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## Integrated Magneto-Electrochemical Sensor for Exosome Analysis

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## Abstract

Extracellular vesicles, including exosomes, are nanoscale vesicles that carry molecular information of parental cells. They are being pursued as biomarkers of cancers that are difficult to detect or serially follow. Here we present a compact sensor technology for rapid, on-site exosome screening. The sensor is based on an integrated magnetic-electrochemical assay: exosomes are immunomagnetically captured from patient samples, and profiled through electrochemical reaction. By combining magnetic enrichment and enzymatic amplification, the approach enables i) highly sensitive, cell-specific exosome detection, and ii) sensor miniaturization and scale-up for high throughput measurements. As a proof-of-concept, we implemented a portable, eight-channel device, and applied it to screen extracellular vesicles in plasma samples from ovarian cancer patients. The sensor allowed for the profiling of multiple protein markers simultaneously within an hour, outperforming conventional methods in assay sensitivity and speed.

## Keywords

extracellular vesicles; electrochemical sensing; cancer; diagnostics; point-of-care

Growing evidence has positioned extracellular vesicles (EVs) as an effective readout of cancer management<sup>1-6</sup>. Exosomes, in particular, have emerged as a potent biomarker. Exosomes are nanoscale vesicles actively secreted by cells. These vesicles carry molecular constituents of their originating cells, including transmembrane and cytosolic proteins<sup>7</sup>,

Conflict of Interest

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mRNA<sup>8</sup>, DNA<sup>9</sup> and microRNA<sup>10</sup>, and can thus serve as cellular surrogates<sup>11</sup>. Combined with their relative abundance and ubiquitous presence in bodily fluids (*e.g.* serum, ascites, urine, CSF)<sup>12-14</sup>, exosomes can offer unique advantages for longitudinal monitoring<sup>15-17</sup>. Exosome analyses are minimally invasive and afford relatively unbiased readouts of the entire tumor burden, less affected by the scarcity of the samples or intra-tumoral heterogeneity<sup>5</sup>.

Analyzing extracellular vesicles in routine clinical settings, however, still remain a difficult task, mainly due to the lack of adequately sensitive and fast assay platforms, particularly for protein analyses. Flow cytometry provides high throughput detection<sup>18</sup>, yet tends to miss small vesicles (<200 nm) because of weak light scattering<sup>19</sup>. New counting approaches based on particle-tracking or dynamic light scattering can offer more accurate particle counts, >10<sup>3</sup>-fold higher than those reported by flow cytometry<sup>20</sup>, but yield limited molecular information. Conversely, conventional molecular assays (*e.g.* Western blot, ELISA) require large amounts of samples, and become impractical for clinical research needs, notably serial analyses, large patient cohorts, or limited specimens in biorepositories.

We have previously developed miniaturized systems to facilitate EV analyses in clinical environments. These systems include microfluidic devices for sample preparation<sup>21, 22</sup> and analytical tools for protein analyses<sup>3, 5</sup>. Ensuing clinical studies with patient samples established EVs' clinical utility for cancer diagnosis and treatment monitoring<sup>3, 5</sup>: tumor-specific exosomes could be identified based on their unique transmembrane protein signature, and exosomal protein changes could be correlated with treatment responses. Translating these systems into routine clinical tests, however, is still limited by i) separate procedures and devices required for exosome isolation and detection; ii) technical complexities involved in device fabrication and operation; and iii) relatively high costs of analytical instruments (*e.g.* NMR detector, spectral microscope).

We reasoned that electrochemical sensing could be an effective detection modality that is easily applicable to clinical settings. Electrochemical sensing could achieve high sensitivity through signal amplification with redox-active reporters<sup>23-28</sup>. The readout system measures electrical currents, and can be realized as a compact and low-power portable device<sup>29, 30</sup>. Harnessing electrochemical sensing for exosome analyses, however, is challenged by both assay design and hardware development: i) combining exosome isolation and detection in a single platform; ii) providing fast, high-throughput assays for multiplexed screening; and iii) making the system affordable and easy-to-use.

