

Mic- Your Personal qPCR cyclers

The world's first Magnetic Induction Cycler.

The box is small. The ideas are big.



SPEED



ACCURACY



SIZE



CONNECTIVITY

MIC optional High Resolution Melting software, **free** UNTIL 31 Dec.2017

Mic's optional HRM analysis characterises DNA samples according to their melt behaviour so you can identify mutations. HRM is the perfect tool for applications including determining allele prevalence, screening for loss of heterozygosity, DNA fingerprinting, DNA methylation, species identification and calculating the ratio of somatic acquired mutations.

Even difficult Class IV SNPs are no problem for Mic – the example here clearly shows the A base allele (red), T base allele (blue) and the heterozygote (purple).

CONTACT US AND REQUEST A DEMO info@sgsh.es