

High-Throughput Sorting and Placement of One-Bead–One-Compound (OBOC) Libraries from Bulk to Single Wells in Organic Solvent

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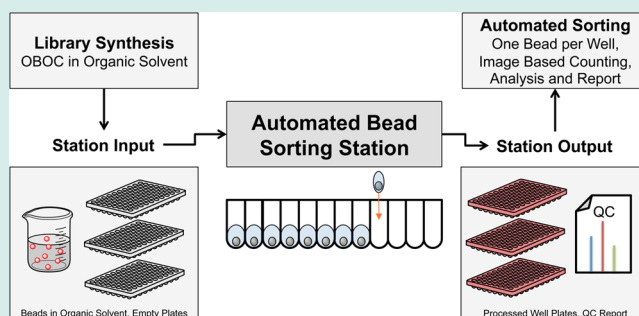
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S Supporting Information

ABSTRACT: One-bead–one-compound (OBOC) solid-phase combinatorial chemistry has been used extensively in drug discovery. However, a major bottleneck has been the sorting of individual beads, while still swollen in organic solvent, into individual wells of a microwell plate. To solve this problem, we have constructed an automated bead sorting system with integrated quality control that is capable of sorting and placing large numbers of beads in bulk to single wells of a 384-well plate, all in an organic solvent. The bead sorter employs a unique, reciprocating fluidic design capable of depositing 1 bead every 1.5 s, with an average accuracy of 97%. We quantified the performance of this instrument by sorting over 8500 beads, followed by cleaving the conjugated compound and confirming the chemical identity of each by liquid chromatography/mass spectrometry (LC/MS). This instrument should enable more efficient screening of combinatorial small molecule libraries without the need to dry beads or otherwise change the chemical environment.

KEYWORDS: bead sorting, OBOC, organic solvents, high-throughput screening, combinatorial libraries



Since the introduction of solid phase supports for the synthetic chemistry of a tetrapeptide by Merrifield in 1963,¹ inert beads have been used to generate hundreds of thousands of unique compounds. Solid supports are commonly used in the “one-bead–one-compound” (OBOC) approach, and they have been used to generate diversity-oriented libraries for nerve targeting dyes,² cancer-targeting peptides,³ and chemical genetics^{4,5} to name just a few.

Split-pool methods, in combination with equipment designed for high-throughput screenings (HTS), have been a popular approach to drug discovery over the past 20 years.^{6–8} However, this approach also comes with the unique challenge of segmenting and analyzing each individual solid-phase bead. Typically, a completed library of beads are segmented into individual wells of a 96-, 384-, or 1496-well plate followed by either cleavage and chemical screening, or direct on-bead screening.⁹ The post-synthesis segmentation of a completed library, although easy for the human hand, is time-consuming and extremely monotonous. Previously described methods for arraying micrometer- to millimeter-sized supports include using a vacuum manifold with dry beads,⁵ droplet deflection using

electrostatics,¹⁰ and fluidic deflection of water compatible beads.¹¹ Although each of these methods have their strengths, no system described to date automates the process of sorting beads into multiple microtiter plates while maintaining the bead in an organic solvent and providing deposit error checking.

These three issues are of practical importance: automation, solvent compatibility, and error checking. Bead segmentation processes that rely too heavily on human intervention have restricted the benefits of the diversity-oriented split pool method. The ability to automate sorting over multiple plates removes the limitation of the human hand and the need for human supervision. A bead sorter with chemically compatible fluidics provides flexibility in choosing common, functionalized polystyrene resins that are otherwise difficult to handle in aqueous environments.¹² Lastly, error checking of sorted plates permits verification of the process while simultaneously