We herein report on a new integrated approach, iMEX (integrated magnetic-electrochemical exosome), for fast, streamlined exosome analyses. The iMEX combines two orthogonal modalities, magnetic selection and electrochemical detection. Magnetic beads are used for exosome capture and labeling; captured exosomes are detected via electrochemical sensing. The iMEX offers distinct advantages: i) cell-specific exosomes can be isolated directly from complex media without need for extensive filtration or centrifugation; ii) the assay can achieve high detection sensitivity through magnetic enrichment and enzymatic amplification; iii) through the electrical detection scheme, sensors can be miniaturized and expanded for parallel measurements. To prove the concept, we implemented a portable

iMEX platform with eight detection channels. The iMEX was able to detect exosomes at a sensitivity of  $<10^5$  vesicles. Importantly, it only used  $10 \,\mu$ L of samples and generated readouts within 1 hour. We further explored iMEX's potential clinical utility by profiling EVs collected from ovarian cancer patients. With its capacities for fast, high-throughput, and on-spot analysis, the iMEX could accelerate the transition of EV and exosome analyses towards routine clinical testing.

## **RESULTS AND DISCUSSION**

#### **iMEX** platform

The miniaturized iMEX system had eight independent channels (Figure 1A). Each channel was equipped with a potentiostat capable of measuring a wide range of current ( $\pm 7.5 \,\mu$ A). The input signal was conditioned by a low-pass filter (cut-off frequency, 5 Hz) to suppress high frequency noise (Supplementary Figure 1). The eight potentiostats were connected to a digital-to-analog converter for potential control, an analog-to-digital converter for signal digitization, a multiplexer for channel selection, and a micro-controller unit for system operation (Figure 1B; Supplementary Figure 2). We packaged the device as a handheld unit (Figure 1C). A card-edge connector was used for the quick attachment of the electrode cartridge. A magnet holder, containing 8 cylindrical magnets, was placed underneath the electrode cartridge. These magnets are used to concentrate magnetic beads to the sensor surface. The overall device cost was <\$50. The system performance was comparable to that of commercial equipment (SP-200, Bio-Logic; Supplementary Figure 1B), however at much smaller footprint and cost. Furthermore, the iMEX effectively provided a simultaneous readout from all electrodes through rapid polling of each channel (50 msec per channel). All data were monitored and analyzed by custom-designed software (Supplementary Figure 3).

**Figure 1D** summarizes the iMEX assay scheme. Exosomes are first captured onto immunomagnetic beads. Secondary antibodies with an oxidizing enzyme (horseradish peroxidase; HRP) are then used, followed by mixing the beads with chromogenic electron-mediators (3,3',5,5'-tetramethylbenzidine; TMB) which generate electrical current when HRP is encountered (**Supplementary Fig. 4**; see **Methods and Experimental** section for details). Using magnetic beads significantly simplifies the assay procedures: excess agents (*e.g.* antibodies, enzymes) can be removed via magnetic washing, and captured exosomes can be magnetically concentrated on the electrodes to improve the detection sensitivity.

#### Assay optimization

We optimized the iMEX assay protocol. We applied the chronoamperometry method for signal detection: the electrical current generated from TMB reduction was monitored while a reduction potential (-100 mV versus Ag/AgCl reference electrode) was applied to a working electrode. The current level (*I*) reached a plateau within 1 minute after the reduction potential was applied (**Figure 2A**). We averaged the current level (*I*) from 40 to 45 seconds as a representative value.

