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1 Formation of assembloids by DNA-mediated 2 synthetic cell self-assembly

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17 biophysics, DNA nanotechnology
18

19 **Abstract:** Approaches in tissue engineering, organoid culture, and organs-on-chip have
20 propelled the development of increasingly sophisticated *in vitro* models of human tissues.
21 However, as they are formed from natural cells, it is challenging to control their molecular
22 composition and biophysical properties, increasing variability, and limiting their robustness. To
23 overcome these limitations, we introduce a self-assembly strategy for synthetic cells that
24 enables the formation of millimeter-sized synthetic constructs based on single synthetic cells.
25 Specifically, we functionalize the lipid membrane of synthetic cells with cholesterol-tagged
26 single-stranded DNA aptamers, which drive programmable intercellular adhesion through
27 sequence-specific hybridization. This allows individual synthetic cells to interconnect into
28 higher order 3D constructs. By varying aptamer complementarity, internal architecture with
29 spatially distinct functional zones and tuneable mechanical properties can be encoded. Most
30 importantly, the DNA-driven self-assembly operates directly in cell culture medium, is
31 compatible with high-throughput microwell formats enabling scalable screening workflows and



1 is reversible by DNA displacement. To demonstrate biological functionality of these synthetic
2 tissues, we incorporate T cell-stimulatory antibodies into spatially segregated tissue regions.
3 This design mimics lymph node organization and supports infiltration of natural primary human
4 T cells, which subsequently expand within the synthetic tissue. Together, these results
5 establish a route to tissue-scale matrices built from synthetic cell collectives and represent a
6 critical step toward functionally integrating living and non-living matter.

7

8 **Introduction**

9 Synthetic tissue constructs are defined as minimalistic replicas of natural tissues, mimicking
10 their functional and structural properties, which are built not from natural but from individual
11 synthetic cells [1]. Similar to natural tissues, their functional properties arise from their
12 biochemical, mechanical and structural characteristics, which are emergent properties from
13 the synthetic cell collective [2]. The research field of synthetic cell engineering received
14 significant attention in the last decades and achieved several critical milestones. For example,
15 giant unilamellar vesicles (GUVs) were designed to act as functional immune-inspired
16 synthetic cells with adoptable lipid membrane [3] or to serve as cargo transport system [4].
17 Colloidosomes, synthetic cells with a semipermeable silica membrane surrounding an
18 aqueous core, have been designed to receive and react to chemical signals as well as to
19 reconstitute minimalistic metabolic processes [5]. Other examples for synthetic cell systems
20 include coacervates, formed by liquid-liquid phase separation, which can act as membraneless
21 synthetic organelles within a membrane-based synthetic cells [6, 7]. In our previous work we
22 developed synthetic cells termed droplet supported lipid bilayer (dsLBs) composed of a lipid
23 bilayer membrane which is supported by a polydimethylsiloxane (PDMS) oil core. DsLBs cover
24 a tuneable stiffness range from ~1 kPa to ~3000 kPa by varying by varying their internal oil
25 core cross-linking degree. The dsLB lipid membrane is a lateral mobile bilayer, as confirmed
26 by nitrobenzoxadiazole assay, cryoTEM and fluorescent recovery after photobleaching. DsLB
27 have sizes of 3.75 μm ($\pm 2.61 \mu\text{m}$) and PDI of 0.58. To achieve biochemical functionality, the



1 dsLB membrane can be functionalized with antibodies and other proteins for controlled
2 interactions with living cells, including activation and proliferation of high-quality T cells and
3 hybrid spheroids [8, 9].

4 Synthetic cells are generally considered to have high translational potential for application in
5 biomanufacturing, advanced cell culturing and tissue engineering [10]. However, following this
6 realm, transforming synthetic cell technologies from dispersed synthetic cell solution to
7 condensed organized synthetic cell collectives in 3D formats, remains a major frontier in the
8 field [1]. Importantly, adding such a 3D characteristic to advanced cell culture systems has
9 previously proven to be a major advantage for natural cell cultures, most prominently in stem
10 cell and oncogenesis analysis [11-14], where organoids have become powerful and highly
11 relevant model systems. The field is now progressing even further, from generating individual
12 organoids that replicate specific tissue structures to creating assembloids, which integrate
13 multiple organoid types into coherent, multifunctional constructs that better capture the
14 complexity of living organs[15].

15 Formation of multicellular 3D synthetic assembloids from synthetic cells remains challenging
16 as this requires three main features that need to be integrated: biochemical functionality,
17 hierarchical organization and physiological biomechanical properties [16]. Looking at natural
18 cell cultures, matrigel has been a gold standard to mimic exclusively the extracellular matrix
19 (ECM) properties of tissues for studies concerning embryonic or malignant cells [17, 18]. Even
20 though this system is tuneable to some degree in terms of integrated growth factors, proteins,
21 enzymes or structural components such as collagen [19], it is limited by batch-to-batch
22 variability [20]. Many naturally derived 3D culture systems share these limitations
23 [21]. Therefore, synthetic cells could meet this need to form synthetic constructs from the
24 bottom-up to introduce more controllability, reliability and tunability to 3D culture systems
25 thereby not only mimicking the ECM component of tissues but also cells in a tissue
26 environment.