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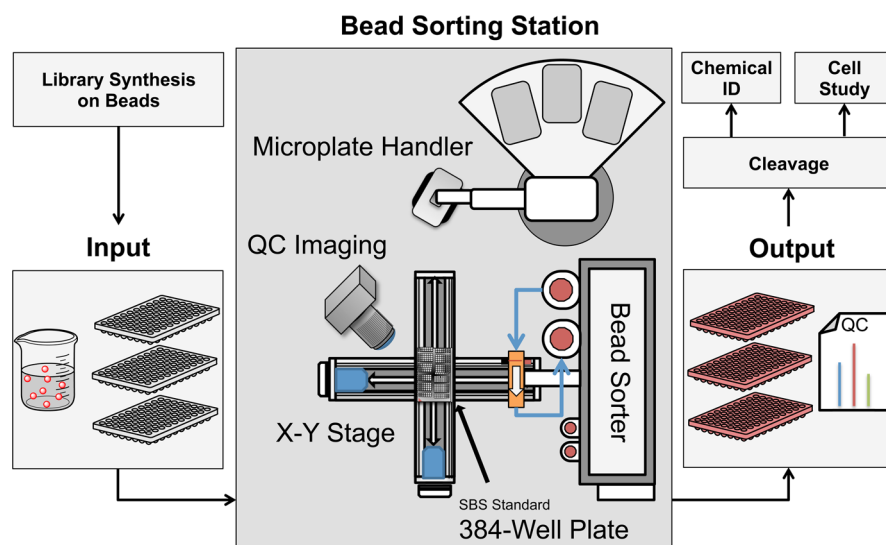


Figure 1. Design and functionality of the automated bead sorting system. Single-bead/single-well sorting was performed after solid phase chemical library synthesis on beads 400–500 μm in diameter. Post-cleavage analysis of processed plates may include chemical identification or cell study assays after compound cleavage from the beads. The bead sorting system is comprised of a microplate handler, a camera for quality control, an X–Y stage, and a bead sorter, arranged as shown.

