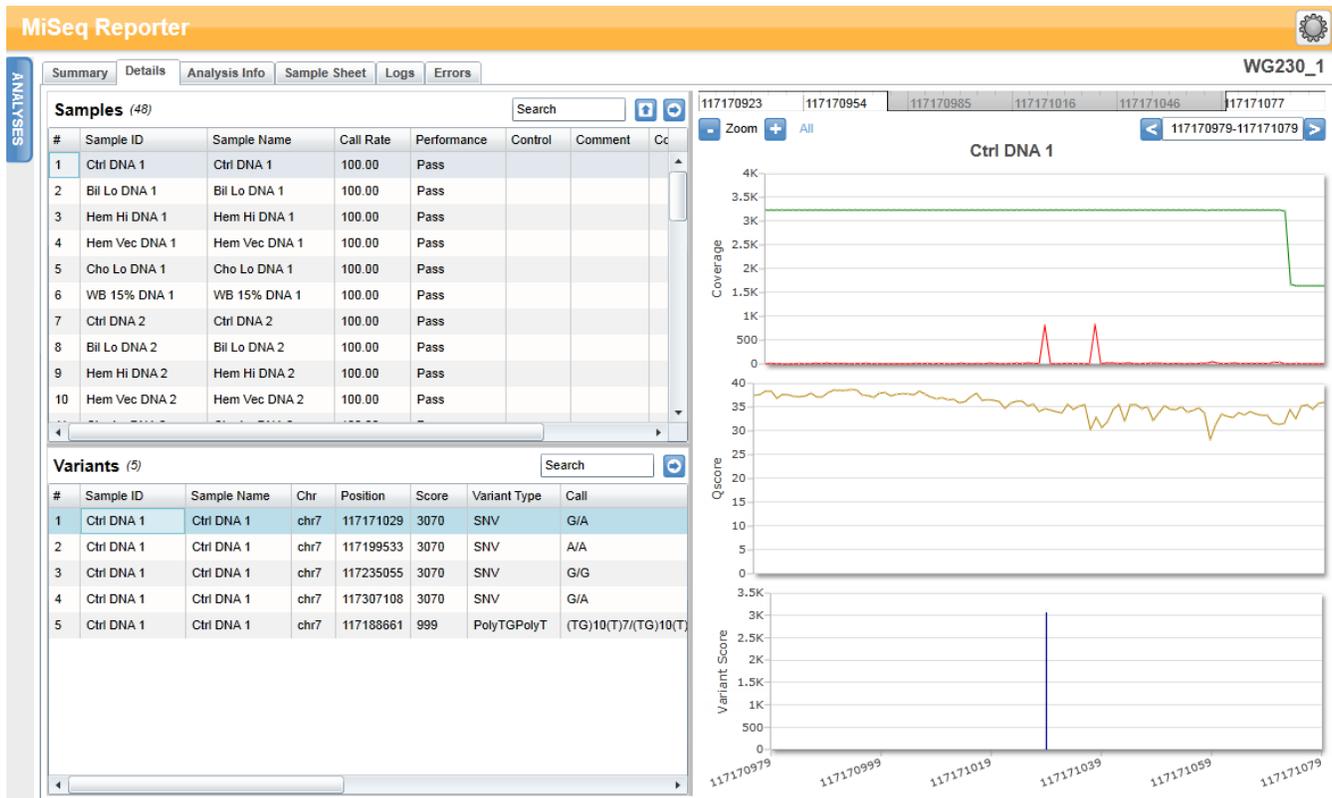


Figure 4: Easy Visualization Using the MiSeq Reporter Software.