1 3D culturing systems are particularly attractive for *ex vivo* expansion of T cells, which is a
2 critical step in adoptive cell therapy. Therefore, several 3D systems have been developed to
3 culture, activate and expand T cells. These include synthetic hydrogels that can be tuned in
4 their crosslinking degree leading to different stiffnesses [22] or equipped with biochemical cues
5 such as immune cell stimulating beads [23]. Even though hydrogels offer many advantages
6 such as their degradability, viscoelasticity and bioprinting compatibility [24-27], hydrogels are
7 bulk polymer networks that omit mimicry of cells within a tissue complex and the hierarchical
8 microanatomy of tissues. We therefore recently engineered dsLB-based syntactic cells to form
9 3D synthetic tissues which are able to interact with natural T cells [28]. The self-assembly is
10 based on cross-linking individual synthetic cells *via* biotin-conjugated lipids in their membrane
11 with streptavidin [21]. This allows to introduce hierarchical structure and tissue-like mechanical
12 properties. When integrated with T cell-agonistic antibodies, the resulting millimeter-sized 3D
13 lymphatic bottom-up tissues (lymphBUTs) act as minimalistic, controllable and tuneable
14 natural lymph node replicas into which T cells can infiltrate, migrate, activate and differentiate
15 into regulatory-like CD8⁺ T cells [21].

16 However, streptavidin-biotin-based lymphBUTs are still limited in three key features. 1) Their
17 size is limited due to the formation process, thus leading to relatively small constructs with
18 diameters of approximately 2-3 mm, which is smaller than many physiological tissue
19 architectures. 2) Routinely used cell culture media contains soluble biotin, which interferes with
20 streptavidin-based synthetic cell cross-linking, conventional lymphBUTs cannot be formed in
21 cell culture medium. This limits their integration with living cell handling technologies based on
22 bioprinting or organs-on-chip. 3) The formation of lymphBUTs with different reactions zones,
23 in analogy to natural tissues, where different cell types form microphysiological environments
24 (e.g. B cell and T cell zones in the lymph node), is only possible by mixing pre-formed synthetic
25 cell clusters without control over the final spatial arrangements and their sizes. This does not
26 allow to form constructs with more than two different zones, much less than most natural
27 tissues.



1 We reasoned that using single stranded DNA (ssDNA) instead of multivalent streptavidin as
2 an intercellular adhesion protein analogue to interlink individual synthetic cells could overcome
3 the limitations detailed above [29, 30]. Moreover, new DNA-based formation techniques allow
4 for continuous growth of constructs to larger sizes. Also, self-assembly of dispersed synthetic
5 cells *via* DNA hybridization can be performed in physiological, high ionic strength buffers such
6 as cell culture media. Furthermore, by using ssDNA strands of varying complementary
7 individual zones can be formed under controlled conditions. Lastly, controlled disassembly of
8 3D constructs was achieved by providing higher affinity displacement strands thus restoring the
9 status of dispersed synthetic cells. Therefore, we established formation of constructs based
10 on ssDNA and evaluated their structure and function as assembloids for *ex vivo* T cell
11 expansion.

12

13 Results

14 Construct formation *via* ssDNA

15 To leverage the advantage of single-stranded DNA (ssDNA) as selective synthetic cell
16 intercellular adhesion protein analogue, we first integrated ssDNA in the dsLB membrane *via*
17 a cholesterol anchor. The 20bp long anchor strand was subsequently coupled to a partly
18 complementary ssDNA (linker strand - linker 1a, linker 1b) (**Figure 1 A**) to interconnect dsLBs
19 and form constructs (**Figure 1 B, C**). During ssDNA-based formation of these constructs we
20 achieved significantly larger sizes of individual constructs than for our previous biotin-based
21 ones. The new assembly strategy, which allows construct formation by sedimentation-based
22 contact formation between individual dsLBs, evades mechanical stress by orbital mixing which
23 allows for the formation of bigger-sized constructs. We compared construct sizes between the
24 two formation approaches, confirming that ssDNA-based constructs (**Figure 1 C**). We first
25 verified the integration of the anchor 20bp-Cy5 strand into the dsLB membrane and
26 subsequent linker 1a 20bp-Cy3 strand hybridization by flow cytometry. Varying ssDNA
27 concentrations were tested in the range of 3 mol% to 15 mol% with the corresponding linker



1 ssDNA in an equimolar ratio. Quantification of the mean fluorescence intensity (MFI) showed
2 robust integration across all concentrations tested (**Figure 1 D**). Integrating the anchor ssDNA
3 did also not lead to phase separation in the dsLB lipid bilayer membrane indicated by a
4 homogenous fluorophore distribution in the membrane shown in the confocal microscopy
5 images (**Figure 1 E**). Most importantly, when ssDNA functionalized dsLBs were incubated in
6 Eppendorf tubes, they spontaneously formed millimeter-sized free-floating constructs (**Figure**
7 **S 1 A**). When dsLBs were incubated without linker strands (anchor strands only), no construct
8 formation was observed, demonstrating the specific DNA-pairing between the individual
9 synthetic cells as a requirement for the self-assembly (**Figure S 1 C**).