To capture exosomes, we used magnetic beads that were coated with antibodies against tetraspanin, transmembrane proteins enriched in exosomes<sup>5, 31, 32</sup>. We first compared signal levels with differently sized beads (diameters, 2.7  $\mu$ m and 8.8  $\mu$ m). When the total surface area of beads was matched to capture similar amount of exosomes, the measured signal levels were nearly identical (**Supplementary Figure 5A**). This result can be explained with diffusivity in porous media: The effective diffusivity ( $D_e$ ) for stacked beads can be expressed as  $D_e = D_0 \bullet \varepsilon^m$ , where  $D_0$  is the diffusivity in free media, and  $\varepsilon$  is the porosity of the structure<sup>48</sup>. In case of uniformly sized beads, both  $\varepsilon$  ( $\geq$ 0.47) and m (= 3/2) are bead-size independent; the iMEX signals are thus expected to remain constant. We opted to use 2.7- $\mu$ m beads; bigger beads tended to sediment, requiring frequent shaking of samples. Compared to the no-enrichment scheme, magnetic enrichment led to ~72% increase in the analytical signal (**Supplementary Figure 5B**).

We next tested three representative tetraspanin proteins (CD63, CD9, CD81) as a target; these markers are reportedly enriched in exosomes<sup>13, 33-39</sup>. We prepared 2.7- $\mu$ m magnetic beads specific to each marker (see **Methods and Experimental** section for details). When applied to exosomes from different cell lines, CD63-based capture showed consistently high signal (**Figure 2B**); we thus opted to use CD63 as a marker for exosome enrichment.

For each target marker (M), we prepared a pair of magnetic beads: one conjugated with antibodies against CD63 (CD63-beads) and the other with antibodies against iso-type matched IgG (IgG-beads). Exosomes were mixed with each bead-type and subsequently labeled with antibodies against a target marker; the net signal difference  $\Delta I_{\rm M}$  (=  $I_{\rm CD63+M} - I_{\rm IgG+M}$ ; Figure 2A) was then obtained. We used  $\Delta I_{\rm CD63}$  to estimate the total exosome load, and defined a normalized metric  $\xi_{\rm M}$  (=  $\Delta I_{\rm M}/\Delta I_{\rm CD63}$ ) as the expression level of a target marker (M). Note that such scaling would compensate for variations in exosome numbers among samples.

#### iMEX validation

We applied the developed iMEX protocol to profile exosomes for transmembrane proteins. For this validation study, we harvested exosomes from cell culture (OV90, OVCA420) through a conventional method (see **Methods and Experimental** section for details), and spiked them into phosphate buffered saline (PBS) solution (~10<sup>9</sup> exosomes/mL). Samples were aliquoted, and processed by iMEX and enzyme-linked immunosorbent assays (ELISA). Comparative analysis showed high correlation between two methods (**Figure 2C**;  $R^2 = 0.931$ ), confirming iMEX's analytical capacity. The iMEX assay, however, was faster (1 hr) and consumed smaller amounts of samples (10  $\mu$ L) than ELISA (5 hr, 100  $\mu$ L).

We further tested iMEX for detecting exosomes in biofluids. Cancer exosomes were collected from cell culture (OV90), and varying numbers of exosomes were spiked into undiluted human plasma. Titration experiments established the limit of detection (LOD) of  $3 \times 10^4$  exosomes, with the dynamic ranges spanning four orders of magnitude (**Figure 2D**). Similar measurements with ELISA required more than  $10^7$  exosomes for reliable detection. Using matched controls (IgG-beads) was important to compensate for background signals from sample-dependent, non-specific exosome binding.

#### Profiling of protein markers in cell derived exosomes

We applied the iMEX to screen exosomal surface markers from a panel of ovarian cancer cell lines. Because iMEX enriches CD63-positive (CD63+) exosomes and labels them for target proteins, we were able to examine how closely CD63+ exosomes reflect their cells of origin. We chose six representative surface markers based on prior studies: epithelial cell adhesion molecule (EpCAM)<sup>40</sup>, CD24<sup>41</sup>, cancer antigen 125 (CA125)<sup>42, 43</sup>, human epidermal growth factor 2 (HER2)<sup>44, 45</sup>, mucin 18 (MUC18)<sup>46</sup>, and epidermal growth factor receptor 2 (EGFR)<sup>47</sup>. The cellular expression levels of these markers were measured with flow cytometry; exosomes were harvested from the conditioned cell-culture media, and profiled with iMEX. The molecular profiles of cells and CD63+ exosomes were highly correlated (**Figure 3**), which supported the use of exosomes as cellular surrogates.

