

# Microfluidic Measurements of Protein Biomarkers from Exhaled Breath

## BEAD-BASED ASSAYS

- Large surface area for capturing analytes from biological samples
  - $>10^4$  beads per sample
  - Amenable to protein or nucleic acid analytes
  - High capture efficiency
- Magnetic core of beads facilitates sample preparation
  - Beads can be controlled with a magnetic field
    - Easy to translate or capture with a simple magnet
- Low sample volume requirement ( $\leq 100 \mu\text{L}$ )
  - Amenable for point-of-care analyses

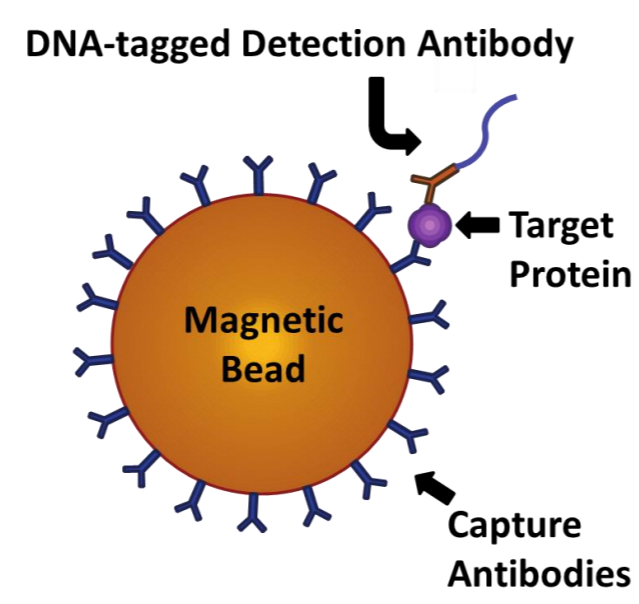


Figure 1. Illustration of an immunoPCR complex.

## MICROWELL ARRAYS

- Compatible with bead-based assays
- Highly sensitive amplification-based detection
  - PCR and ELISA formats
- Enables single molecule detection
  - One bound target produces a locally high dye concentration ( $\sim 1 \mu\text{M}$ ) for facile fluorescence detection
- Digital detection mode provides LODs up to 1,000x superior to conventional bioassays
- Quantitation achieved by determining the percentage of active wells in the array
  - Correlates to the analyte concentration in the sample
    - Poisson statistics