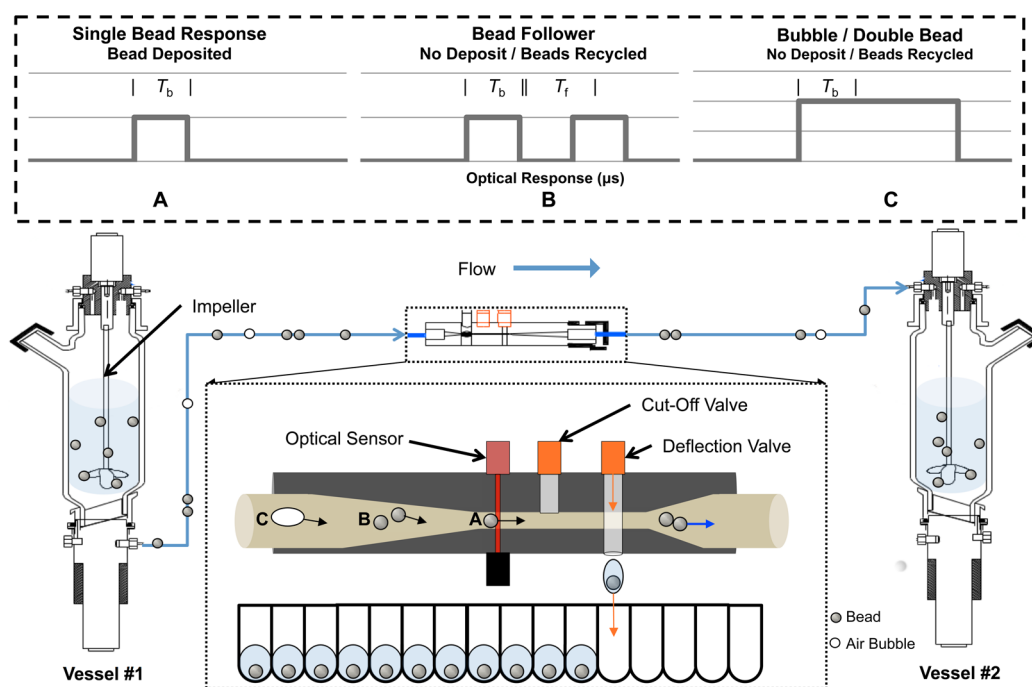


Figure 2. Bead fluidics and optical sensing. Beads pass through an optically monitored flow cell while being transported in an organic solvent from the source vessel (Vessel #1) to the receiver vessel (Vessel #2). The optical sensor data are used in conjunction with single bead deposit parameters (T_b , T_f , etc.) to deflect individual beads from the suspension into a microtiter plate well below. Shown are examples of (A) a single bead deposit response, (B) a pair of beads traveling too close for a successful deposit, and (C) an air bubble in the fluidic line. At the point when the source vessel no longer contains fluid, the vacuum/pressure levels reverse and the receiver vessel becomes the source vessel (and vice versa). This process is repeated until the predetermined number of plates is complete or no successful single bead deposits can be accomplished.

propagating individual well information that is required for post-sorting analysis and screening.

In this study, we hypothesized that an instrument that incorporated chemically resistant fluidics and quality control (QC) algorithms would be capable of sorting OBOC chemical libraries with high speed and high accuracy, without the need to remove the beads from their native chemical environment.

Hardware Integration. As shown in Figure 1 (and in Figures S1 and S2 in the Supporting Information), the bead

sorting station was assembled using three pieces of commercially available equipment and a custom bead sorter. The equipment can be categorized by function: management of SBS plates, management of bead sorting, and quality control. A microplate handler (Caliper Twister II) was used for plate management; its two separate reserve towers were used for the storage of new plates and processed plates. A TechElan Bead Sorter (TeBS) (TechElan, Mountainside, NJ, www.techelan.com), which is a custom piece of equipment designed for