#### Clinical: Direct analyses of plasma from patients with ovarian cancer

The iMEX assay isolates EVs directly from plasma or serum, and allows profiling in a rapid, high-throughput manner — key for successful integration into the clinical workflow. To demonstrate clinical feasibility, we customized the iMEX assay for ovarian cancer EV detection in blood (**Figure 4A**). Clinical plasma samples were aliquoted without any purification, and each aliquot (10  $\mu$ L per marker) was incubated with magnetic beads for EV capture (15 min), followed by magnetic washing. The bead-bound EVs were consecutively labeled for target markers (15 min) and HRP (15 min), and loaded onto the device. With the 8-electrodes independently operated, we were able to *simultaneously* measure four different markers (CD63, EpCAM, CD24, and CA125) along with their respective IgG-controls. The IgG-controls were beneficial to specific detection of target molecules among non-purified clinical samples.

We tested single-time point plasma samples from 11 ovarian cancer patients and five healthy controls. The expression levels of EpCAM and CD24 in EVs were much higher in ovarian cancer patients than healthy controls, and both metrics showed high correlation (**Figure 4B**, **Supplementary Figure 6**). We next examined iMEX's potential for serial EV testing by measuring EpCAM and CD24 in plasma collected at two time points (2 months apart) from four ovarian cancer patients undergoing drug treatment. The iMEX assays were conducted blinded to treatment response. For 'non-responding' patients, expression levels of EpCAM and CD24 increased while 'responding' patients displayed a significant decrease in both markers (**Figure 4C**). The level of CD24 showed steeper increases than that of EpCAM for non-responders.

## CONCLUSION

We developed the iMEX technology to aid the translation of exosome analysis into clinical settings. A unique feature of iMEX is the integration of vesicle isolation and detection into a single platform. The use of magnetic actuation simplifies vesicle isolation and subsequent assay steps, and the electrochemical sensing facilitates high-throughput screening and sensor miniaturization. The current study validated these concepts: i) a portable detection system was implemented with the capacity for parallel measurements; ii) the iMEX enriched exosomes directly from blood, and profiled them for molecular information; iii) the entire

iMEX assay (i.e., exosome isolation, labeling, detection) was completed within 1 hour while consuming only  $10 \,\mu$ L of plasma per marker. We also demonstrated iMEX's clinical potential by profiling EVs within blood collected from ovarian cancer patients.

The bead-based magnetic enrichment brings several advantages in the iMEX measurement. First, the method provides a convenient way of concentrating signal sources on the electrodes, which enhances the detection sensitivity. Second, compared to the surface-based capture wherein antibodies are immobilized on the chip surface, the bead-based method is amenable to reliable and simpler conjugation chemistry, and benefits from faster binding kinetics between antibodies and exosomes. Third, the bead-bound vesicles could be readily recovered for downstream molecular analyses in tandem with iMEX. For instance, beadbound EVs can be eluted<sup>31</sup> or lysed<sup>3</sup> to profile their nucleic acid contents.

The iMEX system can complement the other exosome analysis platforms we have developed (**Supplementary Table 1**), particularly the nano-plasmonic exosome (nPLEX) sensor<sup>3</sup>. The nPELX is well suited to conduct massively parallel exosome screening, potentially for central clinical laboratories; it has high detection sensitivity, down to ~ $10^3$  exosomes, and is equipped with >1000 detection sites<sup>3</sup>. The system complexity and the requirement for nanofabrication, however, is a limiting factor for nPLEX's routine clinical applications. The iMEX system has lower sensitivity and throughput than nPLEX, but is affordable and miniaturized for on-site exosome detection.

