

# Development of a Smartphone-Integrated Reflective Scatterometer for Bacterial Identification

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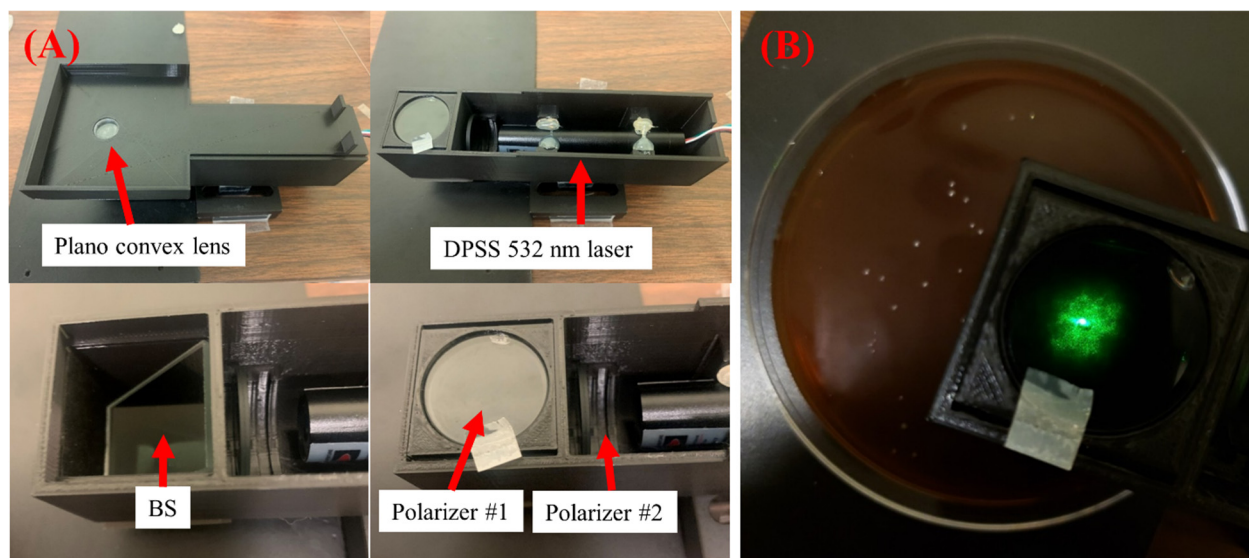
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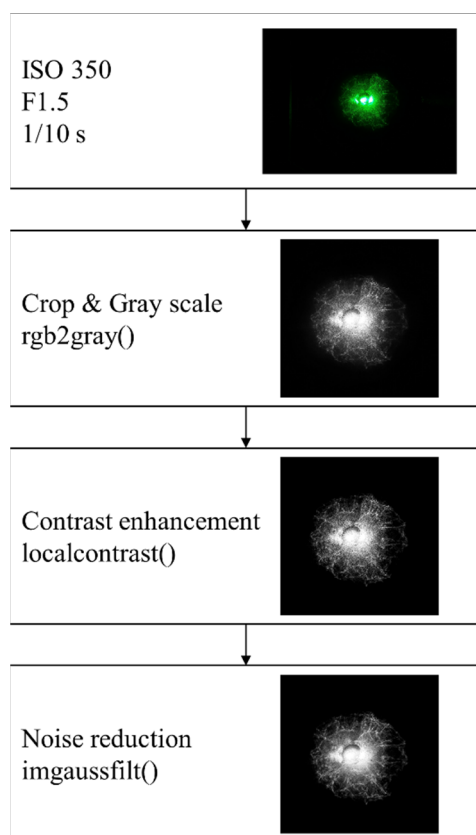
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## Supplementary Materials:

