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2 *Methods Paper*

3 **Droplet Microfluidics with Reagent Micromixing for Investigating 4 Intrinsic Platelet Functionality**

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15 **Abstract**

16 *Introduction*—Precision mapping of the functional structure
17 of platelet populations holds great promise for the identification
18 of hyper-reactive subtypes that are likely to be disease
19 drivers, having value in prognostics and as therapeutic
20 targets. However, the ability to measure the intrinsic functional
21 capacity of individual platelets is confounded by
22 potent paracrine cross-talk, resulting in phenotypic remodeling
23 of the entire platelet population, and in doing so
24 obscuring the identity of hyper-reactive platelets.

25 *Methods*—To address this we have developed a droplet
26 microfluidics strategy for single platelet confinement to
27 exclude paracrine signaling. Consideration of the Poisson
28 distribution was used for high throughput single platelet
29 encapsulation and the preparation of minimal platelet
30 collectives serving as digital models for understanding the
31 role of hyper-reactive platelets coordinating system-level
32 behavior by paracrine signaling. Platelets are retrieved from
33 the droplets for phenotyping using standard flow cytometry.
34 In addition, we have incorporated a staggered herringbone
35 micromixing element for accurate agonist and antibody
36 dispensing in droplets.

37 *Results*—The methodology was used for characterizing
38 sensitivity distributions from healthy blood donors in
39 response to convulxin (agonist of the GPVI receptor, the
40 major platelet receptor for collagen). P-selectin exposure and
41 $\alpha_{IIb}\beta_3$ integrin activation were used as analytical end-points

42 to demonstrate the existence of hyper-reactive platelets that
43 direct 20-fold gains in system level sensitivity.

44 *Conclusions*—The analytical workflow represents an enabling
45 tool for the accurate classification of platelet subtypes and
46 description of their underlying biology.

47 **Keywords**—Platelets, Single cell analysis, Droplet microfluidics, Micromixing, Flow cytometry.

50 **ABBREVIATIONS**

PDMS	Poly(dimethylsiloxane)	51
PRP	Platelet rich plasma	52
HEPES	Sodium 2-(4-(2-hydroxyethyl)piperazin-1-yl)ethanesulfonate	53
BSA	Bovine serum albumin	54
CVX	Convulxin	55
PBS	Phosphate buffered saline	56

69 **INTRODUCTION**

70 Platelets respond to blood vessel damage for coordinated repair, but also play a critical role in cardiovascular disease. Platelet functional diversity has long been a matter of investigation^{5,21} with the enticing possibility of identifying a platelet subtype that drives pathophysiological thrombus formation, a prognostic marker for risk of heart attacks and strokes and a choice therapeutic target.^{2,6,13,21} Key examples are immature, so-called reticulated platelets, distinguished by volume and elevated RNA content correlating with

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disease^{3,10} and the identification of pro-aggregatory and pro-coagulant heterotypic states emerging following dual stimulation with collagen and thrombin.^{1,17} However, definitive platelet function measurements requires stimulation, leading to potent paracrine signaling driving the activation of otherwise unreactive platelets and obscuring the classification of hyper-reactive platelets from less reactive, 'follower' platelets. Methods are therefore needed to compartmentalize single platelets to prevent paracrine cross-talk and preserve their intrinsic functional capacity. Microfluidic methods are suited for single cell manipulations,¹⁴ with droplet microfluidics²³ being the preferred approach for high throughput encapsulations to enable large-scale screening campaigns.⁷ In this contribution, we build on the first demonstration of droplet microfluidics for investigating single platelet functional variability⁹ by providing a detailed description of the droplet confinement approach and the inclusion of a micromixer element for assay reliability and automation.

DROPLET MICROFLUIDICS

Droplet microfluidic devices were prepared using the standard SU-8 to PDMS soft lithography route. Inlet and outlet ports were introduced using a 1-mm-diameter biopsy punch (Miltex) and Scotch® Magic™ Tape (3M™, 810) was used to remove any particulates. Oxygen plasma bonding (Femto, Deiner) was used to secure the PDMS (Sylgard 184) channels to a microscope slide acting both as a window and handle. The microscope slide can be coated with a thin film of PDMS or used in its native state for plasma bonding. Following assembly, devices were flooded from the outlet with 1% (v/v) trichloro(1H,1H,2H,2H-perfluoroctyl)silane (Merck) in HFE-7500™ (3M™) to passivate the PDMS and glass surfaces. This ensures partitioning of the fluoro-oil phase to the microchannel walls for the reliable generation of monodisperse droplets. In addition, glass surfaces activate platelets,^{15,22} but the passivation of the *glassy-like* plasma activated PDMS and glass surfaces with the fluorinated silane prevents direct platelet contact and activation. For droplet generation experiments, 0.75% (v/v) 008-fluorosurfactant (RAN Biotechnologies) was added to a HFE-7500™ carrier fluoro-oil. The surfactant is a non-ionic tri-block copolymer of poly(ethylene glycol) (PEG) and perfluoropolythene (PFPE) which stabilizes aqueous-fluoro-oil emulsions and is biocompatible, preventing the adsorption and interaction of biomolecules at the droplet interface.⁸