In the current work, we focused on profiling CD63+ EV population (exosomes), which was motivated by two factors: i) the signal from CD63 capture was the highest (**Figure 2B**) among the tetraspanin markers tested; and ii) we and others have previously shown that ovarian cancer exosomes are enriched with CD63<sup>3, 35</sup>. The iMEX profiling found a high correlation in protein expression between CD63-positive exosomes and their parent cells (**Figure 3**); this result validated the potential use of CD63-positive exosomes as cellular surrogates. However, we note the need to extend the exosome-capture strategy, considering that diverse EV types (*e.g.* CD63-negative) may exist in patient samples. Examining these populations could yield more precise information to capture tumor heterogeneity. The iMEX method can be readily adopted for such purposes by changing capture antibodies.

We envision multiple directions to further advance the iMEX technology. First, the assay throughput can be improved by increasing the number of detection sites. Electrochemical sensing is ideally suited for such a scale-up: the sensing elements (electrodes) can be readily microfabricated into a large array format, and signals (electrical currents) can be read out by compact electronics with high-speed multiplexing. Second, the detection sensitivity could be improved by exploring new designs for electrochemical signal detection. The signal level is correlated with the surface area of a sensor and the amount of enzyme bound to target entities; thus, higher sensitivity can be achieved by using a nanostructured sensor surface or multi-label nanoparticles<sup>24, 49, 50</sup>. Third, detection targets can be expanded to include other exosomal constituents. For example, exosomes carry various nucleic acids (*e.g.* mRNA, microRNA); analyzing nucleic acids along with exosomal proteins would provide more accurate snapshot of tumor states<sup>6, 8, 9, 51</sup>. Electrochemical sensing has been applied to detect a trace amount of nucleic acids (<1 pM) without PCR amplification<sup>23, 28, 52</sup>. We

expect similar approaches could be adopted to profile exosomal nucleic acids. The resulting iMEX could be a powerful clinical tool for affordable, scalable, and comprehensive exosome analyses, thereby deepening our insights into tumor biology and accelerating effective cancer management.

## METHODS AND EXPERIMENTAL

#### Fabrication of the iMEX system

The device consists of a micro-controller (Atmega328, Atmel Corporation), a digital-toanalog converter (DAC8552, Texas Instruments), an analog-to-digital converter (ADC161S626, Texas Instruments), a multiplexer (ADG708, Analog Devices), and eight potentiostats. Each potentiostat consists of two operational amplifiers (AD8606, Analog Devices): one amplifier maintains the potential difference between a working electrode and a reference electrode, and the other one works as a transimpedance amplifier to convert a current to a voltage signal. The current measuring range of the transimpedance amplifier was  $\pm 7.5 \ \mu$ A. The eight-channel electrodes are commercially available (DropSens, Spain).

#### Preparation of immunomagnetic beads

5 mg of magnetic beads coated with epoxy groups (Dynabeads M-270 Epoxy, Invitrogen) were suspended in 1 mL of 0.1M sodium phosphate solution at room temperature for 10 minutes. The magnetic beads were separated from the solution with a permanent magnet and re-suspended in 100  $\mu$ L of the same solution. 100  $\mu$ g of antibodies against CD63 (Ancell) or respective IgG (Ancell) were added and mixed thoroughly. 100  $\mu$ L of 3M ammonium sulfate solution was added, and the whole mixture was incubated overnight at 4°C with slow tilt rotation. The beads were washed twice with phosphate buffer saline (PBS) solution and finally re-suspended in 2 mL of PBS with 1% bovine serum albumin (BSA). More details can be found in the manual provided by the manufacturer of the magnetic beads.

#### **Biotinylation of labeling antibodies**

10 mM Sulfo-NHS-Biotin (Pierce) solution in PBS was incubated with antibodies for two hours at room temperature. Unreacted Sulfo-NHS-Biotin was removed using Zeba spin desalting column, 7K MWCO (Thermo Scientific). Antibodies were kept at 4 °C until use.

