

The Effect of White Pigmentation in 96-well Plates During QPCR

Saima Naveed Nayab¹, Philip N. Harries², Simon C. Baker² and Ian Kavanagh^{1†}

¹: Thermo Fisher Scientific, ABgene House, Blenheim Road, Epsom, Surrey, KT19 9AP
²: School of Life Sciences, Oxford Brookes University, Gypsy Lane, Oxford, OX3 0BP.



Introduction

QPCR instruments are able to monitor amplicon quantity in real-time during PCR reactions by detecting fluorescence signals and recording fluorescence data. The majority of fluorescence is reflected out of the QPCR plate either by the polypropylene itself or by the walls of the thermal cycling block, when natural polypropylene is used. QPCR plates containing a white pigment improve reflection towards the fluorescence detector inside the thermocycler during a QPCR run (Figure 1 and Figure 2).

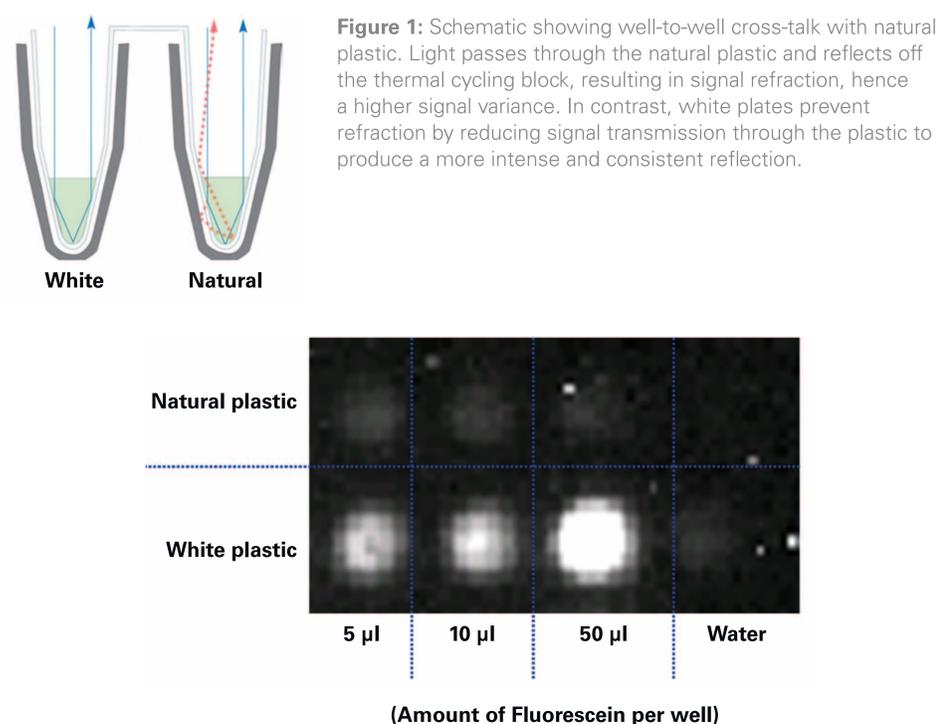


Figure 2: White plastic generates greater reflection of light back to the detector compared to natural plates, resulting in higher signal-to-background ratio. Three dilutions of fluorescein were added to each well and detected using a CCD camera.

Aim

To investigate if white 96-well plates allow better fluorescence detection of PCR products during QPCR assays than natural 96-well plates.

Method

A QPCR assay targeting a 94 bp region of human GAPDH was selected to assess QPCR performance of natural versus Thermo Scientific white 96-well QPCR plates (AB-0600). In order to avoid inter-run variation, both types of plates were cut in half and sealed (Figure 3) using a Thermo Scientific ALPS 50V manual heat sealer, for simultaneous assay in a Stratagene Mx3005P QPCR instrument (Stratagene, Leicester).

Human genomic DNA (Sigma Aldrich, Dorset) was serially diluted to 100 ng, 10 ng, 1 ng and 100 pg in the final reaction. Eight replicate samples were used for each template dilution and the experiment was repeated 3 times. Thermo Scientific ABsolute™ QPCR SYBR® Green Mix (AB-1162) was used for all experiments. Both GAPDH primers (forward 5' ACAGTCAGCCGCATCTTCTT 3' and reverse 5' ACGACCAATCCGTTGACTC 3' (Thermo Scientific) were used at a final concentration of 70 nM.

