

## Supporting Information

### **A low cost mobile phone dark-field microscope for nanoparticle-based quantitative studies**

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## Experimental Section

### Exosome quantification assay.

The AuNR probe was generated by adding 40  $\mu\text{L}$  of neutravidin-labeled AuNRs (Nanopartz C12-25-650-TN-50,  $7 \times 10^{-9}$  M) to 200  $\mu\text{L}$  PBS, then added 20  $\mu\text{L}$  biotinylated CD9 antibody (NB110-81616, Novus, 0.5mg/mL), and sonicated the mixture for 8 minutes at 80% amplitude using a 5 second on/off cycle (Q500 Sonicator, Qsonica) to accelerate conjugation. AuNR samples were then centrifuged at  $4000 \times g$  for 10 min, aspirated, resuspended in 400  $\mu\text{L}$  PBS and washed twice more using the same protocol, then resuspended in 200  $\mu\text{L}$  PBS and stored at  $4^\circ\text{C}$  until use. Protein A/G-modified 192-well slides (Arrayit) were incubated with 1  $\mu\text{L}$ /well anti-LAM IgG (courtesy of BEI Resources) and sonicated at  $37.5^\circ\text{C}$  for 8 minutes at 80% amplitude using a 5 second on/off cycle (Q500 Sonicator, Qsonica) to accelerate IgG binding. Slides were then blocked with 1  $\mu\text{L}$ /well 5% BSA (Thermo Scientific) for 1 hour at  $37.5^\circ\text{C}$  and PBS-washed 3 times before use. Slides were loaded with 1  $\mu\text{L}$ /well of 2-fold PBS-diluted serum, PBS-washed times, and incubated with anti-CD9 IgG-conjugated-AuNRs for 1 hour at  $37.5^\circ\text{C}$  in a humidified chamber, then washed for 10 min at  $25^\circ\text{C}$  with PBS and 0.01% Tween-20 (PBST, pH 7.0), rinsed with deionized water, and air-dried for DFM imagery.

## Tables

**Table S1.** Cost list of the MDFM.

Item	Supplier	Cost (\$)
Objective Lens (10×/20×)*	Olympus	200 / 400
Condenser	Olympus	920
Case	3dhubs.com	60
Mobile Phone	Motorola	180
Total		1360 / 1560

\* Lens details in Table S3

**Table S2.** MDFM and DDFM inter-assay and intra-assay reproducibility.

Assay	Concentration μg/μL	Precision (% CV)			
		Inter-assay		Intra-assay	
		MDFM	DDFM	MDFM	DDFM
Binding affinity	0.091	13.65	5.51	5.11	5.52
	0.023	11.69	3.86	3.76	4.20
	0.005	9.83	3.89	3.20	1.60
Protein quantification	0.083	8.06	6.02	3.06	2.14
	0.021	10.89	5.74	7.37	2.30
	0.005	13.32	8.02	7.79	7.72

**Table S3.** Objective lens specification for MDFM.

	10×	20×
Manufacturer	Olympus	Olympus
Model	CACHN10XPHP	LCACHN20XPHP
Numerical Aperture	0.25	0.4
Working Distance (mm)	8.8	3.2
F Number	22	22

**Table S4.** Demographic and Clinical data of the exosome analysis cohort. The HTI cohort that served as a source of case and control samples in this study were the subjects of a large, population-based TB surveillance study that performed active surveillance of all confirmed TB cases in Houston/Harris County, Texas between 1995 and 2004.