Microchannels were fabricated to a height of 20 μm . Passive filters having 15- μm -gaps between structures were placed downstream of the fluoro-oil, ago-

nist and antibody inlets to reduce particle and fibre fouling, prolonging reliable device operation. The pinched droplet generation junction had a width of 22 μm , abruptly expanding to 50 μm for the generation of picolitre droplets. Droplet size can be modulated by shear rate (velocity) and also the fluoro-oil:aqueous flow ratio. In practice, a 4:1 ratio was used involving the delivery of 20 $\mu\text{L}/\text{min}$ fluoro-oil-surfactant with 2 $\mu\text{L}/\text{min}$ of both agonist and antibody solutions and 1 $\mu\text{L}/\text{min}$ platelet suspensions. Having independent platelet and antibody inputs prevents extravagant antibody consumption and is necessary to attain optimal flow cytometry signal to noise levels (see Electronic Supplementary Material, ESM Fig. 1). Samples and reagents were housed in syringes bearing 25G needles and interfaced to the microfluidic device using medical grade, sterile polythene tubing (ID 0.38 mm; OD 1.09 mm). This tubing is tightly secured over the needle and, by straightforward insertion, fits securely within the microfluidic ports (prepared using the 1-mm-diameter biopsy punch). Flow rates were managed using syringe pumps (Fusion 200, Chemyx). Droplet generation was imaged using a high-speed camera (2500 fps, Miro eX2, Phantom) to allow real-time quality control analysis, necessary to ensure flows had stabilised for the reliable generation of monodisperse droplets before collection for downstream analysis. The flow arrangements for monodisperse droplet generation are shown in Figs. 1a and 1b allowing the generation of monodisperse 25- μm -diameter droplets ($\text{CV} \pm 0.95\%$) with a volume of $\sim 8 \text{ pL}$ at a frequency of 10.4 kHz (Figs. 1c and 1d). Platelets are characteristically shear sensitive.¹¹ These conditions produce a junction mean flow velocity of $\sim 1 \text{ m/s}$, imparting shear rates and stresses which exceed physiological conditions.²⁰ Critically, however, junction transit times are of the order $\sim 50 \text{ } \mu\text{s}$, with droplets exiting the device within 20 ms. These shear durations are insufficient for platelet activation as evidenced using vehicle controls.

ENCAPSULATION STATISTICS

Platelets are near neutrally buoyant, remaining in suspension throughout experimental durations. This allows the continuous delivery of platelets to the microfluidic device for encapsulation within droplets. Unlike other cells which appreciably sediment this avoids the requirement for stirring mechanisms¹² which pose the risk of inadvertent platelet activation. Platelets are delivered randomly, requiring consideration of the Poisson statistic for controlled platelet encapsulation. The probability of platelets being encapsulated in a droplet is described as follows;