10



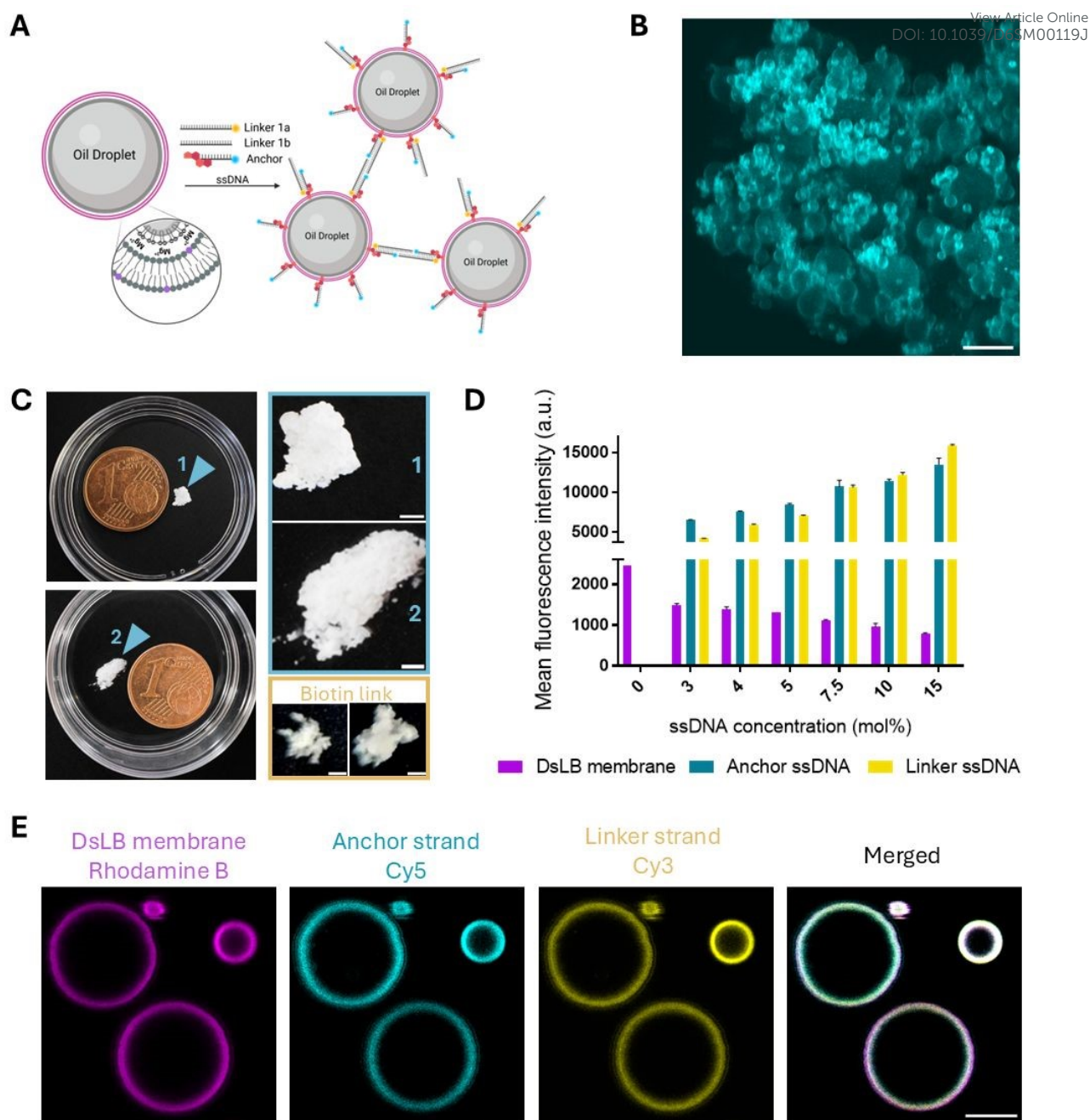


Figure 1: Establishment of a construct formation strategy using ssDNA **A)** Schematic illustration of inter-dsLB binding *via* partly complementary ssDNA. **B)** Representative confocal microscopy maximal z-projection showing a construct assembled from individual dsLBs using 20bp ssDNA (dsLB membrane: Cy5). Scale bar is 20 μm . **C)** Top view of representative constructs formed *via* ssDNA hybridization (blue) or biotin-streptavidin bond (yellow) [21] next to a 16.25 mm diameter 1 Eurocent for size. Scale bars 1 mm. **D)** Flow cytometry-based quantification of mean fluorescence intensities of the dsLB membrane (Atto 488 lipids), anchor 20bp ssDNA (Cy5) and linker 1a 20bp ssDNA (Cy3) at different concentration of ssDNA including 3 mol%, 4 mol%, 5 mol%, 7.5 mol%, 10 mol% and 15 mol%. Results are shown as mean SD, from 2 replicates. **E)** Representative confocal microscopy image of single dsLBs (Rhodamine B lipid in membrane) decorated with anchor 20bp ssDNA (cyan – Cy5) and linker 1a 20 bp ssDNA (yellow – Cy3). Scale bar is 5 μm .

