Amplification efficiency and thermal stability of qPCR instrumentation: Current landscape and future perspectives

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18 Introduction

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20 Quantitative polymerase chain reaction (qPCR), the method by 21 which a small sample of genetic material can be exponentially 22 amplified and quantitatively measured in real time, is now a 23 mainstay of research and medical laboratories. As the process 24 has evolved, the applications for qPCR have increased rapidly, 25 and include the detection of infectious diseases, paternity identification, forensic analysis and food processing. The PCR 26 27 process necessitates the cycling of test samples through a temperature profile, typically 95, 55 and 72°C, multiple times. 28 29 The time taken to change the temperature of the samples 30 between these levels is a key determinant of the speed of the 31 process and thus of the duration of a test (1). A typical 40-cycle 32 PCR can take ~2 h to complete and improvements in that time 33 have not been achieved as rapidly as the advances in other

Abstract. Quantitative polymerase chain reaction (qPCR)

is a method of amplifying and detecting small samples of

genetic material in real time and is in routine use across many

laboratories. Speed and thermal uniformity, two important

factors in a qPCR test, are in direct conflict with one another

in conventional peltier-driven thermal cyclers. To overcome

this, companies are developing novel thermal systems for

qPCR testing. More recently, qPCR technology has developed

to enable its use in point-of-care testing (POCT), where the

test is administered and results are obtained in a single visit

to a health provider, particularly in developing countries. For

a system to be suitable for POCT it must be rapid and reliable.

In the present study, the speed and thermal uniformity of four

qPCR thermal cyclers currently available were compared, two

of which use the conventional peltier/block heating method

and two of which use novel heating and cooling methods.

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areas. Therefore, some of the potential benefits of the qPCR 34 process remain limited by speed. 35

Thermal uniformity, the absence of which can cause 36 discrepancies in the cycling conditions between different 37 samples on the same plate, is directly linked to speed. In many 38 PCR instruments, conductive blocks are used to connect the 39 heating or cooling source(s) to the test samples. When heating 40 and cooling the system, it is necessary to drive heat into and 41 out of the block. Temperature gradients are eliminated by the 42 natural flow of heat within the blocks; therefore, over time the 43 same conditions should be delivered to all the test samples. 44 However, block-based systems are vulnerable to greater heat 45 losses on the edges and surfaces that tend to distort the thermal 46 distribution. The conductivity of the blocks affects the rate of 47 heat flow and thus the uniformity of heating of the samples. In 48 addition, the larger the thermal mass of the block, the greater 49 the amount of heat to be transferred and the longer this will 50 take. The faster heat is driven into or out of the system, the 51 less time is available for the temperature distribution of the 52 conductive block to even out and for thermal uniformity to 53 be maintained. Ultimately, such a system can only maintain 54 thermal uniformity if the rate of change of temperature is 55 slower than the time it takes for the temperature of the conduc-56 tive block to even out. To achieve quick cycle times, large 57 temperature gradients are applied to the block, which can lead 58 to the target temperatures of samples being over- or undershot. 59 Thus, in these types of systems, the requirement for uniformity 60 of temperature directly conflicts with the desire for speed; they 61 are able to deliver one feature or the other but not both (2). 62

The ABI Prism 7900HT (Applied Biosystems, Thermo 63 Fisher Scientific, Waltham, MA, USA) is perhaps the industry 64 standard peltier/block-based thermal cycler. The CFX96 65 (Bio-Rad Laboratories, Inc., Hercules, CA, USA) provided an 66 upgrade to the conventional system by reducing the thermal 67 mass of the block. Alternatives to the block-based system have 68 also been developed. The Rotor-Gene Q (Qiagen, Hilden, 69 70 Germany) combines a centrifugal set-up with an air-based thermal system. Ensuring that samples are continuously 71 rotated through heated air removes the edge effect to provide 72 superior thermal uniformity. xxpress® (BJS Biotechnologies, 73 Perivale, UK) employs a different system in which an 'active 74 heating' method is combined with a block of low thermal 75 76 mass, precisely controlling the amount and location of additional heating to avoid temperature discrepancies (Table I). 77 The present study investigated and compared the efficiency 78

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Table I. Ramp rate and thermal uniformity of aPCR instruments.

| PCR platform | Thermal system | Advertised fastest ramp rate (°C/sec) | Advertised thermal uniformity (° |
|------------------------------|-------------------|---------------------------------------|--|
| ABI Prism 7900HT | Block/peltier | 1.5 | ± 0.5 (measured 30 sec after timing starts) |
| Bio-Rad CFX96 | Block/peltier | 3.3 (average) | ±0.4 (well-to-well within 10 sec of reaching 90°C) |
| Qiagen Rotor-Gene Q | Air | 15 (peak) | ±0.02 |
| BJS biotechnologies xxpress® | Resistive heating | 10 | ±0.3 |

and thermal uniformity of four of the qPCR thermal cyclers currently available that use the conventional block/peltier system or novel methods.