#### iMEX assay

 $10 \,\mu\text{L}$  of exosomes-spiked PBS solution (or plasma) was mixed with  $50 \,\mu\text{L}$  of the immunomagnetic bead solution for 15 minutes at room temperature. The bead concentration was determined according to the following criterion:

 $\left[C_b \times V_b \times 4\pi R_b^2\right] / \left[C_e \times V_e \times \pi R_e^2\right] > 100$ , where  $C_b$  and  $C_e$  are the bead and the exosome concentrations, respectively;  $V_b$  and  $V_e$  are the volume of the bead solution and the exosome-spiked solution (or plasma), respectively;  $R_b$  and  $R_e$  are the mean radius of beads and exosomes, respectively. This requirement ensured that sufficient bead surface was available for exosome capture. In our experiment condition,  $R_e \sim 50$  nm,  $R_b = 1.4 \,\mu$ m, and  $C_e \sim 10^{10}$  /mL. Therefore, we adjusted the bead concentration to  $\sim 10^8$  /mL. The magnetic beads were separated from the solution with a permanent magnet and re-suspended in 80  $\mu$ L

of PBS (1% BSA). After 5 seconds of vortexing, the beads were separated and re-suspended in 80  $\mu$ L of PBS (1% BSA). 10  $\mu$ L of antibodies of interest (20  $\mu$ g/mL in PBS) were mixed with the beads for 15 minutes at room temperature. The magnetic beads were separated and washed as described before, and they were re-suspended in 50  $\mu$ L of PBS (1% BSA). 5  $\mu$ L of streptavidin-conjugated HRP enzymes (1:100 diluted in PBS) were mixed with the beads for 15 minutes at room temperature. The magnetic beads were separated and washed as described before, and they were re-suspended in 7  $\mu$ L of PBS. The prepared bead solution and 20  $\mu$ L of UltraTMB solution (ThermoFisher Scientific) were loaded on top of the screen-printed electrode. After 3 minutes, chronoamperometry measurement was started with the electrochemical sensor. The current levels in the range of 40-45 seconds were averaged.

### Enzyme-linked immunosorbent assay (ELISA)

CD63 antibody (Ancell) and IgG1 antibody (Ancell) were dilute to 5  $\mu$ g/mL concentration in PBS and added to the Maxisorp 96 well plate (Nunc) for overnight incubation at 4°C. After washing with PBS, 2% BSA in PBS blocking solution was added to the plate for 1 hour incubation at room temperature. Subsequently, 10<sup>8</sup> exosomes in 100  $\mu$ L PBS were added to each well for 1 hour incubation at room temperature. After discarding the blocking solution, antibodies (1  $\mu$ g/mL) against various markers were added to each well and incubated at room temperature for 1 hour. Unbounded antibodies were washed with PBS three times. Streptavidin-HRP molecules were added to the each well for 1 hour at room temperature. After washout with PBS, the chemiluminescence signal was measured.

#### Flow cytometry

 $5 \times 10^5$  cells per antibody were used for flow cytometry experiments. Cells were fixed with 4% paraformaldehyde for 10 minutes at room temperature, and then washed with PBS (0.5% BSA). Subsequently, cells were blocked with BSA (0.5% in PBS) and then incubated with primary antibodies (4 µg/mL). After primary antibody incubation, cells were washed, incubated with fluorophore conjugated secondary antibody (2 µg/mL; Abcam) and washed. The fluorescence signals from the labeled cells were measured using BD LSRII Flow Cytometer (BD Biosciences). Mean fluorescent intensities (MFIs) recorded were normalized using the following formula [(signal-IgG isotype control)/secondary]. Blocking and incubation with antibodies (primary and secondary) were preformed for 30 min each at room temperature. Every washing step comprised of three 5-min washes at 300 g with PBS (0.5% BSA).