12

13 We next evaluate the integration of ssDNA with varying length by testing 15bp and 25bp
 14 strands. We define the ssDNA length by the number of complementary nucleotides between



1 anchor and linker strands (**Table 1**). The MFIs of the dsLB membrane and the anchor ssDNA
2 were measured by flow cytometry. Individual synthetic cells demonstrate successful ssDNA
3 integration in both cases (**Figure 2 A**). Of note, the shorter ssDNA strands (15bp) showed
4 lower MFI for the ssDNA, while displaying higher MFI for the Rhodamine B fluorophore
5 attached to the head group of 1 mol% of the lipids in the dsLB membrane (**Figure S 1 B**). This
6 is indicative for a static quenching effect between DNA and the membrane fluorophore due to
7 proximity [31, 32]. Integration of ssDNA of both lengths was further confirmed by confocal
8 microscopy images (**Figure 2 B**).

9 Importantly, construct formation was observed with ssDNA of both lengths (15bp and 25bp)
10 while no formation was detected with only the corresponding anchor stands (**Figure 2 C**,
11 **Figure S 1 D**). The main objective of using DNA as a synthetic cell cross-linker is that
12 complementarity and oligo length can be applied to program the self-assembly as well as
13 generating constructs and assembloids with spatially defined biochemical and potential
14 biomechanical properties. We reasoned that the number of inter-dsLB connections (ssDNA
15 concentration) could impact the overall synthetic tissue stiffness as we previously observed for
16 biotin-streptavidin connections [21]. Therefore, we evaluated the mechanical properties of the
17 constructs by microplate parallel compression analysis. For this, constructs formed with either
18 15bp, 20bp or 25bp and 3 mol%, 5 mol% or 10 mol% ssDNA were produced. The maximal
19 compression force required to compress 10% of the total construct height was measured
20 (**Figure 2 D**). The measurements indicate that ssDNA concentrations between 3 mol% and 10
21 mol% have a minor impact on the mechanical properties of the constructs. However, constructs
22 assembled with shorter ssDNA strands (15bp) tend to increase the stiffness with higher ssDNA
23 concentration beginning at 10 mol% in the dsLB membrane. Contrary, the ssDNA
24 concentration does not affect the stiffness of constructs formed with longer ssDNA (20bp and
25 25bp) (**Figure 2 E - G**). We additionally tested constructs formed with 20bp ssDNA and ssDNA
26 concentrations of 2 mol% and 20 mol% and detected significantly lower compression force
27 needed to compress 10% of the total construct height, indicating reduced total construct
28 stiffness (**Figure S 1 E**). To optimally balance cost efficacy and construct stability, we chose 5



1 mol% and 20bp ssDNA length for next experiments. The decreased stiffness for 2 and 20
2 mol% is potentially rooted in the fact that compressibility is mostly a function of the number of
3 interactions between individual dsLBs in the construct and not necessarily of their binding
4 strength [21]. Of note, the observed variations in compression force in all conditions are,
5 among other factors, a result of the construct orientation.

6 We further aimed to verify that construct formation *via* ssDNA can be performed in serum-
7 supplemented cell culture medium, in contrast to streptavidin-based lymphBUTs. For this, the
8 ssDNA functionalized synthetic cell suspensions were diluted into RPMI 1640 cell culture
9 medium and construct formation was observed (**Figure S 1 F**). This enables the integration of
10 natural cells into biochemically functionalized constructs further, referred to as synthetic
11 tissues, under physiological environments and opens the door for cell handling procedures
12 such as bioprinting or organs-on-chip.

13 Next, we aimed to explore the potential to perform DNA-based operations with the constructs
14 to alter structure or function *in situ*. Several operations, including allosteric gates, hybridization
15 chain reactions or strand displacement can be performed with DNA. We implemented strand
16 displacement for switchable disassembly of constructs. For this we added competing linker
17 DNA with higher affinity (dis strand) to the constructs formed with anchor 20bp + linker 11bp
18 1a/1b or 13bp mm 1a/1b to displace 200% of the original linker DNA. By confocal time lapse
19 microscopy, we observed total disassembly of the constructs within 10 min (**Figure S 1 G**).