Group	Labels	Age	Gender	TB Diagnosis	Culture Positive	Skin Test Positive
TB	1153.01	16	M	Yes	No	Yes
	1150.01	13	F	Yes	No	Yes
	1133.01	17	M	Yes	Yes	Yes
	1121.01	9	M	Yes	No	Yes
	1109.01	16	F	Yes	Yes	Yes
Healthy	1113.05	9	M	No	N/A	N/A
	1139.04	17	F	No	N/A	N/A
	1076.10	11	M	No	N/A	N/A
	1170.05	8	F	No	N/A	N/A
	1182.04	4	F	No	N/A	N/A

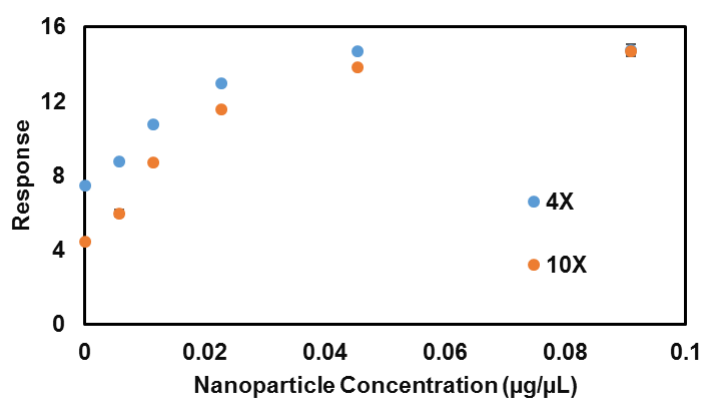
**Table S5** Summary of mobile phone microscope studies.

#	Researchers	Target	Application	Illumination	Field
<b>Pathogens</b>					
1	(Pirnstill and Coté, 2015)	Malaria parasites	I	BF	NF
2	(Koydemir et al., 2015)	Giardia parasites	I & Q	BF	FF
3	(D'Ambrosio et al., 2015)	Filarial parasites	I & Q	BF	FF
4	(Ame et al., 2013)	Helminth parasites	I	BF	FF
<b>Cells</b>					
5	(Phillips et al., 2015)	Cells	I	BF & DF	FF
6	(Switz et al., 2014)	Epithelial cell	I	BF	FF
7	(Skandarajah et al., 2014)	Red blood cells	I & Q	BF	FF
8	(Lee and Yang, 2014)	Cells	I	BF	FF
9	(Breslauer et al., 2009)	Blood cells	I	BF	FF
<b>Biomolecules</b>					
10	(Wang et al., 2017)	BSA	I & Q	BF	FF
11	(Archibong et al., 2017)	Hemoglobin	I & Q	BF	FF
12	(Lee et al., 2017)	estradiol	I & Q	BF	FF
13	(Khan et al., 2015)	Glucose	Q	BF	FF
14	(Berg et al., 2015)	HSV-1 IgG	Q	BF	FF
15	(Lee et al., 2011)	DNA	Q	BF	FF
16	(Zhang et al., 2011)	DNA	I & Q	BF	FF
<b>Particles</b>					