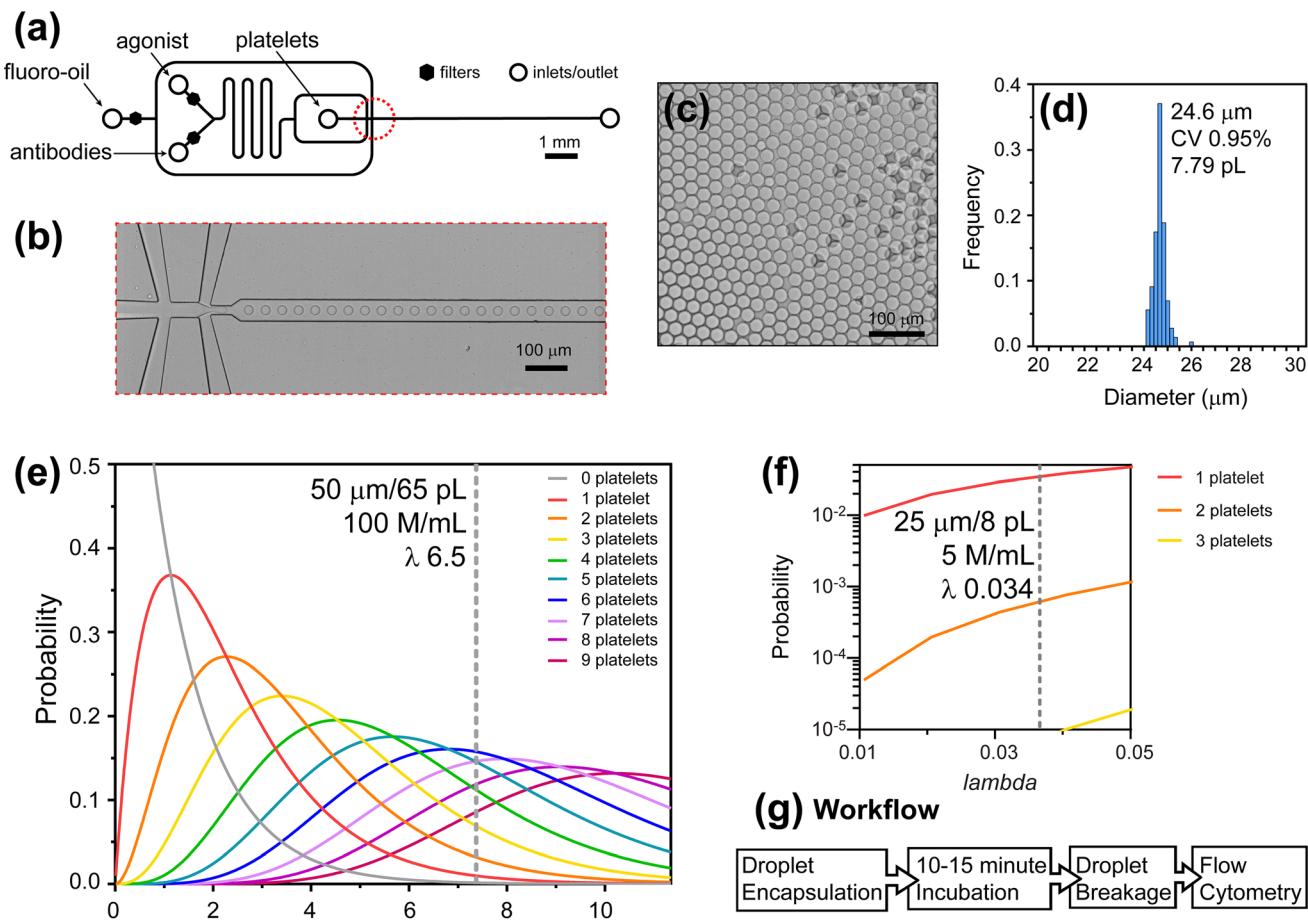


FIGURE 1. Microfluidics and Poisson distribution considerations for platelet encapsulation in droplets. Microfluidic device layout with aqueous and oil inputs and droplet generation junction identified with a red dashed circle (a). Droplets generated at 10.4 kHz (b) and resulting monodisperse droplets (c) with ImageJ-based size analysis involving >1000 droplets (d). Poisson distribution probabilities for given λ values to define encapsulation conditions for platelets with chosen $\lambda = 6.5$ conditions for multi-platelet encapsulation highlighted with a grey dashed line (e). Single platelet encapsulation probabilities (note \log_{10} y-axis scaling) with chosen $\lambda = 0.034$ conditions for single platelet encapsulation highlighted with a grey dashed line (f). Overall workflow for measuring single platelet intrinsic functional capacity (g). Figure elements adapted from Jongen *et al*⁹ (CC BY 4.0)

$$P_{(X)} = \frac{\lambda^X e^{-\lambda}}{X!}$$

where $P_{(X)}$ is the probability of X number of platelets being packaged in a droplet and λ the mean number of platelets per droplet. For the encapsulation of single platelets in the ~8 pL droplets we used a conservative platelet input concentration of $25 \times 10^6/\text{mL}$, that is diluted 5-fold upon combination with agonist and antibody streams. This yields a λ value of 0.034, or 3.34% of droplets containing a single platelet, 0.08% containing multiples and with the majority of droplets empty. In this case, single platelet encapsulations outnumbered multiple platelet encapsulations 42-fold, with flow cytometric forward scatter area vs height gating used to further exclude platelet multiples (aggregating upon activation) from analysis. Single platelet encapsulation eliminates paracrine cross-talk, but

this at the expense of throughput. A strength of droplet microfluidics is the kHz droplet generation rates, so even with >96% empty droplets, >100,000 single platelets can be encapsulated in 5 min, sufficient to effectively survey diversity in the platelet population.