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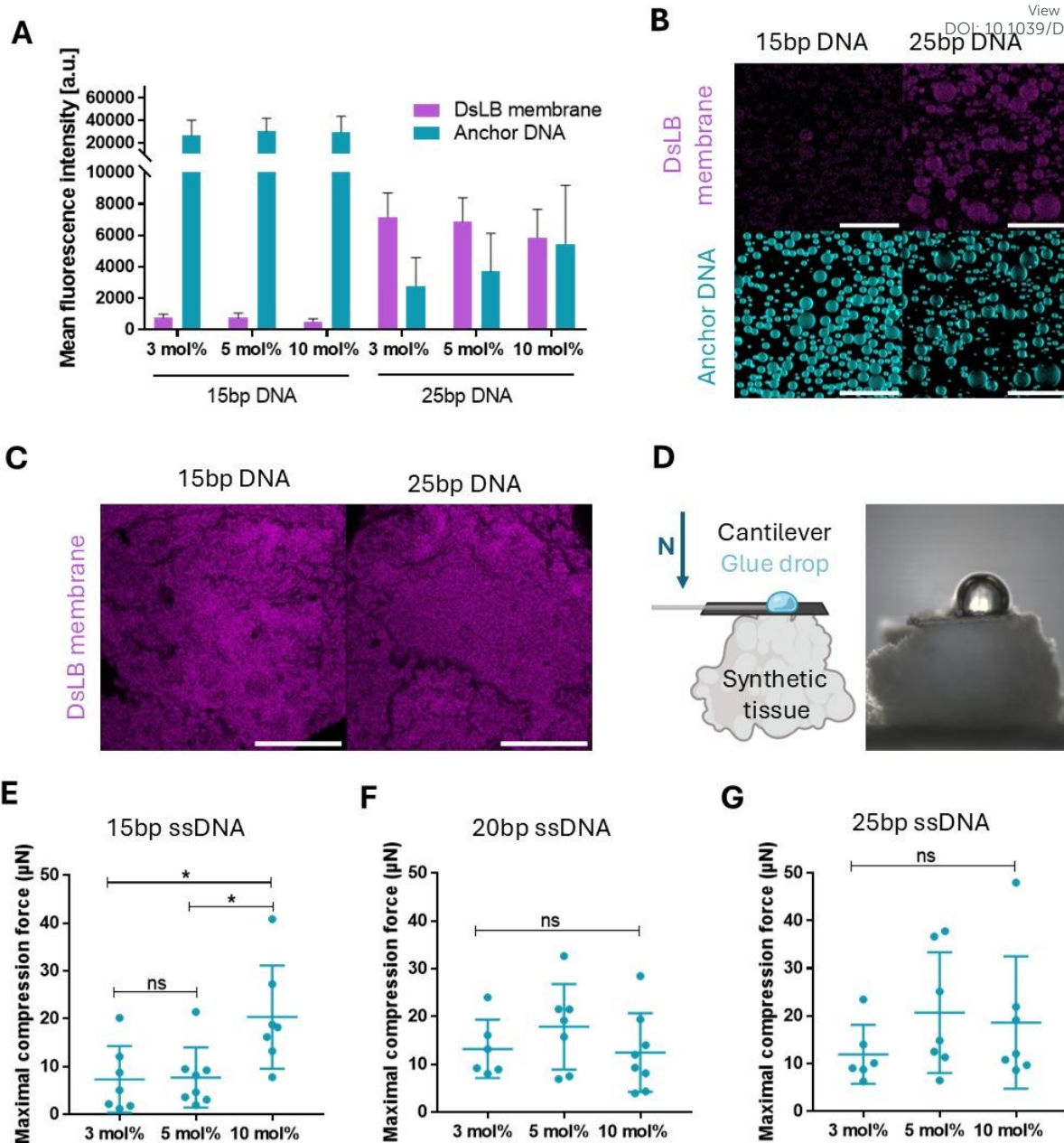


Figure 2: Mechanical properties of ssDNA-based constructs **A**) Flow cytometry measurements showing the mean fluorescence intensity of dsLBs (magenta) equipped with anchor ssDNA with 15bp and 25bp at different ssDNA concentrations of 3 mol%, 5 mol% and 10 mol%. **B**) Representative confocal microscopy z-projection of dsLBs (magenta) and anchor 15bp and 25bp ssDNA (cyan). Scale bar is 50 μm . **C**) Representative confocal microscopy z-projection of dsLB (magenta) assembly into constructs by anchor 15bp and 25bp ssDNA strands + corresponding linker 1a/1b. Scale bar is 500 μm . **D**) Representative bright field image of a construct assembled with 5 mol% 20bp ssDNA in the parallel plate micro-compression setup and the corresponding schematic illustration. **E**) – **G**) Maximal compression force measurement of constructs assembled from ssDNA with **E**) 15bp **F**) 20bp and **G**) 25bp at different ssDNA concentrations of 3 mol%, 5 mol% and 10 mol% using parallel micro-compression. Results are shown as \pm SD of $n \leq 6$ constructs. p values were calculated using two-tailed t test, ns = not significant $p > 0.05$, $^*p < 0.05$.