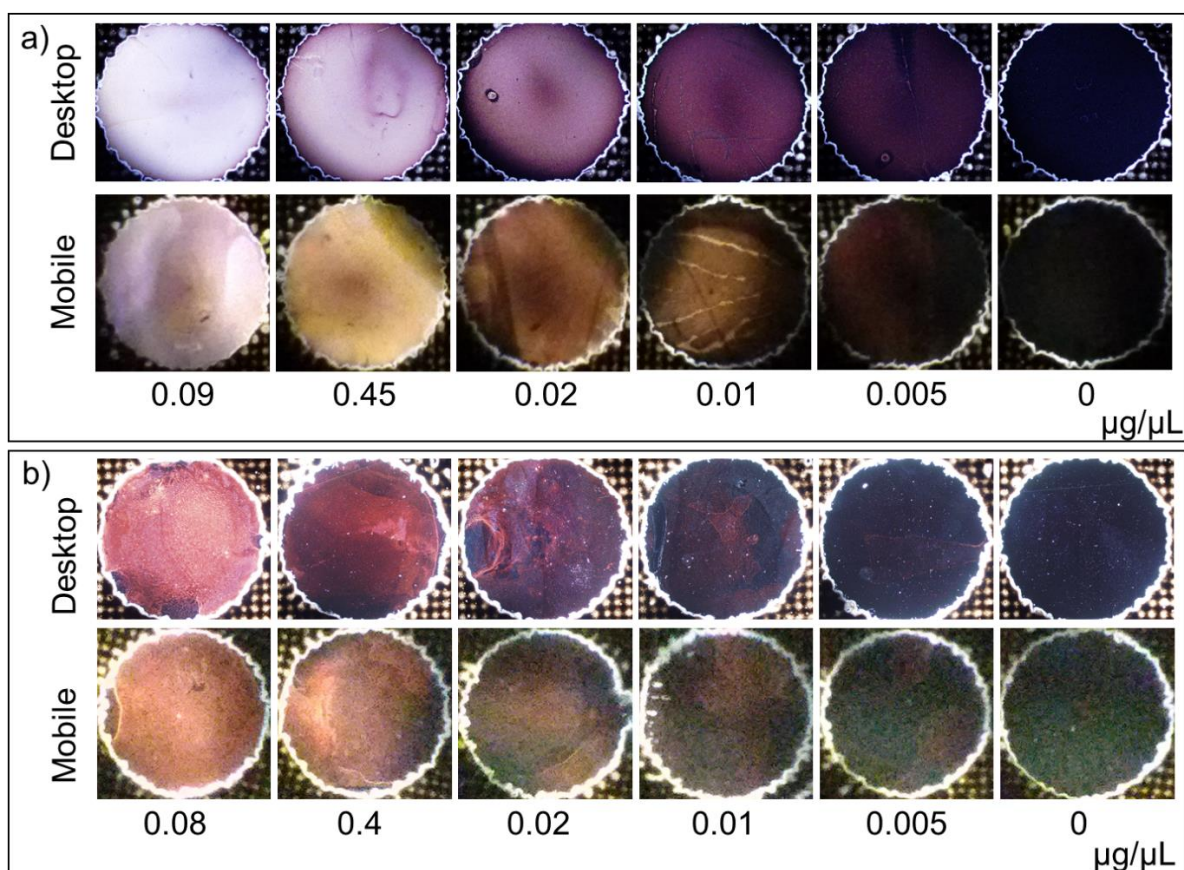
17	(Wei et al., 2013)	Fluorescent particle	I & Q	BF	NF
18	(Preechaburana et al., 2011)	Gold Particle	I & Q	BF	FF
19	(Lu et al., 2009)	Gold Particle	I & Q	BF	FF

I: imaging; Q: quantification; BF: bright-field; DF: dark-field; NF: near-field; Far-field: FF