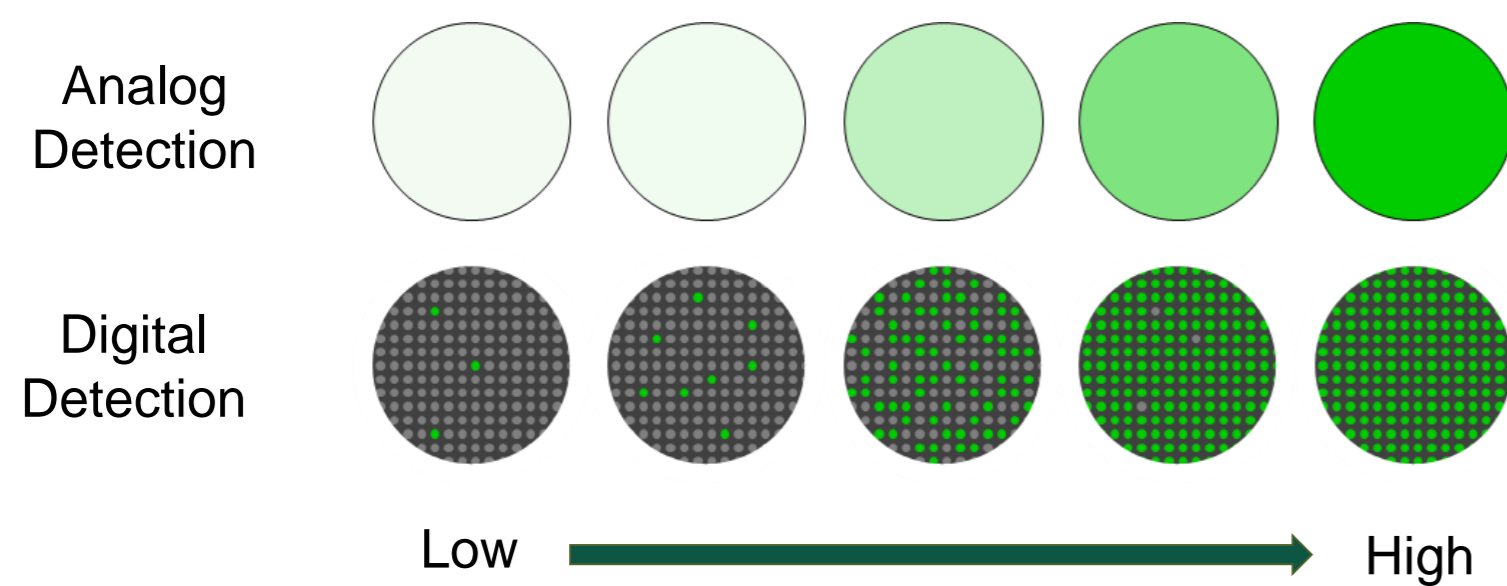


Figure 2. Illustration comparing a simulated fluorescent signal observed as a function of analyte concentration. Analog detection (top) yields a continuum response whereas digital detection (bottom) produces increasing numbers of individual "active" wells. Adapted from *Anal. Chem.* **2013**, *85*, 1258-1263.

## MICROCHIP DESIGN

- Microwells etched into a silicon substrate
- A glass coverplate is etched to dictate channel height
- Wells densely packed into the array
  - $\sim 10^6$  wells/cm<sup>2</sup>
- Wells engineered to hold only a single bead
  - $3.0 \mu\text{m}$  diameter regions for bead loading
- Bead loading regions separated from signal acquisition regions
  - $70 \text{ fL}$  fluidic region adjoined for signal integration
  - Prevents bead autofluorescence from interfering with assay signal

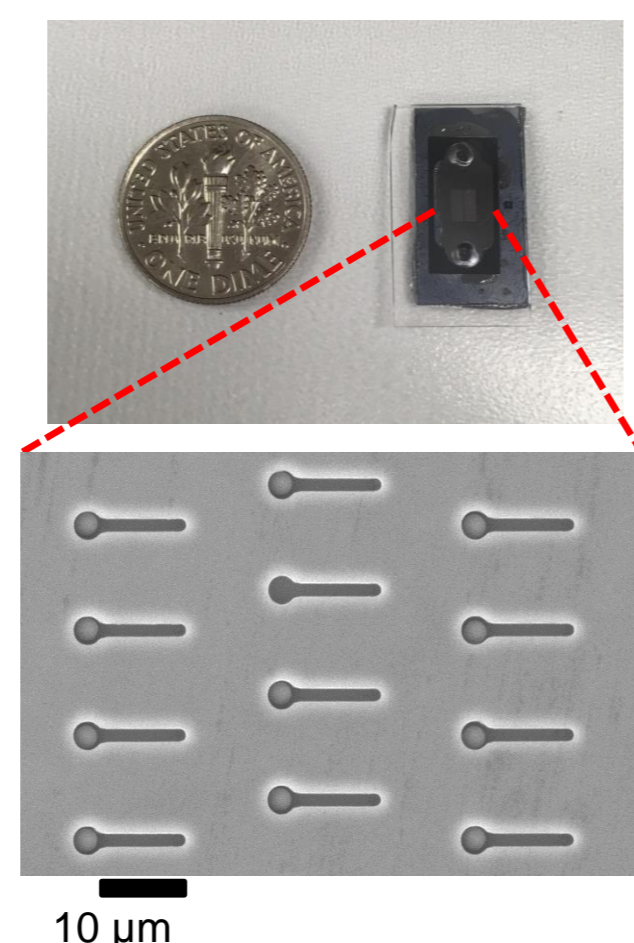


Figure 3. Photo of a microfluidic microwell array device. Inset shows a SEM image of beads loaded into the array. Adapted from *Anal. Bioanal. Chem.* **2020**, *412*, 6917-6926