#### **Cell culture**

OV90, OVCAR3, OCVA420, and TIOSE6 cells were grown in RPMI-1640 medium (Cellgro). CaOV3 were cultured in Dulbecco's modified essential medium (DMEM, Cellgro). All media were supplemented with 10% FBS and penicillin-streptomycin (Cellgro). All cell lines were tested and were free of mycoplasma contamination (MycoAlert Mycoplasma Detection Kit, Lonza, LT07-418).

#### Exosome isolation from cultured cells

We used a conventional method to harvest exosomes from cell culture media. Cells at passages 1–15 were cultured in vesicle-depleted medium (with 5% depleted FBS) for 48 hours. Conditioned medium from ~10<sup>7</sup> cells was collected, centrifuged at 300g for five minutes. Supernatant was filtered through a 0.2- $\mu$ m membrane filter (Millipore) and concentrated by 100,000g for one hour. After the supernatant was removed, exosome pellet was washed with PBS and centrifuged at 100,000g for 1 hour. Exosome pellet was resuspended in PBS.

#### Clinical samples preparation

The study was approved by the Institutional Review Board at the Dana-Farber/Harvard Cancer Center (PI: Castro), and the procedures followed were in accordance with institutional guidelines. Informed consent was obtained from all subjects (n = 11). Peripheral blood was withdrawn (~15 mL) from patients with ovarian cancer and centrifuged at 400 g for 15 minutes to separate plasma from red blood cells and buffy coat. 10  $\mu$ L of plasma was used for each surface marker analysis.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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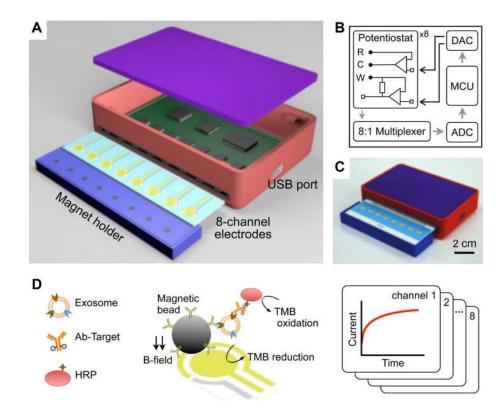
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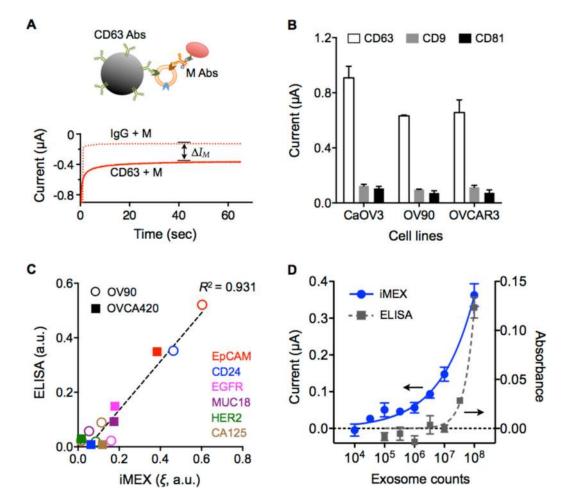
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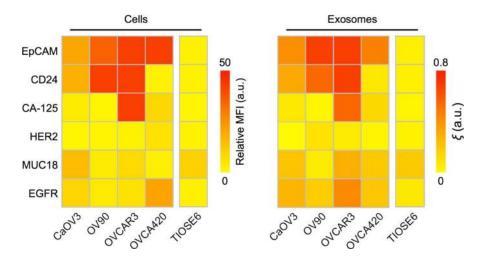
#### Figure 1. Integrated magnetic-electrochemical exosome (iMEX) platform