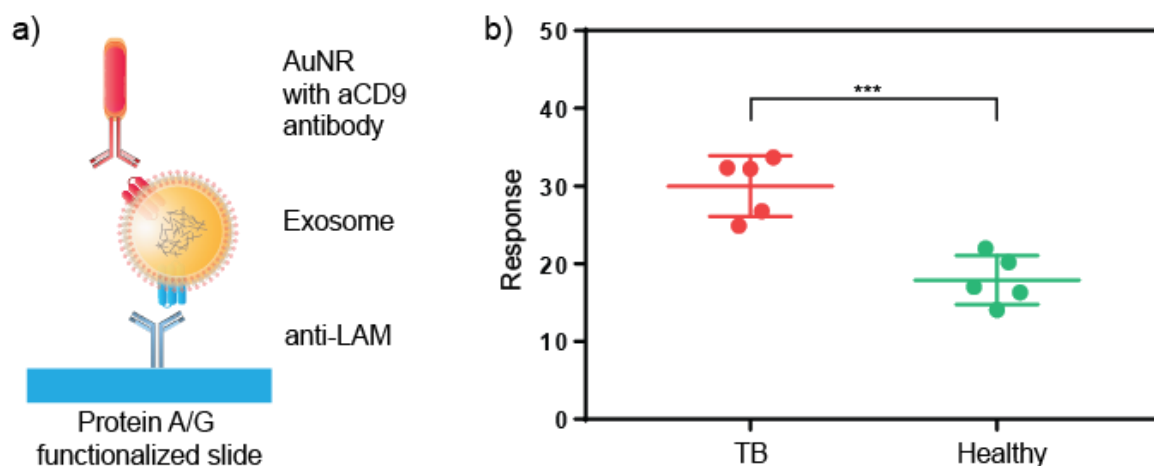
## Figures



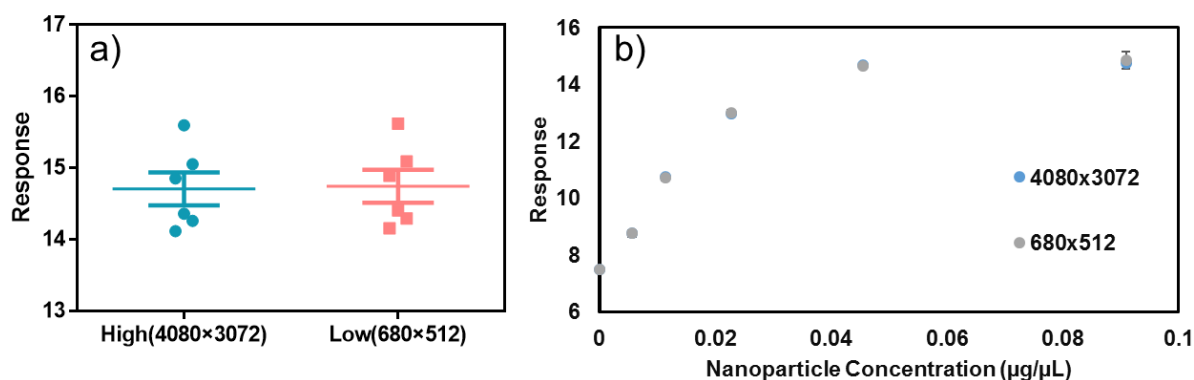
**Figure S1.** DSM response comparison of different magnification (4X and 10X objective lens) under DFM high resolution (4080x3072 px) for binding affinity assay same as shown in Figure 4a.



**Figure S2.** Representative images by MDFM and DDFM for a) binding affinity assay (b) protein quantification assay along with nanoparticle and target concentration gradient respectively.



**Figure S3.** Exosome quantification scheme. Antibody-labeled detection wells bind exosomes derived from cells that have internalized *Mycobacterium tuberculosis* (*Mtb*) bacilli by binding the *Mtb* antigen lipoarabinomannan (LAM). These *Mtb*-derived exosomes are mixed with a nanoparticle-labeled probe that binds CD9, a pan-specific exosome biomarker, and quantified by DFM image analysis. (b) DSM-quantified *Mtb* exosome concentrations in serum samples of pediatric patients with tuberculosis (TB) and healthy controls. Data are presented as mean  $\pm$  SEM, N=5/group; \*\*\*p=0.0007 by two-sided Mann-Whitney U-test. Table S4 is the demographic information of the cohort.



**Figure S4.** Comparison of quantification results from high (4080x3072) and low (680x512) resolution DDFM images at (a) a single concentration point (0.023  $\mu\text{g}/\mu\text{L}$ ) and (b) across multiple concentration values (0  $\mu\text{g}/\mu\text{L}$  to 0.09  $\mu\text{g}/\mu\text{L}$ ) for the assay scheme depicted in Figure 4a. Graphs depict Mean  $\pm$  SD values for each concentration value. The mean ratios of paired low-resolution to high-resolution response values for a) and b) were  $0.9976 \pm 0.0005$  and  $0.9992 \pm 0.0036$ , respectively.

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