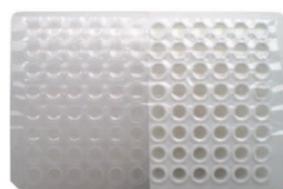


Figure 3: A digital image showing the natural and white plates (AB-0600) cut in half and heat sealed together in order to assess their QPCR performance in a single experiment.

Results

Results presented in Table 1 and Figure 3 show that compared to the natural plates, white QPCR plates generate earlier C_T and higher end-point fluorescence values. The GAPDH amplification plots are consistently superior across all four template dilutions with white plates. Figure 4 shows notably more refined SYBR Green melting profiles for the GAPDH amplicon using white plates in comparison with those produced using natural plates.

Table 1: Mean C_T Values for GAPDH				
DNA (ng)	100	10	1	0.1
Natural Plates	25.45	28.63	31.63	35.05
White Plates	23.86	26.98	30.62	34.30
ΔC_T (White-Natural)	-1.59	-1.64	-1.01	-0.75

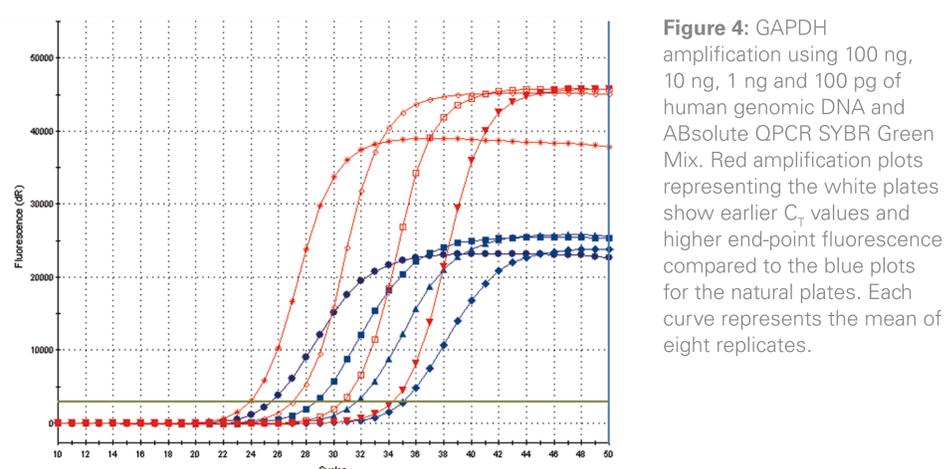


Figure 4: GAPDH amplification using 100 ng, 10 ng, 1 ng and 100 pg of human genomic DNA and Absolute QPCR SYBR Green Mix. Red amplification plots representing the white plates show earlier C_T values and higher end-point fluorescence compared to the blue plots for the natural plates. Each curve represents the mean of eight replicates.

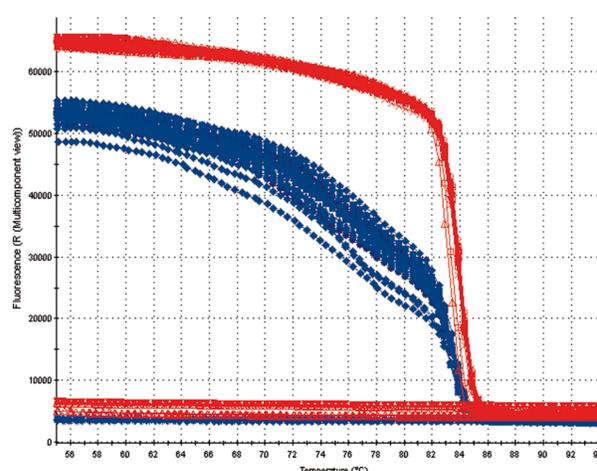


Figure 5: Melting profiles of GAPDH amplicons in white plates (red) and natural plates (blue) across four 10-fold dilutions of human genomic DNA.

Conclusion

- Earlier C_T and higher end-point fluorescence values using the white plates, indicate greater QPCR sensitivity due to higher signal-to-background ratio and minimal well-to-well cross-talk compared to natural plates.
- A significant enhancement in shape and precision of the melt curve shows greater QPCR specificity.
- The use of white QPCR plates for high resolution melt analysis may result in more sensitive SNP discrimination, and improved detection of low copy number targets in comparison with natural plates.

References

1. Nayab SN, Jones G and Kavanagh I (2008) Absolute™ Fast QPCR Master Mix: Minimizing Protocol Time without Compromising Performance. *Nature Methods* December 5 (12) an2.

† For more information contact ian.kavanagh@thermofisher.com