In addition to using droplets to measure single platelet intrinsic functionality, the analysis of small platelet collectives has value for understanding the role of paracrine cooperativity: The presence of a single hypersensitive platelet in confinement is sufficient to activate other platelets in the same droplet, whereas the absence of hypersensitive platelets in a droplet prevents the platelets from being activated. In doing so, a digital responsive:unresponsive pattern is observed when analysing the droplets by fluorescent microscopy. For the preparation of small, minimal collectives the droplet microfluidics device was up-scaled with a $50 \times 50 \mu\text{m}$ droplet generation junction and operation using 80

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209 μ L/min fluoro-oil, 4 μ L/min undiluted PRP ($\sim 5 \times 10^8$ mL), and 8 μ L/min agonist and antibody flows. This
 210 produced 50- μ m-diameter droplets ($CV \pm 0.89\%$), with
 211 the 65 pL volume (and dilution by reagents) leading to
 212 a μ value of 6.5, or 0–15 platelets per droplet. Poisson
 213 distribution theory and the described single and mul-
 214 tiple platelet encapsulation conditions are described in
 215 Figs. 1e and 1f.

217 PLATELET SENSITIVITY MEASUREMENTS

218 Controlled platelet encapsulation requires a platelet
 219 count during sample preparation. This was achieved
 220 using a method involving platelet-specific CD61 anti-
 221 body labelling for analysis using an Accuri C6 (BD
 222 Biosciences) instrument.¹⁶ The overall sampling pro-
 223 cess involved obtaining blood from healthy volunteers
 224 (free from anti-platelet medication) with informed
 225 consent and ethical approvals. Wide-bore venepunc-
 226 ture using a 21G needle was used to collect blood (with
 227 the first 4 mL discarded) into vacuum tubes containing
 228 1:10 v/v 3.2% trisodium citrate. Gentle tube inversion
 229 was followed by centrifugation at 240 g for 15 min
 230 without brake for the preparation of PRP that was
 231 rested for 30 min before experiments. Using the pla-
 232 telet count, the PRP was adjusted to a final concen-
 233 tration of 25×10^6 /mL in HEPES buffer (136 mM
 234 NaCl, 2.7 mM KCl, 10 mM HEPES, 2 mM MgCl₂,
 235 0.1% (w/v) glucose and 1% (w/v) BSA (pH 7.45) for
 236 single platelet measurements or left undiluted for the
 237 minimal collective measurements.

238 The overall analytical workflow involved the
 239 encapsulation of single platelets with agonist and flu-
 240 orescent reporter antibodies in droplets, followed by
 241 incubation, then releasing and fixing the platelets for
 242 flow cytometry (Fig. 1g). Platelets were stimulated
 243 with the potent platelet agonist convulxin (CVX,
 244 (Enzo Life Sciences) a snake venom toxin serving as a
 245 soluble proxy for ECM proteins including collagen
 246 that activate platelets via the GPVI receptor^{4,18} to
 247 drive global platelet activation with notable end-points
 248 being $\alpha_{IIb}\beta_3$ integrin activation²⁴ (inside-out signalling)
 249 and P-selectin exposure, a marker for alpha granule
 250 secretion. These activation markers were detected
 251 using fluorescent antibodies from BD Biosciences;
 252 fluorescein isothiocyanate (FITC)-conjugated PAC-1
 253 mAb (PAC-1 clone; 1.25 ng/ μ L) that recognizes the
 254 active conformation of $\alpha_{IIb}\beta_3$, and allophycocyanin
 255 (APC)-conjugated anti-CD62P mAb (AK-4 clone; 0.63
 256 ng/ μ L) for P-selectin. In addition, R-phycocerythrin
 257 (PE)-conjugated anti-CD42b mAb (HIP1 clone; 1.25
 258 ng/ μ L) was used as a platelet identifier for gating. A
 259 10–15 min incubation (room temperature, dark) pro-
 260 duced an optimal signal to noise, with this temporal

261 window resulting from continuous droplet formation
 262 for a 5 min period required to collect $\sim 100,000$ droplet-
 263 confined platelets in a tube followed by incubation for
 264 10 min at room temperature. Technical developments
 265 are envisaged for continuous encapsulation and con-
 266 tinuous platelet release following an in-tubing dynamic
 267 incubation to improve the platelet-platelet incubation
 268 time reproducibility and allow assay automation.
 269 Further assay refinements consider the use of a mineral
 270 oil overlay to eliminate, albeit rare, droplet coalescence
 271 at the emulsion:air interface. For the larger, 50 μ m
 272 droplets a 60 min incubation was used to allow
 273 aggregation to conclude at room temperature in the
 274 absence of stirring (see ESM Fig. 2). This allows bio-
 275 marker presentation with an appreciable signal to
 276 noise.