20 Materials and methods

22 *qPCR*. The expression of 18S rRNA in human genomic DNA 23 was assessed and compared by qPCR using an ABI Prism 24 7900HT, a Bio-Rad CFX96 System, a Qiagen Rotor-Gene Q 25 and a BJS Biotechnologies xxpress®. Human genomic DNA was purchased from Bioline (Meridian Bioscience, Cincinnati, 26 27 OH, USA) and input in concentrations of 100, 10, 1, 0.1 and 28 0.01 ng/ μ l to give final concentrations of 5, 0.5, 0.05, 0.005 29 and 0.0005 ng/ μ l, generating a standard curve. Eukaryotic 30 18S rRNA gene primers were used as follows: forward, 31 3'-AAACGGCTACCACATCCAAG-5' and reverse, 3'-CCTC-32 CAATGGATCCTCGTTA-5'. SYBR FAST qPCR master mix 33 (Kapa Biosystems, Wilmington, MA, USA) was used across 34 all platforms using the following thermal profile: 20 sec hot start at 95°C followed by 40 cycles of 95°C for 1 sec and 60°C 35 for 10 sec. Heating and cooling rates and all other parameters 36 37 were at the manufacturers' pre-set levels.

38 Thermal variability was assessed in qPCR by measuring 39 the amplification of 18S rRNA in a selection of wells 40 covering all areas of the sample plate on ABI Prism 7900HT, Bio-Rad CFX96 System, Qiagen Rotor-Gene Q and BJS 41 42 Biotechnologies xxpress instruments. Human genomic DNA 43 at 100 ng/ μ l (final concentration, 5 ng/ μ l) was used with the protocol detailed above. 44

Statistical analysis. Statistical tests commonly used to 46 determine the reliability and accuracy of a quantitative PCR 47 assay include performing a standard curve experiment with 48 49 each dilution series run in triplicate. The Cq value was plotted 50 against the log of the nucleic acid input level to generate a 51 linear graph. The slope or gradient of this graph was used to 52 determine the PCR reaction efficiency and a linear regression 53 analysis with a correlation coefficient or R² value was included 54 to determine the accuracy and repeatability of the standard 55 curve. The ideal result is a PCR reaction efficiency of 100% and an \mathbb{R}^2 value of 1. An efficiency of <90 or \geq 110% is unac-56 57 ceptable and indicates that further optimisation is required. If 58 the R^2 value is ≤ 0.985 , this raises questions about assay reli-59 ability with respect to pipetting accuracy and the range of the 60 assay (3).

Results

Amplification efficiency. A standard curve was generated by 78 amplifying 18S rRNA in human genomic DNA at concentra-79 tions of 5, 0.5, 0.05, 0.005 and 0.0005 ng/ μ l and plotting C_t 80 against log concentration. Efficiency was calculated by the 81 following equation: Efficiency = $10^{(-1/\text{slope})-1}$. Efficiency of reac-82 tion values between 90 and 110% are considered acceptable 83 for qPCR reactions. The fastest instrument was the xxpress[®], 84 which completed 40 cycles in 12 min (Fig. 1). 85

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Thermal variability. Thermal variability was assessed by 87 measuring the amplification of 18S rRNA in 5 ng/ μ l human 88 genomic DNA in a selection of wells covering all areas of the 89 sample plate on ABI Prism 7900HT, Bio-Rad CFX96 System, 90 Qiagen Rotor-Gene Q and BJS Biotechnologies xxpress 91 instruments (Fig. 2). The average Ct, Ct spread and Ct standard 92 deviation were for CFX: 16.0, 1.315 and 0.34; for xxpress: 13.6, 93 1.2 and 0.29; for Prism 7900HT: 14.4, 4.526, and 1.91; and for 94 Rotor-Gene: 16.8, 1.319, and 0.43 (Fig. 3). 95

Discussion

qPCR instrumentation is rapidly evolving not only to meet 99 the needs of basic science but also in an attempt to address 100 some of the needs of the current healthcare system, in 101 terms of diagnosis as well as prognosis. For example, qPCR 102 technology has been widely used in the field of molecular 103 diagnostics for a number of infectious diseases (4). Food 104 and Drug Administration-approved qPCR-based screening 105 tests include group A Streptococcus and methicillin-resistant 106 Staphylococcus aureus (MRSA), HIV-1, human metapneumo- 107 virus and H1N1 influenza virus (5,6). More recently, Qiagen 108 received FDA approval of a therascreen® KRAS RGQ PCR 109 kit, paired with a colorectal cancer drug. KRAS mutations 110 occur in $\sim 40\%$ of colorectal cancer patients (7,8). Therefore, 111 screening patients by PCR will detect the most frequent muta- 112 tions in the KRAS gene and should aid with the selection of 113 therapeutic interventions. 114

Over the past decade, there has been a shift from testing in 115 reference hospitals/centres to clinical/diagnostic laboratories 116 worldwide (4). Point of care testing (POCT) allows a test to be 117 carried out and results obtained in a single visit to a primary 118 or secondary care health provider (9). In developing countries, 119 POCT is perhaps even more effective. The requirement for 120