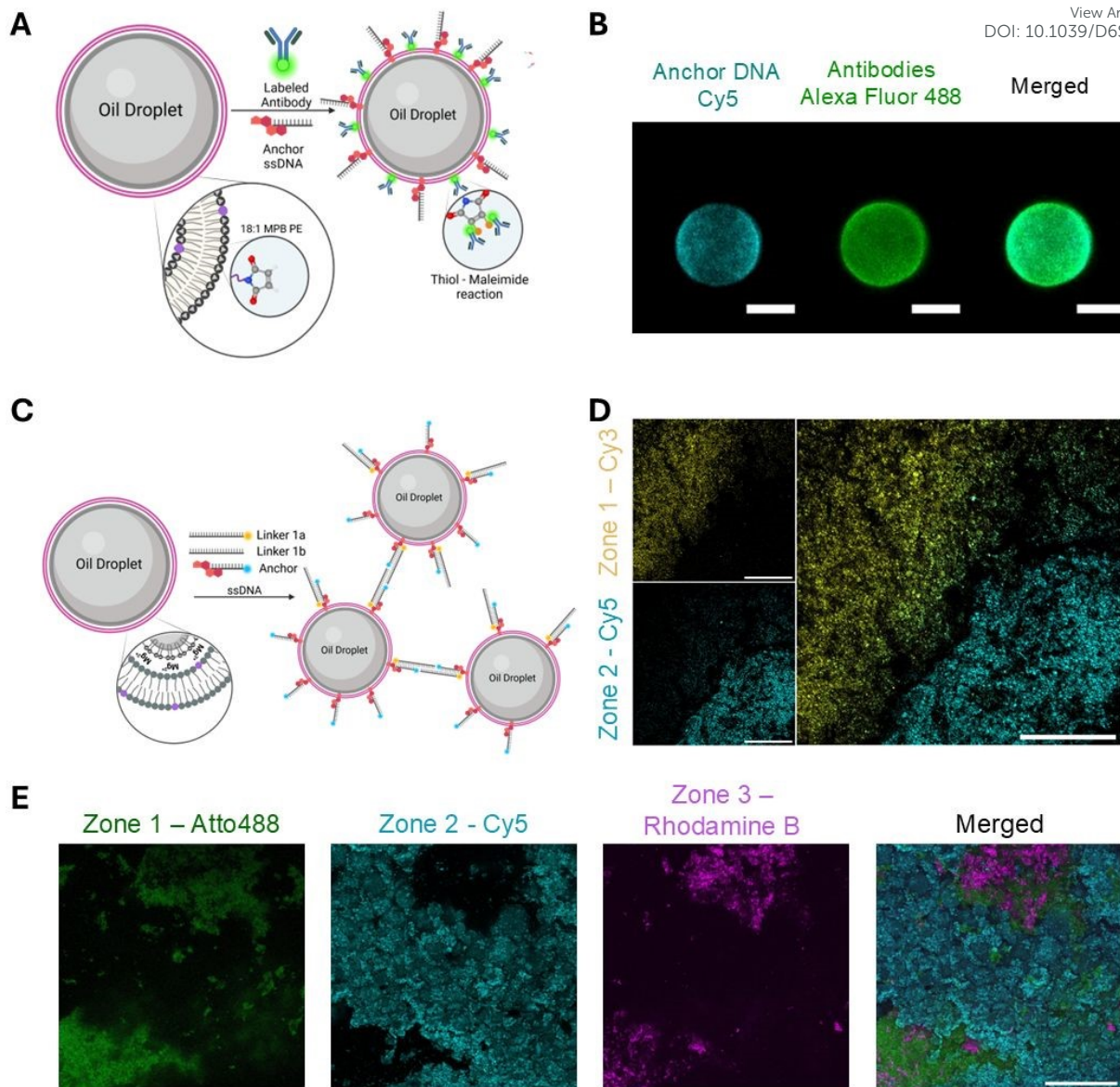
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14 Towards introducing biochemical functionality into the constructs, we next coupled T cell-
 15 stimulating IgG antibodies, specifically anti-CD3 and anti-CD28, to the dsLB membrane via



1 lipids with maleimide headgroups (**Figure 3 A**). These immune-functionalized constructs are
2 further referred to as synthetic tissues. Confocal microscopy imaging of an AlexaFluor488-
3 conjugated IgG confirmed successful functionalization of the synthetic cell surface and no
4 competitive binding of ssDNA and antibodies (**Figure 3 B**). The functionalization of dsLB
5 membranes with ssDNA enabled the formation of complex architectures based on the innate
6 high specificity of DNA hybridization, allowing highly directed integration of multiple dsLB
7 systems (plain, immune-stimulating, etc.), regardless of membrane components, into a single
8 multi-zone synthetic tissue (**Figure 3 C**). Again, confocal microscopy imaging confirmed the
9 formation of synthetic tissues with multiple zones of different constructs visualized by the
10 ssDNA combination of anchor 20bp + linker 1b+1a-Cy3 (yellow) or linker 2b+2a-Cy5 (cyan)
11 (**Figure 3 D**). Of note, we observed preserved structural integrity of constructs over a period
12 of 2 weeks at 4 °C (**Figure S 2 A**). This approach leverages the high specificity of DNA
13 hybridization and programmable ssDNA design to enable the construction of multicomponent,
14 hierarchical constructs. These mimic the morphology and microanatomy of real tissues and
15 comprise multiple distinct zones (**Figure 3 E**).
16