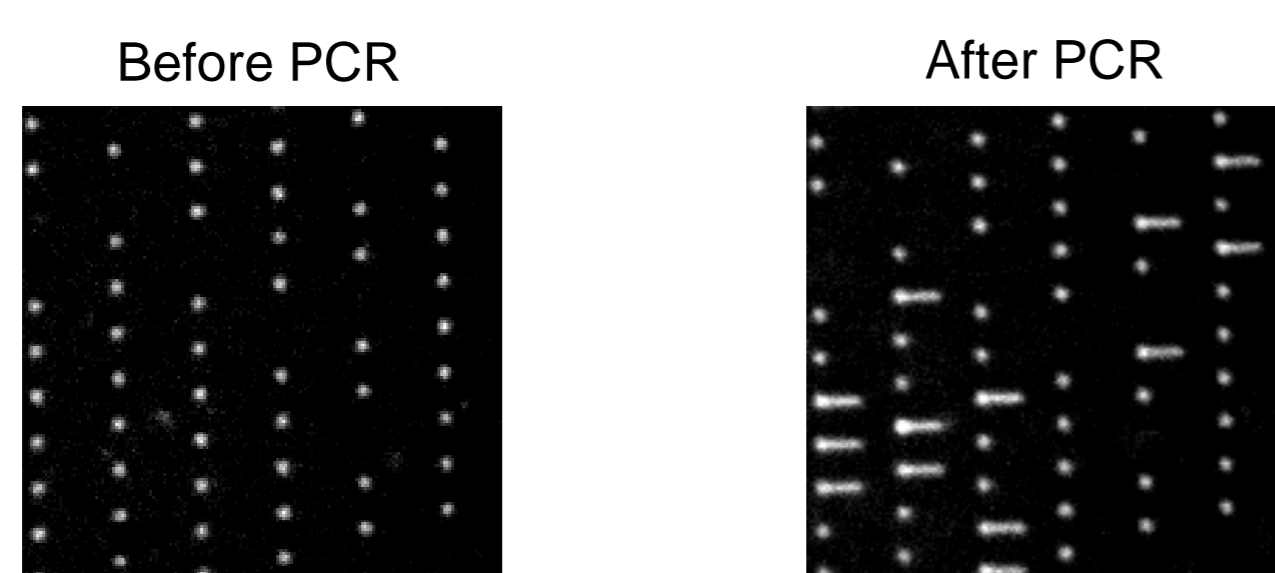


Figure 4. Images of a zoomed in region of a device before and after PCR amplification. Wells containing a single target biomarker exhibit intense fluorescence after PCR.

## PROTEIN ASSAYS

- The cytokines GM-CSF, IL-6, IL-36 $\gamma$  were used as model proteins
  - Involved in inflammation pathways and upregulated in certain cancers
- Limits of detection lower than conventional assays
  - GM-CSF:  $4 \text{ fM}$
  - IL-6:  $30 \text{ fM}$
- Multiplexing achieved by encoding beads targeting different cytokines with different amounts of fluorescent dye
  - Used a fluorescent protein for encoding
    - R-phycoerythrin
    - Small molecule dyes found to inhibit PCR amplification
- Achieved three distinct bead populations
  - No overlap in fluorescence between populations
- Encoded beads compatible with PCR
  - Beads encoded with  $50 \text{ nM}$  PE exhibited strong PCR amplification