277 Platelets were released from the droplets and fixed
 278 by breaking the emulsion. This involved removal of the
 279 fluoro-oil layer beneath the emulsion layer, the addi-
 280 tion of 200 μ L CellFixTM fixative (BD Biosciences)
 281 then 200 μ L 1H,1H,2H,2H-perfluoro-1-octanol (PFO,
 282 Merck), a weak but neat surfactant that exchanges
 283 with the PEG-PFPE surfactant to coalesce the droplets
 284 with fixative to promptly preserve platelet activation
 285 states. Careful pipetting is then needed to retrieve the
 286 platelet volume while avoiding contamination with
 287 PFO. As a consequence many platelets are discarded,
 288 and this is further exacerbated when using a mineral oil
 289 overlay. Nevertheless, the extreme throughput of dro-
 290 let microfluidics allows 10's of thousands of platelets
 291 (from the original 100,000) to be retrieved for analysis.
 292 Finally samples were measured using an Accuri C6
 293 cytometer (BD Biosciences). In the case of the minimal
 294 collectives, samples were first diluted with PBS to
 295 substantially reduce co-incident events during flow
 296 cytometry.

297 The droplet microfluidics approach for capturing
 298 intrinsic single platelet sensitivities was test-driven
 299 using a CVX dose response experiment. Key data are
 300 presented in Figs. 2a and 2b, with vehicle controls
 301 compared to maximal stimulations with 100 ng/mL
 302 CVX and an intermediate 1 ng/mL CVX stimulation
 303 producing a distinct bimodal activation distribution
 304 for both $\alpha_{IIb}\beta_3$ active conformation and P-selectin
 305 expression level end-points. Results are compared with
 306 platelet collectives able to communicate in bulk solu-
 307 tion, but otherwise treated to the same methodology as
 308 droplet-confined single platelets. The platelet collect-
 309 ives demonstrate substantial (~20-fold) median sensi-
 310 tivity gains. This behaviour can be attributed to
 311 hypersensitive platelets (identified by single platelet
 312 experiments) activating and coordinating the activa-
 313 tion of neighbouring platelets by paracrine signalling.
 314 The overall collective dose response behaviour⁹ indi-
 315 cates a broad-scale sensitivity continuum (not hetero-