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2 **Figure 3: Zone formation for constructs and immune-functionalized synthetic tissues** **A)** Schematic
3 illustration of biochemical dsLB functionalization with antibodies *via* Thiol-Maleimide reactions. **B)** Representative
4 confocal microscopy z-projection of anchor 20bp-Cy5 ssDNA strand (cyan) and Alexa Fluor 488 labelled antibodies
5 (green). Scale bar is 5 μm . **C)** Schematic illustration of multi-zone construct formation using different linker ssDNA
6 connections. **D)** Representative confocal microscopy z-projection of constructs formed and visualized by the ssDNA
7 combination of anchor 20bp + linker 1b+1a-Cy3 (yellow) forming zone 1 or linker 2b+2a-Cy5 (cyan) forming zone
8 2. Scale bar is 250 μm . **E)** Representative confocal microscopy z-projection of multi-zone construct formation
9 visualized by fluorophore-labelled lipids in the dsLB membrane: zone 1 – Atto 488 (dsLB membrane: green) formed
10 with anchor 25bp + linker 1a/1b 25bp, zone 2 – Cy5 (dsLB membrane: cyan) formed with anchor 20bp + linker
11 1a/1b 20bp and zone 3 – Rhodamine B (dsLB membrane: magenta) formed with anchor 20bp + linker 2a/2b 20bp.
12 Scale bar is 250 μm .

13

14 For high-throughput screenings or potential industrial applications, a more parallelized method
15 for construct formation is required. To address this, we implemented a microwell plate
16 formation approach. For this, each well of a 96-well plate served as a template for a “pancake”
17 like construct covering the entire well (**Figure 4 A**). This strategy presents a more robust



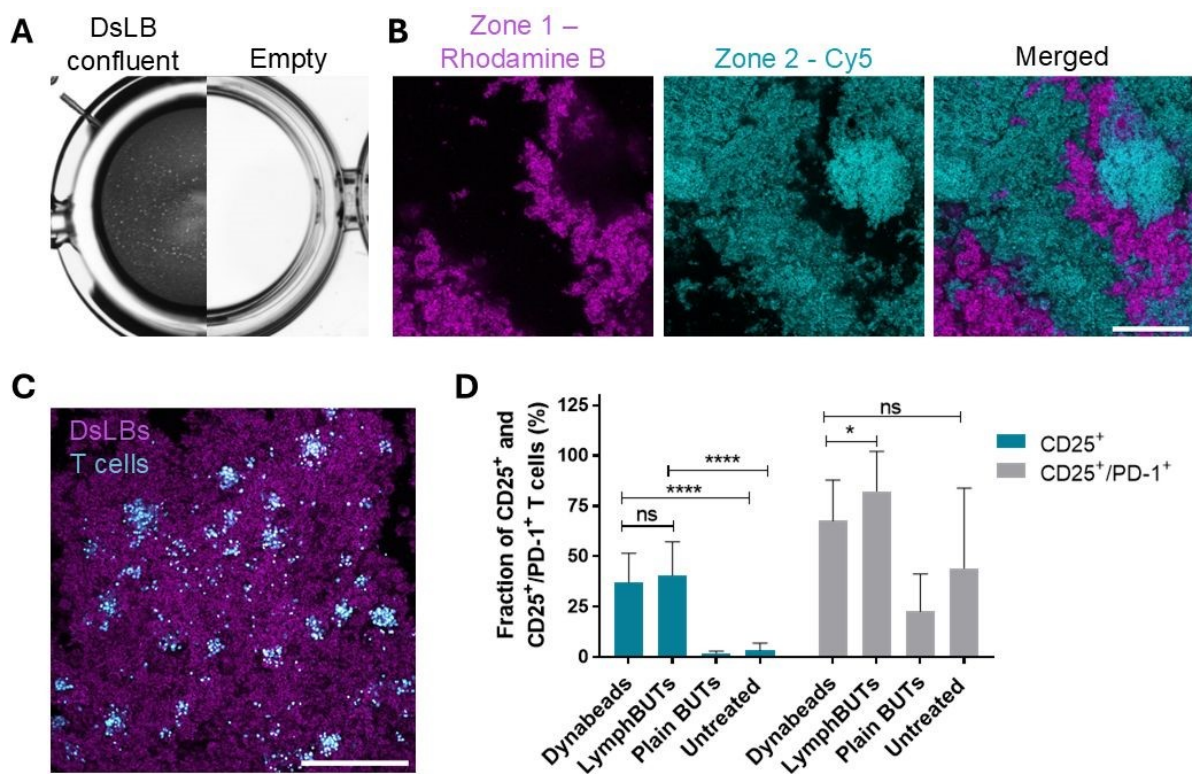
1 method and allows to maximize contact formation between natural cells incubated within the
2 synthetic tissue. Synthetic tissue formation and zone organization was confirmed for this setup
3 by stereo- and confocal microscopy (**Figure 4 A, Figure 4 B**).