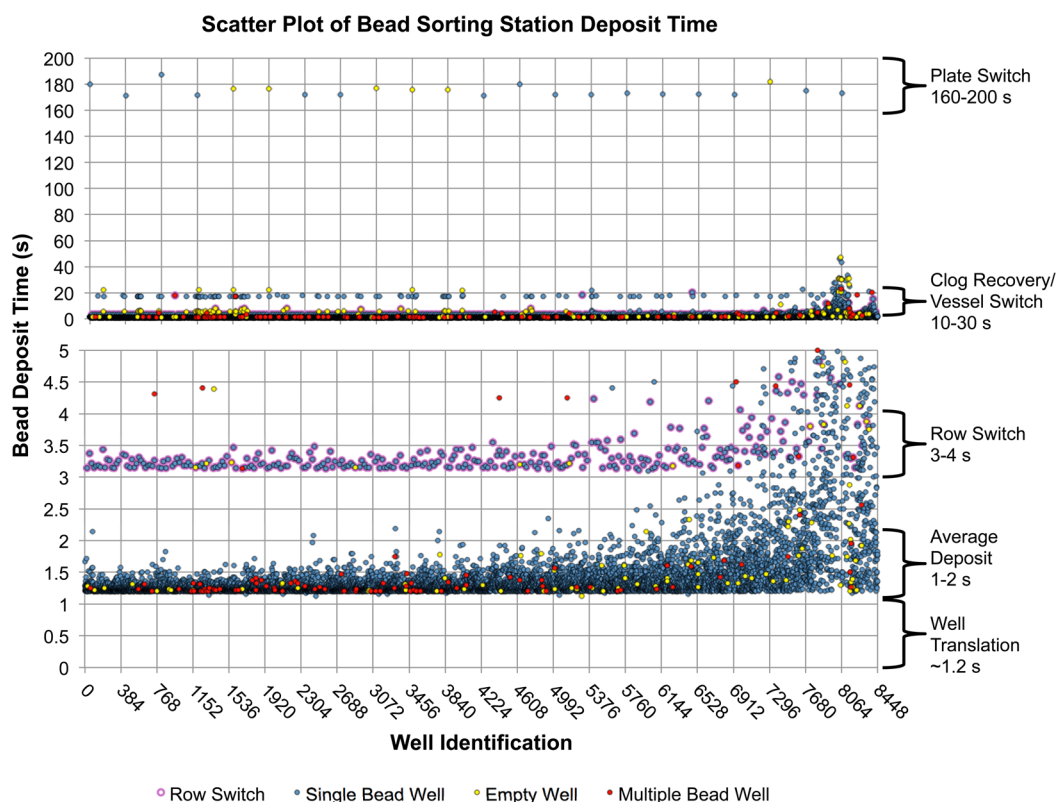


Figure 3. Sorting and deposit statistics. Shown are the bead sorting and deposit statistics for ~ 8500 $400\text{--}450$ μm CIT beads conjugated to fluorescein in DMSO deposited into 22 384-well plates. Bead deposit time is measured as the time between two “deposit successful” signals from the bead sorter and can include (1) dead time from the X–Y stage advancing to the next well, (2) monitoring time until a single bead condition is met, (3) dead time from the X–Y stage advancing to the next row, (4) vessel reversal time, (5) clog recovery time, and (6) plate switching time. Expressed through the color of each deposit spot is the image-processed counting result for that well, which may be empty, contain a single bead, or contain multiple beads. In addition, every 25th well is highlighted in purple outline to indicate the extra time needed to perform a plate row switch.

depositing single beads from a suspension of thousands, was selected for bead management. Bead counting was initially accomplished with individual well imaging via a dark-field camera assembled from Microvideo Instruments (Avon, MA). However, the use of a flatbed scanner (Canon, Model CanoScan 8800F) was chosen to provide whole-plate analysis, which benefits the system with reduced processing time. A translation stage was assembled using two lead screw-driven actuators (Velmex BiSlide) for transporting plates between the areas of plate management, bead management, and quality control. As detailed in Figure 1, the assembly of these four pieces of equipment was the first step in a multistep process, starting from library synthesis and ending with a completed screening assay.

Bead Sorter. The TeBS was selected for bead sorting because of its infinite-loop design. As shown in Figure 2, the beads flow from the source vessel to the receiver vessel while passing through an optically monitored flow cell. A total of 10 timing parameters were determined empirically for high accuracy of single bead deposits. These parameters include signal processing triggers that are related to (1) the size of the bead, (2) the separation between beads, and (3) the presence of double beads and bubbles. The most important of these parameters are (1) the bead width time (T_b), which is used to exclude double beads and air bubbles, and (2) the bead follower time (T_f), which negates the deposit if another bead is observed to be following too close to the bead of interest. The custom flow cell of the TeBS is suited for beads $400\text{--}800$ μm in size.

Software Integration. The equipment described above was assembled and fully controlled via a custom-made software application written in C# and based on proprietary DLL libraries from National Instruments (NI-DAQ, NI-IMAQdx, and NI-VISION) and Caliper Instruments (iLink Pro). The TeBS communication with the software is accomplished with three digital I/O signals, which reflect the “request bead,” “successful deposit,” and “wait for request” TTL signals to the PLC controller within the TeBS. The “request bead” signal is used by the software to initiate the process of a single bead deposit by the TeBS only if the “wait for request” signal is not active. This “wait for request” TTL is used as feedback to the software to indicate that the TeBS is currently unavailable to deposit a bead, which occurs during routine processes such as vessel switching or fluidic line cleaning. The “successful deposit” signal is the output of the TeBS to confirm a bead deposit, which then triggers the advancement of the well plate by the Velmex stage. Since this station was designed to operate without human intervention, the most challenging aspect was maintaining alignment of the equipment. Positional calibration of the microplate handler was completed through the provided iLinkPro software and bead sorting positional alignment was completed using a three-well calibration within the station’s software.

Automated Bead Sorting in Organic Solvent. Using the customized bead sorter (Figure 2), ~ 8500 fluorescein-conjugated beads in DMSO were sorted automatically into 22 384-well plates (Figure 3). The average bead deposit time was 1.3–1.5 s per bead, which is due largely to the ~ 1.2 s dead

between “successful deposit” signals of the TeBS. The well deposit time takes into account the dead time of the X – Y translation from the previous well, the optical monitoring time, and other dead time such as automated clog detection/recovery, or vessel switching.