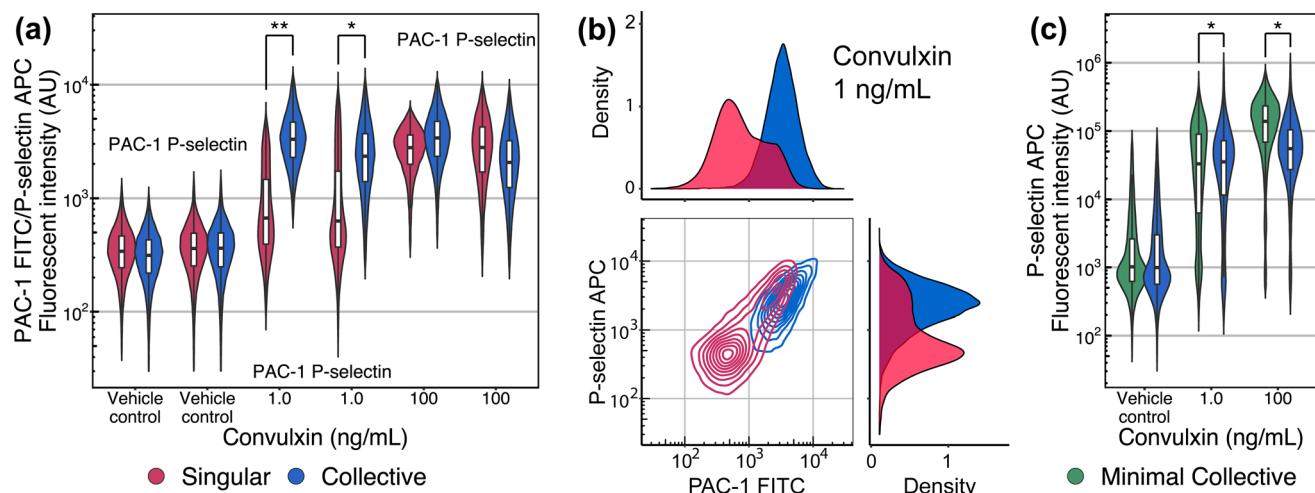


FIGURE 2. Single and collective platelet responses to CVX stimulation. Signals from vehicle controls, 1 ng/mL and 100 ng/mL CVX stimulations of single droplet-confined platelets and bulk, platelet collectives, with PAC-1 and P-selectin end-points (a). Contour and density plots of the fully activated collective condition and the bimodal activation distribution of single platelets stimulated with 1 ng/mL CVX (b). Minimal collective (0–15 platelets) and bulk, collective platelet vehicle controls and responses to 1 ng/mL and 100 ng/mL CVX stimulations using the P-selectin end-point (c). The relative risk was used to ascertain significance (* >2 , ** >10). For single platelet conditions, $n = 10,000$ –36,000 platelet events were measured, and $n \approx 40,000$ –55,000 for the minimal collective and collective conditions. Violins were plotted using the 1–99th percentiles. Figure adapted from Jongen *et al.*⁹ (CC BY 4.0)

typic sensitivity states). With low CVX doses (0.1 and 0.3 ng/mL), rare, hyper-reactive platelets drive the activation of some neighbours and with increasing doses progressively more platelets are directly activated by CVX and activate other neighbours until sufficient numbers of platelets are activated to drive the entire population into a fully activated state. Other experimental scenarios and outcomes may indicate the existence of heterotypic populations. However, platelet variability will likely obscure the boundaries between platelet subtypes. In this event, peak fitting software from the field of spectroscopy is well suited for identifying different platelet subtypes within the density plots.

The collective sensitivity gains are attributed to paracrine cross-talk between platelets. Minimal platelet collectives with 0–15 platelets encapsulated within 50- μ m-diameter droplets (~65 pL) evidence this inference. The minimal collectives produced platelet population responses similar to platelet collectives (size-based gating was removed to capture platelet aggregates). However, complete activation produced P-selectin expression signals exceeding those of the platelet collectives. This reveals the consequences of confinement, in which secretion products accumulate, driving platelets to higher activation states. With intermediate 1 ng/mL CVX stimulations the P-selectin expression signals span 4 orders of magnitude (see Fig. 2c). This is a result of variable platelet numbers (0–15 platelets), dictated by the Poisson distribution during encapsulation, but also the generation of two population

states: by extending analysis to fluorescent imaging using P-selectin and CD42b biomarkers allows the identification of droplets containing aggregates and others with multiple platelets but without aggregates, indicating the presence of hypersensitive platelets in the former but not the latter. These considerations can be tested using experiments involving paracrine inhibitors. Taken, together the data emerging from droplet-confined single platelet and minimal collective measurements holds promise for the development of mathematical models for predicting community cooperation and resultant platelet collective behaviours originating from imbalance in the platelet population structure.

MICROMIXING

Investigating single cell phenotypic diversity demands assay precision and reliability to prevent the population structure from being artificially distorted. For single platelet sensitivity measurements, each platelet must be encapsulated with defined agonist and antibody quantities. While droplet microfluidics represents an appealing route to automation, occasional flow oscillations alter droplet contents (see Fig. 3a). In practice, this requires continuous supervision to discard these droplets from analysis.

As a step towards unattended automation we have integrated a micromixing element for the homogenization of agonist and antibody inputs upstream of