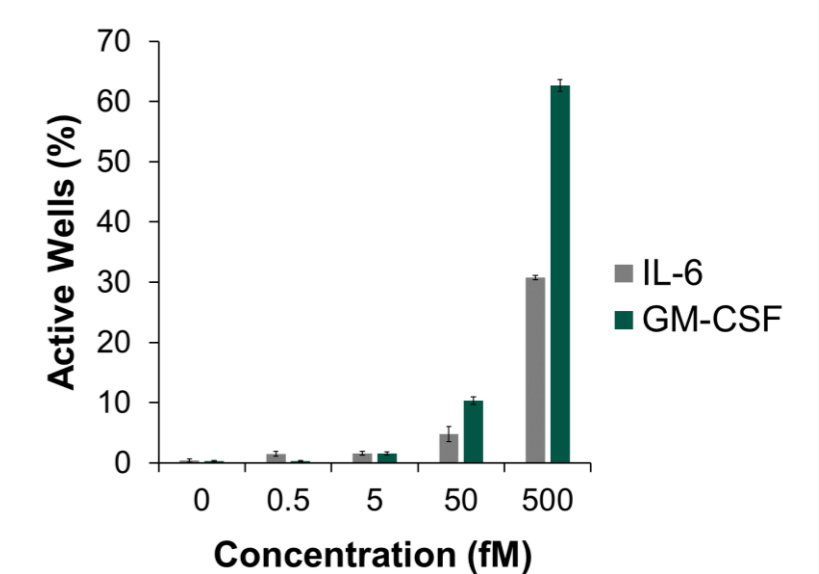


Figure 5. Standard curves for GM-CSF and IL-6 analyzed using iPCR.

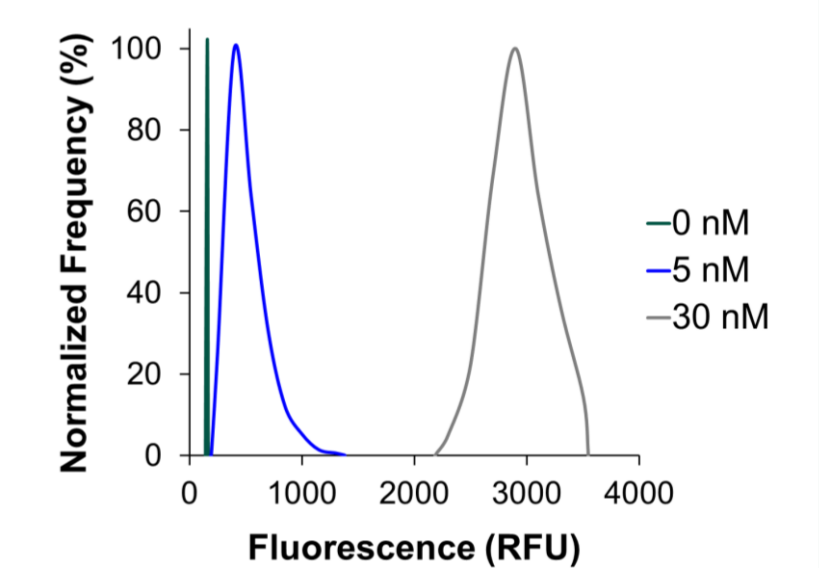


Figure 6. Fluorescence distributions from encoded bead populations.

## EBC ANALYSIS

- Exhaled breath condensate (EBC) was collected in chilled vials
  - $100 \mu\text{L}$  collected from healthy volunteers
- Sample validation required to screen for saliva contamination
- Amylase assays showed that EBC contained no detectable amylase, unlike saliva
  - Verified that sample originated from the lungs, not the mouth



Figure 7. Exhaled breath condensate was collected for analysis.

