Precision Melt Analysis software is a convenient, easy-to-use application that imports and analyzes data files generated from the CFX96™ or CFX384™ real-time PCR detection system to genotype samples based on the thermal denaturation properties of double-stranded DNA. The software can be used for a variety of genotyping applications, including scanning for new gene variants, screening DNA samples for SNPs, identifying insertions/deletions or other unknown mutations, and determining the percentage of methylated DNA in unknown samples.

Precision Melt Analysis software makes it easy for you to:

- Streamline your data analysis using the customizable default analysis settings
- Utilize the multiple data view options to manually assign sample genotypes by tailoring the software to the appropriate analysis
- Examine results from a number of melt files, without having to export data, using the Melt Study module
- Analyze multiple experiments from a single plate using the Well Groups feature
- Publish your data in several formats by easily exporting data and images
Efficient and Effective Analysis

Precision Melt Analysis software saves analysis time by assigning sample genotypes automatically based on cluster analysis, or manually using multiple data view options to tailor the software to the appropriate analysis. Use the normalized melt curves plot feature to generate a basic representation of the different clusters based on curve shifting (for homozygotes) and curve shape change (for heterozygotes). Difference curve plots of a sample fluorescence versus a selected control at each temperature transition provide a convenient visual aid to interpret the data.

Precision Melt Analysis software enables data comparison between multiple file runs by combining data into a single Melt Study. Develop a standard library of melt curve runs to analyze an unlimited number of melt experiments without having to export data.

SNP genotyping by Precision Melt Analysis software using data generated by the CFX384 real-time PCR detection system. Discrimination of human hemochromatosis S65C SNP genotypes (A to T substitution). Data from homozygous wild type (■), mutant (■), and heterozygote (■) samples are shown on a normalized melt curve plot, RFU, relative fluorescence units.

Multifile melt study analysis. Compare melt curves from different files using the Melt Study module in Precision Melt Analysis software. RFU, relative fluorescence units.

Ordering Information

Catalog # Description
184-5025 Precision Melt Analysis Software, includes 2 user licenses, installation CD, 2 HASP HL keys, melt calibration kit
185-5096 CFX96 Real-Time PCR Detection System, includes C1000™ thermal cycler chassis, CFX96 optical reaction module, CFX Manager™ software, license for qbase+™ software, communication cable, reagents, consumables
185-5384 CFX384 Real-Time PCR Detection System, includes C1000 thermal cycler chassis, CFX384 optical reaction module, CFX Manager software, license for qbase+™ software, communication cable, reagents, consumables

A representative difference curve plot generated from data normalized to a mutant sample. Data from homozygous wild type (■), mutant (■), and heterozygote (■) samples are shown. RFU, relative fluorescence units.

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