(A) Sensor schematic. The sensor can simultaneously measure signals from eight electrodes. Small cylindrical magnets are located below the electrodes to concentrate immunomagnetically captured exosomes. (B) Circuit diagram. The sensor system has eight potentiostats, an 8-to-1 multiplexer, an analog-to-digital converter (ADC), a digital-toanalog converter (DAC), and a micro-controller unit (MCU). Each potentiostat has three electrodes: reference (R), counter (C), and working (W). (C) A packaged device. The device has a small form factor ( $9 \times 6 \times 2$  cm<sup>3</sup>). (D) Schematic of iMEX assay. Exosomes are captured on magnetic beads directly in plasma and labeled with HRP enzyme for electrochemical detection. The magnetic beads were coated with antibodies against CD63, an enriched surface marker in exosomes. The working (W) and the counter (C) electrodes were made of gold (Au), and the reference electrode (R) was made of silver/silver chloride (Ag/AgCl). Eight channels are simultaneously monitored for high-throughput analysis. HRP, horseradish peroxidase; TMB, 3,3',5,5'-tetramethylbenzidine.

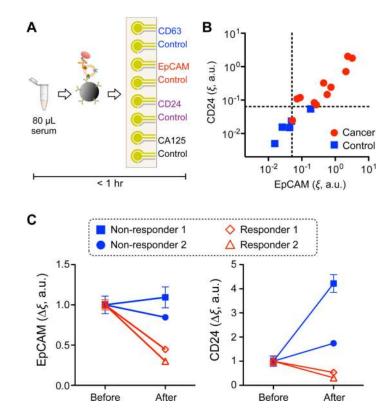


#### Figure 2. Characteristics of the iMEX diagnostic platform

(A) Schematic of the electrochemical measurement in the iMEX assay. With –100 mV reduction potential, the current level was reached a plateau within 1 minute. The current difference between the CD63-bead and IgG-bead samples ( $\Delta I_M$ ) was used as a representative value of a target protein marker. Abs, antibodies. M, marker. (B) Signal comparison of three tetraspanin markers (CD63, CD9, and CD81) in cancer exosomes. Signals from CD63 were much higher than those from other markers in exosomes collected from ovarian cancer cell lines (CaOV3, OV90, and OVCAR3). (C) Comparison between iMEX and ELISA. Six surface proteins were profiled in two ovarian cancer cell lines (OV90 and OVCA420). The results showed high correlation ( $R^2 = 0.931$ ). a.u., arbitrary unit. (D) Varying number of exosomes were spiked into human plasma and assayed by iMEX and ELISA. The detection limits were  $3 \times 10^4$  (iMEX) and  $3 \times 10^7$  (ELISA). All measurements were performed in triplicate, and the data are displayed as mean ± SD.



**Figure 3. Profiling of surface proteins in ovarian cancer cells and their secreting exosomes** Four ovarian cancer cell lines (CaOV3, OV90, OVCAR3, and OVCA420) and one normal cell line (TIOSE6) were screened for six putative cancer markers (via flow cytometry, left panel). Cell-derived exosomes were immunomagnetically captured (CD63-specific) and assayed by iMEX (right panel). The profiling data showed a good match between cells and CD63-positive exosomes. The iMEX assay was in duplicate, and the mean values are displayed.



#### Figure 4. iMEX for clinical applications

(A) iMEX assay for clinical sample analysis. 10  $\mu$ L of plasma per marker was used, and the entire assay was completed within 1 hour without filtration and centrifugation processes. (B) Plasma samples from ovarian cancer patients (n = 11) and healthy controls (n = 5) were analyzed with the iMEX assay. EpCAM and CD24 levels were much higher in cancer patients. The EpCAM and CD24 expression levels ( $\xi_{EpCAM}$  vs.  $\xi_{CD24}$ ) were highly correlated ( $R^2 = 0.870$ ). (C) Longitudinal monitoring of drug treatment responses. Plasma samples from four ovarian cancer patients were analyzed with the iMEX assay. EpCAM and CD24 levels in responders were decreased significantly, but their levels in non-responders were stable (EpCAM) or increased (CD24) after treatment. All measurements were in duplicate. a.u., arbitrary unit.