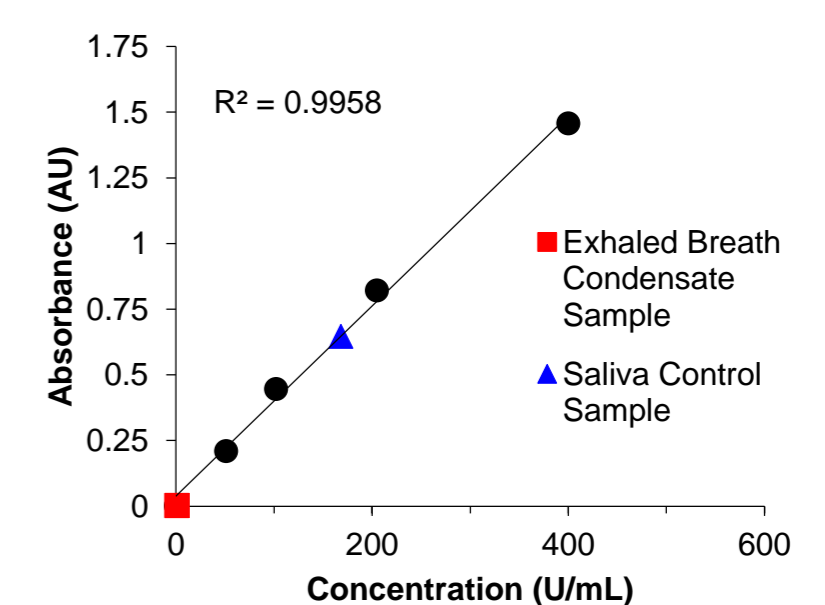


Figure 8. Amylase analysis to screen for saliva contamination.

- Multiplexed bead assays were conducted to measure each analyte in exhaled breath
  - Incubate  $80 \mu\text{L}$  EBC with beads for capture
- Beads in each well were identified by measuring fluorescence at the decoding wavelength
  - Three encoded bead populations for three different target proteins
- Cytokines were quantified from EBC samples
  - GM-CSF:  $110 \text{ fM}$
  - IL-6:  $5 \text{ fM}$
  - IL-36 $\gamma$ :  $400 \text{ fM}$
- Cytokines in EBC below the detection limit of conventional methods
  - LOD ELISA:  $500 \text{ fM}$

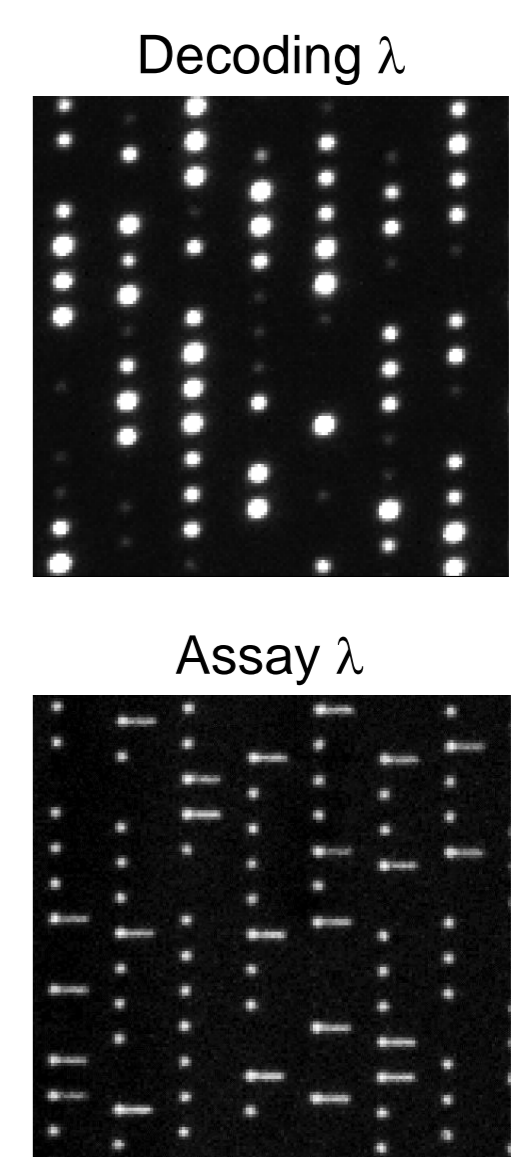


Figure 9. Images of the same region of a microwell array device imaged at the decoding (top) and assay (bottom) wavelengths.

## AKNOWLEDGEMENTS

- Society for Analytical Chemists of Pittsburgh Starter Grant
- Wayne State University startup funding