Image-Based Bead Quantification. Post-sorting imaging of the well plates is used for QC, which quantifies the number of beads in each well. Images were collected either on a per-well basis, using a dark-field camera (Microvideo Instruments), or a per-plate basis, using a Canon flatbed scanner equipped with a back light. The image-processing algorithm utilizes feature detection and a counting classification, which includes measuring any color attributes of the bead (Color Plane Extraction) and extracting an ROI for each well. The image is then processed using a binary threshold for segmentation, analyzed, and discriminated based on the “centroid,” “area,” and “circularity” features of each blob particle. Finally, classification of the blobs is accomplished through an empirically determined area and circularity relating to the single, double, or triple beads per well. Counting results are then merged with deposit timing results acquired from the TeBS. Example images and a counting flow-chart are provided in Figure S3 in the Supporting Information.

Chemical Identification. After sorting into individual wells, the chemical identity of each bead was determined by preparing a parent plate consisting of conjugated dye cleaved from the resin using an automated liquid handler (Caliper SciClone ALH2000). Child plates were generated by an automated process using the same liquid handler by diluting and transferring the parent plate to desired concentrations suitable for UPLC-MS (Waters Acquity UPLC–Xevo G2 QTOF), MALDI-TOF (Bruker MALDI-TOF-TOF), and plate scanning (Gemini XS, Molecular Devices, Sunnyvale, CA). In the case of the child plate prepared for UV–vis absorbance and fluorescence, sodium hydroxide was used to increase the pH to a suitable, basic environment.

The fluorophore was identified during LC-MS by fluorescence at 521 nm and ESI-MS at 331 m/z . Plate scanning spectra were collected at wavelengths of 480–680 nm with an excitation of 450 nm. MALDI-TOF identification was completed with observed negative ions at 287 m/z and 353 m/z , corresponding to $IM-COOH^-$ and $IM-2H+Na^-$. Additional analysis details are provided in Figure S5 in the Supporting Information.

Chemicals, Reagents, and Disposables. Standard SBS-sized 384-well plates with 100 μ L round-bottom polypropylene wells were used in this study (Corning, CoStar 3657). Chloro-(2'-chloro)trityl (CIT) polystyrene resin was purchased through Rapp Polymere (Tuebingen, Germany) in sizes of 400–450 μ m (Catalog No. H40045033) and 500–560 μ m (Catalog No. H50056033) before swelling in an organic solvent. Fluorescein was purchased from Sigma–Aldrich (Catalog No. 32615-25G-R).

■ ASSOCIATED CONTENT

● Supporting Information

Further experimental details, as well as supplementary figures for the bead sorting system. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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Author Contributions

M.W.B., C.J.C., K.B., J.H.L., I.F., A.B.C., and R.O. performed the experiments. M.W.B., R.O., I.F., J.V.F., and H.S.C. reviewed, analyzed, and interpreted the data. M.W.B., H.S.C., and J.V.F. wrote the paper. All authors discussed the results and commented on the manuscript.

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Notes

The authors declare the following competing financial interest(s): TeBS Bead Sorter is commercially available from TechElan, a for-profit company. Ilya Feygin is the Vice President of Engineering at TechElan.