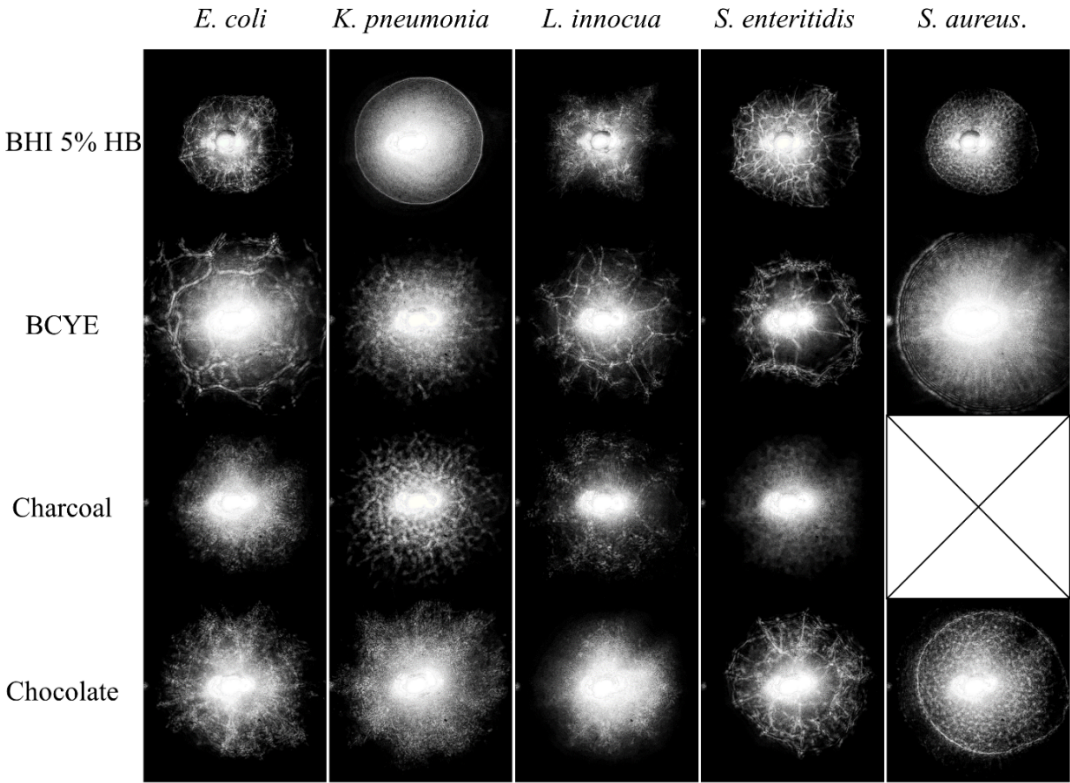


**Figure S1.** More pictures of the proposed instrument. (A) Pictures showing each component of the proposed instrument, (B) A picture of the instrument without the smartphone and top cover, measuring the reflective ELS pattern of *E. coli* colony on BHI with 5% horse blood agar.

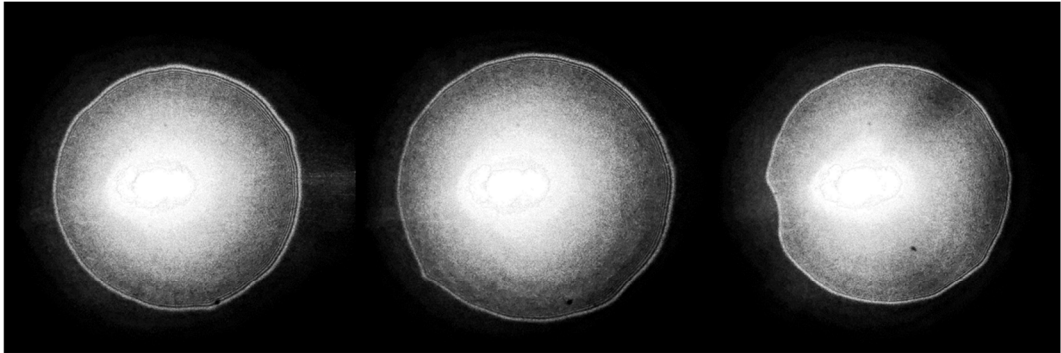


**Figure S2.** Image pre-processing procedure and corresponding setting or MATLAB functions utilized. Example pattern images showing the changes made from each step are attached.

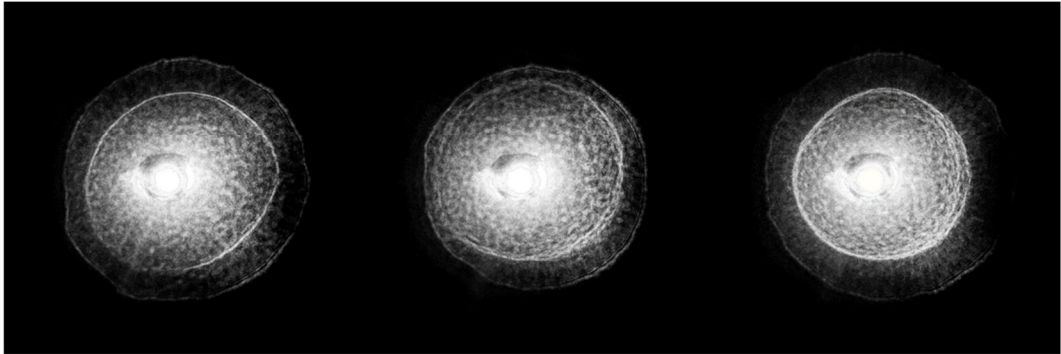
Matlab function, “`rgb2gray(RGB)`” converts the RGB image to grayscale image by eliminating the hue and saturation information while retaining the luminance. “`localcontrast()`”, is an edge-aware local contrast manipulation of images, which enhances or flattens the local contrast of an image by increasing or smoothing details while leaving strong edges unchanged. “`imgaussfilt()`” is a 2-D Gaussian filtering function that filters an image with a 2-D Gaussian smoothing kernel with specified standard deviation. Details of the Matlab functions can be found on Matlab website.