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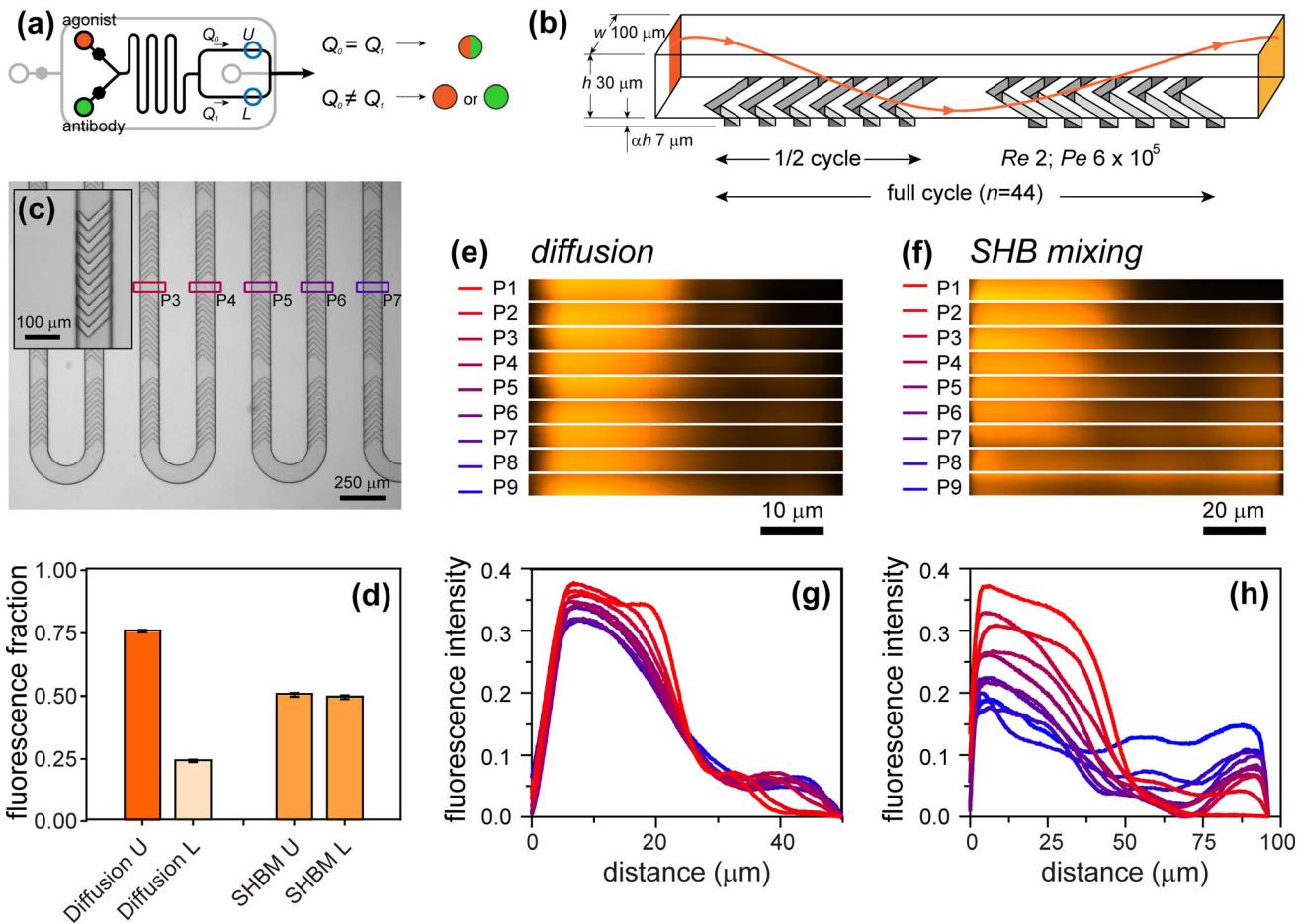


FIGURE 3. Staggered herringbone micromixing. Device layout with illustration of flow instability consequences for droplet loading (a). Staggered herringbone (SHB) micromixing concept with flow and mixing conditions (b). SHB microfabricated ridges and distribution within the serpentine channel, with P3–7 indicating fluorescent confocal microscopy measurement positions (c). Fluorescence distributions (mean \pm SD, $n = 3$) emerging from the upper (U) and lower (L) outlets (see (a), blue circles) at the end of the serpentine channel for diffusion-limited mixing and SHB micromixing (d). 3D fluorescent cross-sections and intensity profiles of diffusion-limited (e, g) and SHB micromixing (f, h) at the different P1–P9 measurement positions.

droplet generation to ensure the accurate dosing of each droplet. To this end, we have used topographic patterning in the form of a staggered herringbone (SHB) micromixer¹⁹ (see Fig. 3b). In a first prototype, devices were fabricated to a height of 30 μm , requiring 40 $\mu\text{L}/\text{min}$ fluoro-oil, 2 $\mu\text{L}/\text{min}$ platelets, and 4 $\mu\text{L}/\text{min}$ agonist and antibody flows for the reliable production of ~ 14 pL droplets at ~ 12 kHz. Herringbone structures in repeats of 6 were alternated in 44 cycles along the 45-mm-long serpentine channel (see Fig. 3c and ESM CAD) and fabricated to a depth of 7 μm within the base of the microchannel. The serpentine channel width was increased from 50 to 100 μm producing equivalent Reynolds number ($Re = 2$) and Péclet numbers for the original and herringbone micromixer systems. The mixing performance of lateral diffusion was compared with herringbone-driven micromixing using an experiment combining PBS with a second PBS stream containing 100 $\mu\text{g}/\text{mL}$ PLL-FITC (Merck, $M_{\text{wt}} = 50$ kDa;