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■ REFERENCES

- (1) Merrifield, R. B. Solid phase peptide synthesis. I. The synthesis of a tetrapeptide. *J. Am. Chem. Soc.* **1963**, *85* (14), 2149–2154.
- (2) Bao, K.; Yoon, J. S.; Bordo, M. W.; Cross, C. J.; Lee, J. H.; Oketokoun, R.; Jeong, M. Y.; Choi, H. S. An automated robotic chemistry system for developing nerve targeting agents. Manuscript in review, 2015.
- (3) Aina, O. H.; Liu, R.; Sutcliffe, J. L.; Marik, J.; Pan, C. X.; Lam, K. S. From combinatorial chemistry to cancer-targeting peptides. *Mol. Pharmaceutics* **2007**, *4* (5), 631–651.
- (4) Blackwell, H. E.; Perez, L.; Stavenger, R. A.; Tallarico, J. A.; Cope Eatough, E.; Foley, M. A.; Schreiber, S. L. A one-bead, one-stock solution approach to chemical genetics: Part 1. *Chem. Biol.* **2001**, *8* (12), 1167–1182.
- (5) Clemons, P. A.; Koehler, A. N.; Wagner, B. K.; Springings, T. G.; Spring, D. R.; King, R. W.; Schreiber, S. L.; Foley, M. A. A one-bead, one-stock solution approach to chemical genetics: Part 2. *Chem. Biol.* **2001**, *8* (12), 1183–1195.
- (6) Lam, K. S.; Lebl, M.; Krchnak, V. The “One-Bead-One-Compound” Combinatorial Library Method. *Chem. Rev.* **1997**, *97* (2), 411–448.
- (7) Stockwell, B. R.; Haggarty, S. J.; Schreiber, S. L. High-throughput screening of small molecules in miniaturized mammalian cell-based assays involving post-translational modifications. *Chem. Biol.* **1999**, *6* (2), 71–83.
- (8) Wagner, B. K.; Carrinski, H. A.; Ahn, Y. H.; Kim, Y. K.; Gilbert, T. J.; Fomina, D. A.; Schreiber, S. L.; Chang, Y. T.; Clemons, P. A. Small-molecule fluorophores to detect cell-state switching in the

context of high-throughput screening. *J. Am. Chem. Soc.* **2008**, *130* (13), 4208–4209.

(9) Cho, C. F.; Behnam Azad, B.; Luyt, L. G.; Lewis, J. D. High-throughput screening of one-bead-one-compound peptide libraries using intact cells. *ACS Comb. Sci.* **2013**, *15* (8), 393–400.

(10) Asano, K.; Funayama, Y.; Yatsuzuka, K.; Higashiyama, Y. Spherical particle sorting by using droplet deflection technology. *J. Electrostat.* **1995**, *35* (1), 3–12.

(11) Christensen, C.; Groth, T.; Bruun Schiødt, C.; Tækker Foged, N.; Meldal, M. Automated sorting of beads from a “one-bead-two-compounds” combinatorial library of metalloproteinase inhibitors. *QSAR Comb. Sci.* **2003**, *22* (7), 737–744.

(12) Gibbs, S. L.; Xie, Y.; Goodwill, H. L.; Nasr, K. A.; Ashitate, Y.; Madigan, V. J.; Siclovan, T. M.; Zavodszky, M.; Tan Hehir, C. A.; Frangioni, J. V. Structure-activity relationship of nerve-highlighting fluorophores. *PLoS One* **2013**, *8* (9), No. e73493.

(13) Hyun, H.; Bordo, M. W.; Nasr, K.; Feith, D.; Lee, J. H.; Kim, S. H.; Ashitate, Y.; Moffitt, L. A.; Rosenberg, M.; Henary, M.; Choi, H. S.; Frangioni, J. V. cGMP-compatible preparative scale synthesis of near-infrared fluorophores. *Contrast Media Mol. Imaging* **2012**, *7* (6), 516–524.

(14) Lee, J. H.; Choi, H. S.; Nasr, K. A.; Ha, M.; Kim, Y.; Frangioni, J. V. High-throughput small molecule identification using MALDI-TOF and a nanolayered substrate. *Anal. Chem.* **2011**, *83* (13), 5283–5289.

(15) Lee, J. H.; Hyun, H.; Cross, C. J.; Henary, M.; Nasr, K. A.; Oketokoun, R.; Choi, H. S.; Frangioni, J. V. Rapid and Facile Microwave-assisted surface chemistry for functionalized microarray slides. *Adv. Funct. Mater.* **2012**, *22* (4), 872–878.

(16) Lee, J. H.; Park, S.; Hyun, H.; Bordo, M. W.; Oketokoun, R.; Nasr, K. A.; Frangioni, J. V.; Choi, H. S. High-throughput screening of small molecule ligands targeted to live bacteria surface. *Anal. Chem.* **2013**, *85* (7), 3508–3514.