**Figure S3.** Representative reflective ELS pattern images of the five bacteria grown on various nutrition media. The rows represent the media type, and the columns represent the bacterial species.



**(A)**



**(B)**

**Figure S4.** The comparison between mucoid and non-mucoid bacterial colonies with their reflective ELS patterns. The reflective ELS patterns of *K. pneumoniae* that formed **(A)** mucoid and **(B)** non-mucoid colonies.

**Table S1.** List of the optical scattering/diffraction technologies for bacterial identification and their specifications.

	Laser wavelength	Pattern type	Target sample	Detector	Features	Classifier	Ref.
<i>Bae et al</i>	635 nm	Transmitted	Foodborne pathogen	CMOS	Haralick texture Zernike	SVM	[3]
<i>Marcoux et al</i>	543.5 nm	Transmitted	Clinical	CCD	Zernike invariant	Bayes Network Naïve Bayes SMO	[4]
<i>Minori et al</i>	Not specified	Transmitted	Urinary tract pathogen	Not specified	NA	NA	[5]
<i>Buzdlewicz et al</i>	635 nm	Transmitted	Clinical	CCD	Mean Entropy Radius Standard deviation Flatness Skewness	LDA QDA SVM	[6]
<i>Pan et al</i>	980 nm	Transmitted	Foodborne pathogen	Screen NIR camera	Zernike invariant	LDA KNN PLSDA BPANN	[8]
<i>Kim et al</i>	532 nm	Reflected	Foodborne pathogen	Screen DSLR camera	Haralick texture Zernike invariant	SVM	[9]

**Table S2.** The cross-validation matrices show the classification result of the five bacteria grown on BHI with 5% horse blood, BCYE, and chocolate agar plates.

BHI 5% HB	<i>E. coli</i>	<i>K.pneumoniae</i>	<i>L. innocua</i>	<i>S. enteritidis</i>	<i>S. aureus</i>
<i>E. coli</i>	97.1	0	0	2.9	0
<i>K. pneumonia</i>	0	100	0	0	0
<i>L. innocua</i>	0	0	100	0	0
<i>S. enteritidis</i>	3.3	0	0	96.7	0
<i>S. aureus</i>	0	0	0	0	100
BCYE	<i>E. coli</i>	<i>K.pneumoniae</i>	<i>L. innocua</i>	<i>S. enteritidis</i>	<i>S. aureus</i>
<i>E. coli</i>	83.9	0	9.2	0	6.9
<i>K. pneumonia</i>	0	93.7	1.8	0	4.5
<i>L. innocua</i>	9.8	0	90	0	0.2
<i>S. enteritidis</i>	0	0	2.1	97.9	0
<i>S. aureus</i>	0	0	0	0	100
Chocolate	<i>E. coli</i>	<i>K.pneumoniae</i>	<i>L. innocua</i>	<i>S. enteritidis</i>	<i>S. aureus</i>
<i>E. coli</i>	90.6	0.8	0	3.8	4.9
<i>K. pneumonia</i>	8.7	90.4	0	0.9	0
<i>L. innocua</i>	0	0.4	98.9	0.1	0.6
<i>S. enteritidis</i>	6.7	5	0	85.5	2.8
<i>S. aureus</i>	2.1	0	0	0.7	97.2

**Table S3.** The cross-validation matrix and the five statistical parameters for the four bacteria grown on charcoal agar plates. Since *S. aureus* did not grow on this media, only four bacteria were classified.

	<i>E. coli</i>	<i>K.pneumoniae</i>	<i>L. innocua</i>	<i>S. enteritidis</i>
<i>E. coli</i>	90.9	0	9.1	0
<i>K. pneumonia</i>	0	100	0	0
<i>L. innocua</i>	0	0.1	99.9	0
<i>S. enteritidis</i>	0	0	9.6	90.4
	<i>E. coli</i>	<i>K.pneumoniae</i>	<i>L. innocua</i>	<i>S. enteritidis</i>
Accuracy	97.7	100	95.3	97.6
Sensitivity	90.9	100	99.9	90.4
Specificity	100	100	93.8	100
PPV	100	99.9	84.2	100
NPV	97.1	100	100	96.9