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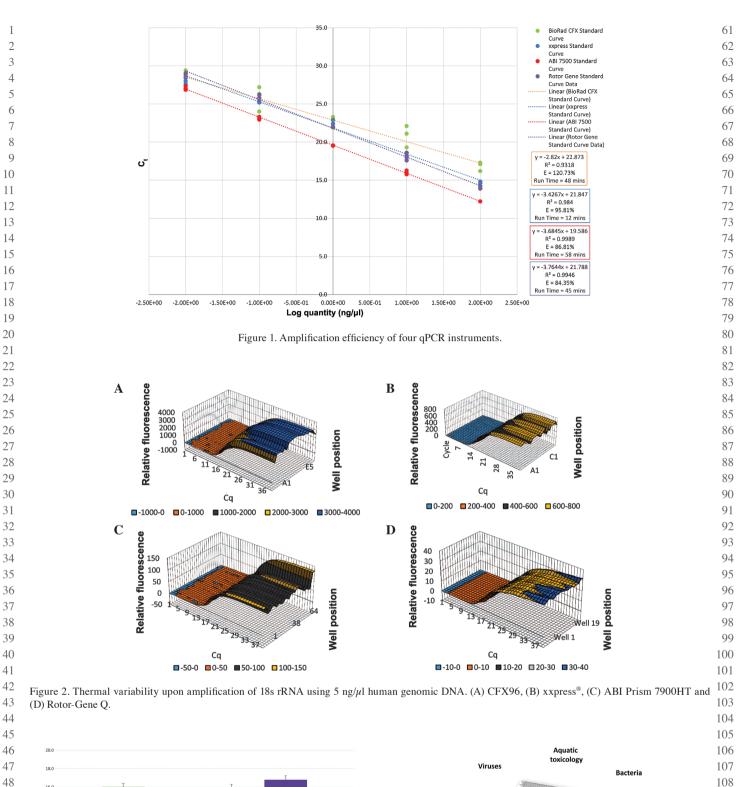
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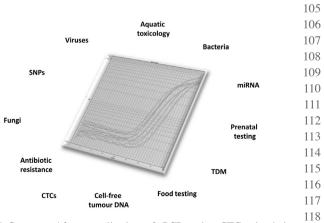


Figure 4. Current and future applications of qPCR testing. CTCs, circulating 119 tumour cells; SNPs, single nucleotide polymorphisms; TDM, therapeutic 120 drug monitoring; miRNA, microRNA.

expensive, central laboratories, highly trained technicians 1 and a reliable method of specimen and data transport can 2 all be removed with the implementation of a well-designed, 3 4 multifunctional POCT system. Bringing the test into the clinic 5 allows treatment to commence without delay and, in areas of 6 high displacement, reduces the likelihood of losing patient 7 contact before the condition has been effectively treated. This 8 is particularly important for communicable diseases such as 9 HIV/AIDS, measles and typhoid fever (10). An effective POC 10 test in a low resource setting is inexpensive to use and maintain. The test must be easy to operate, requiring little to no 11 12 training or specialist knowledge to both generate and interpret 13 results. In a recent study of sub-Saharan Africa, only 34% 14 of hospitals had reliable electricity access (11). Since energy 15 access for healthcare facilities in this region varies markedly, and as electrical sources may be unreliable, low electrical 16 consumption or even the ability to run on battery or solar 17 18 power is desirable.

The results of the present study demonstrate that the 19 20 performance of new technologies in qPCR instrumentation 21 such as Rotor-Gene Q and xxpress, is equally as good as, 22 or even better than, that of conventional qPCR instruments, 23 in terms of amplification efficiency and thermal uniformity. 24 However, a major advantage is that this instrument can deliver 25 40-cycle qPCR in <10 min. Rapid testing can be lifesaving. For example, rapid diagnostic tests can help in the diagnosis and 26 management of patients who present with signs and symptoms 27 compatible with influenza. These technologies can cut down 28 29 the time from 3-10 days for conventional viral cell cultures to 30 less than minutes (12). Infections with MRSA are known to 31 be associated with considerable morbidity and mortality (13). 32 Current sample preparation/testing times based on blood samples can take up to 5 h. However, in an emergency situation 33 34 this process might be too long if the patient admitted is posi-35 tive for MRSA and therefore has the potential to infect others. Equally, an early diagnosis of tuberculosis will assist not only 36 37 in the initiation of appropriate treatment but also limit the spread of this highly contagious disease (14). A test that could 38 39 be administered either at admission to the clinic, or even in an 40 ambulance on the way to the hospital, and takes only 10 min 41 could be of real benefit. Moreover, given the unreliability of 42 electricity in the developing world, diagnostic instrumentation 43 that is rapid is vital.

To date, qPCR-based diagnosis is often associated with high cost, time-consuming procedures, scientists and clinicians trained in qPCR analysis, lack of specificity and sensitivity or even standardisation for certain tests. In the future (Fig. 4), a standardised, rapid, scalable, affordable and easy-to-use48qPCR platform for use in POCT should provide an invalu-49able platform in the field of diagnostic/prognostic testing that50will complement the current conventional methods, including51microscopy, cell culture and immunological-based methods.52

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