Supplementary Information Robert Stedtfeld



Supplementary figures

Fig S1. Results from initial experiments examining optics with Gene-Z. (a) Determination of the signal acquisition timing settings. The coefficient of variation (%) for 4 replicate readings (y-axis on log scale) taken within the same reaction channel versus time required to obtain a signal (settings for signal acquisition are listed in table 2 supplemental material). (b) Comparison of a low and high brightness LED. Amplification profiles (Δ Rn on logarithmic scale) with \$0.60 LED (0.7 lumen, gray lines) and \$20 LED (145 lumen, solid black lines) with 20 µm of SYTO 81 dye, and using 0.7 lumen blue LED with calcein (dash black lines). LAMP was performed with approximately 7,600 copies of genomic DNA from *S. aureus* with a primer set targeting the *mecA* gene.



Fig S2. Schematic of a single reaction channel.



Fig S3. Pictorial of the chip fabrication process.



Fig S4. Rendering of the aluminum heater (also serves for alignment of optical components when chips are inserted) with embedded heater (dark gray), LEDs (green), and optical fibers (yellow). The inset shows the positing of the LEDS and optical fibers with respect to the shelled features of the chip.



Fig S5. Rendering of the aluminum heater shows placement of four thermocouple to test temperature uniformity. T₁ is the where the set-point temperature is measured for all experiments described in this study. The graph shows temperature of the four thermocouples as the heater in the device is ramped to reaction temperature and the average temperature once the set-point is reached. It should be noted that the standard error of measurement, determined by placing thermocouples propinquity on a heat plate is 0.22 °C.



Fig S6. Screen shots of iPod application menus. (a) Home screen. (b) Screen to help user select chip assay. (c) Example screen shot showing tests that can be selected manually. (d)

Manual selection of temperature and time to run the reaction. (e) Example screen shot showing amplification profiles plotted in real-time. (f) Example screen shot showing analyzed

data.



Fig S7. Time lapse images of chip filled with sample (colored dye used to aid in visualization).

Supplementary tables

Table S1. Results from experiment to determine the time required to obtain a reproducible voltage reading from the photodiode.

Number of readings per measurement	Reading frequency (Hz)	Total time per measurement (ms)	Coefficient of variation for 4 replicate
			measurements (%)
10^{4}	10^{5}	234	0.017
10^{3}	10^{4}	156	0.037
5×10^{3}	10^{4}	140	0.082
5×10^{2}	10^{4}	93.0	0.087
10 ³	10^{5}	78.0	5.87
10	10^{3}	62.5	5.63
100	10^{4}	46.8	2.33
10	10^{4}	46.8	6.12
10	10^{4}	31.3	4.60

^a The output of a single measurement is calculated as the average of the different readings per measurement.

Table S2. LAMP primers used in this study.

Pathogen (Gene)	Primer	Sequence	
S. aureus (vicK)	F3	GTCAAGAAATTGGAGAAATTCG	
	B3	ACCAATTACCTTTTTATCGACTT	
	FIP	GGACAGAACTATCATTCGCTTTTTGGACCAAATTATTATTGCGACGAC	
	BIP	GCACTATCACTAGGACAATCAAACGATATTATATACCCAGACACGGT	
	LF	ATTGATTAGACTACGGTTAGACTGC	
	LB	TTAAAAGATTATGGCGGTGGTAAGG	
S. aureus (mecA)	F3	AAAAACGAGTAGATGCTCAA	
	B3	TGGCCAATTCCACATTGT	
	FIP	TCCCAATCTAACTTCCACATACCATAAAAACAAACTACGGTAACATTGA	
	BIP	CATAGCGTCATTATTCCAGGAATGCCGGTCTAAAATTTTACCACGT	
	LF	TTTAACAAAATTAAATTGAACGTTGCGA	
	LB	AGAAAGACCAAAGCATACATATTGAAAA	
E. coli (stx2)	F3	GAGATATCGACCCCTCTTG	
	B3	AATCTGAAA AACGGT AGA AAGT	
	FIP	TCCACAGCA AAATAACTGCCCAAC ATATATCTCAGGGGACCA	
	BIP	GATGTCTATCAGGCGCGTTTTGCCGTATTAACGAACCCG G	
	LF	TGT GGTTAATAACAGACACCGATG	
	LB	ACCATCTTCGTCTGATTATTGAGC	
E. coli (eaeA)	F3	AGCTCTAACAATGTACAGCT	
	B3	AGTTGCAGTTCCTGA AACA	
	FIP	GTCTTATCCGCCGTA AAGTCCGCCGTTCTGTCGAATGGTC	
	BIP	CTAAAGCGGATA ACGCCGATACCCAGGGACATTAGCCTGAG	
	LF	CCCAACCTGGTCGACAACTT	
	LB	ATTACTTATACCGCGACGGTGAA	