$Pe = 6 \times 10^5$) which served as an antibody model with a similar diffusion coefficient ($D = 3 \times 10^8 \text{ cm}^2/\text{s}$). 394
Outputs from either side of the ring structure were 395
measured using a fluorescent plate reader (BMG 396
LabTech FLUOstar Optima). 397

The herringbone structures introduce asymmetric 398 drag, creating flow anisotropy which results in the 399 lateral circulation and folding of the two input laminar 400 streams.¹⁹ The structures are grouped creating a 401 half cycle to cumulatively drive circulation in one 402 direction, then mirrored for circulation in the 403 opposite direction to complete a cycle. The 404 staggered herringbone micromixer delivered complete 405 mixing, with equivalent 406 PLL-FITC levels delivered to the upper and lower 407 outlets of the ring structure at the end of the 408 serpentine channel. In contrast, diffusion-limited 409 PLL-FITC homogenization produced a 3-fold 410 difference in the 411 PLL-FITC levels (see Fig. 3d). Standard 412 fluorescent imaging was used to compare diffusion-limited

homogenization with SHB micromixing throughout the 45-mm-length of the serpentine channel (see ESM Fig. 3), corroborating the fluorescent fractions from the upper and lower outputs.

Diffusive transport is slow, whereas SHB micromixing produced layers in the flow. To image PLL-FITC dispersion throughout the serpentine channel cross-section fluorescent confocal microscopy was used (Leica SP8, 5x objective (14x zoom), 0.15 NA, 488 nm excitation, 500–600 nm emission, 0.5 μm z step, 0.28 pinhole). The time dependent PLL-FITC profile for lateral diffusion is compared with SHB micromixing in Figs. 3e–3h. As predicted with diffusion-limited mass transfer, the fluorescent gradient diminishes along the channel. There is also the emergence of a low fluorescent signal at the far side of the channel, perhaps indicating small magnitude secondary Dean flows ($\kappa \approx 0.3$; 150 μm radius of curvature) generated at the 11 hairpins along the serpentine. With the SHB micromixer the PLL-FITC stream is alternately folded, stretching and thinning the fluid layers with diffusion concluding mixing between the narrowing layers. Although mixing is achieved at the outlets, we have incorporated ~3-fold more SHB mixing cycles than used in the first demonstration by Stroock *et al.*¹⁹ We largely attribute this to the lower number of ridges in each of our half cycle patterns ($6 < 10$). Therefore, we predict this prototype can be improved by increasing the herringbone repeats for extensive lateral transport before flow reversal in the following half cycle. In doing so, the serpentine can be shortened to produce a smaller footprint device with lower back pressure and faster flow equilibration. Nevertheless, the current staggered herringbone mixing element ensures accurate reagent dispensing in droplets and can readily be applied to other microfluidic applications requiring on-board mixing.

CONCLUSIONS

In summary, the droplet microfluidics and cytometry workflow provides a high throughput approach for measuring intrinsic platelet functionality. The inclusion of a herringbone micromixer ensures the accurate delivery of agonist and antibody reagents to the droplets and represents a step towards assay automation for reliably profiling platelet functional diversity.

SUPPLEMENTARY INFORMATION

The online version of this article (<https://doi.org/10.1007/s12195-020-00665-6>) contains supplementary material, which is available to authorized users.

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AUTHOR CONTRIBUTIONS

M.S.A.J., S.I.R.L., P.M.H. and J.W. undertook the research. M.S.A.J., S.I.R.L. and J.W. prepared the figures and manuscript. P.M.H., N.A.E. and O.J.T.M. revised and edited the manuscript. All authors approved the final version of the manuscript.

CONFLICT OF INTEREST

M.S.A.J., S.I.R.L., P.M.H., N.A.E., O.J.T.M. and J.W. declare no conflicts of interest.

RESEARCH INVOLVING HUMAN AND ANIMAL RIGHTS

All human subject research was carried out in accordance with institutional guidelines approved by the University of Southampton Ethics Committee ((ERGO 5538) and South Central – Hampshire B National Research Ethics Service (REC: 14/SC/0211)). No animal studies were carried out by the authors for this article.

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