4 Next, we evaluated the applicability of DNA-based synthetic tissues as culturing system for
5 living human immune cells from healthy donors. For this, we determined the DNA-based
6 construct stability in cell culture medium. Confocal microscopy confirmed structural integrity of
7 the constructs after 4 days in fully supplemented cell culture medium (**Figure S 2 B**).
8 Furthermore, to verify the structural integrity of the constructs under cell culture conditions (i.e.
9 cell culture medium enriched with DNase from serum components), we incubated constructs
10 varying concentrations of DNase. We observed no construct disassembly at low (100 $\mu\text{g}/\text{mL}$)
11 DNase concentrations after 4 days incubation at 37 °C in fully supplemented T cell medium.
12 However, with higher (500 $\mu\text{g}/\text{mL}$) DNase concentrations the constructs disassembly. During
13 construct disassembly, single dsLBs return to the dispersed state, thus increasing the apparent
14 construct size and total dsLB covered area measured by microscopy. The total construct area
15 doubled upon addition of 500 $\mu\text{g}/\text{mL}$ DNase after 4 days of incubation which indicated
16 decreased structural integrity. (**Figure S 2 C**). Additionally, immune-functionalized synthetic
17 tissues remain structurally stable over 4 days of incubation in PBS at 4 °C or fully
18 supplemented medium at 37 °C indicated by no changes in construct area (**Figure S 2 D**).

19 Finally, primary human CD8⁺ T cells were seeded on the DNA-based immune-functionalized
20 synthetic tissues and incubated for 4 days. The immune-functionalized synthetic tissues which
21 host natural cells are also referred to as assembloids. Synthetic tissues with two distinct zones,
22 one antibody-functionalized zone and one unfunctionalized scaffolding zone for structural
23 support, were applied. The functionalized zones comprised 75% of the synthetic cell mix and
24 the scaffolding zone 25%. For both zones, 5 mol% density of 20bp ssDNA on the synthetic
25 cells was used. Staining for T cell nuclei after the incubation and inspection by confocal
26 microscopy reveals clonal expansion clusters of T cell proliferated within the 3D assembloids
27 after 4 days (**Figure 4 C**). Of note, our previous study revealed preferential co-localization of
28 T cells in immune-functionalized zones [21]. Flow cytometry analysis of CD25 expression



1 confirmed the activation of T cells within the assembloids. The CD25 expression was
 2 benchmarked against the industry gold standard, Dynabeads. Both T cell activation
 3 approaches exhibited comparable activation rates of about 40% of the total T cell population.
 4 These results align with the T cell activation we observed in biotin-streptavidin lymphBUTs in
 5 our previous [8, 21], while T cells expanded in the assembloids also exhibited increased
 6 expression of the immunosuppressive receptor PD-1 (**Figure 4 D**).



9 **Figure 4: Synthetic tissue standardization and assembloid application** **A)** Representative stereo microscope
 10 image of a dsLB confluent (“carpet” antibody functionalized constructs) and an empty 96 well. **B)** Representative
 11 confocal microscopy z-projection of multi-zone synthetic tissues visualized by fluorophore-labelled lipids in the dsLB
 12 membrane: zone 1 Rhodamine B (dsLB membrane: magenta) formed with anchor 20bp + linker 1a/1b and zone 2
 13 Cy5 (dsLB membrane: cyan) formed with anchor 20bp + linker 2a/2b. Scale bar is 250 μ m. **C)** Exemplary
 14 representative confocal microscopy z-projection of T cells (nuclei in cyan) infiltrated and formed clonal proliferation
 15 cluster within an assembloid formed with anchor 20bp + linker 1a/ab (dsLB membrane: magenta) after 4 days of
 16 co-cultivation. **D)** Quantification of the CD25⁺ and CD25⁺/PD-1⁺ expressing T cell fraction expanded with Dynabeads
 17 or DNA-based assembloids measured by flow cytometry. Results are shown as mean \pm SD of four donors in n > 2
 18 technical replicates. p values were calculated using two-tailed t test. ns = not significant p > 0.05, *p < 0.05,
 19 ****p < 0.0001.

20

21



1 **Summary and discussion**

2 In this study, we introduce a new approach for assembloid formation that enables controlled
3 and reliable intercellular connections between individual synthetic cells. Inspired by
4 approaches to engineer intercellular binding *via* ssDNA nanotechnology [33, 34], we designed
5 multizonal synthetic tissues. Contrary to our previously established biotin-streptavidin
6 approach, an almost covalent bond that is not programmable in structural self-assembly,
7 partially complementary ssDNA enables the formation of multiple spatial zones in more defined
8 and modular manner. Each zone can be equipped with a distinct protein or antibody portfolio
9 precisely matching the biological requirements of the co-cultured natural cells, ultimately
10 forming millimetre-sized synthetic tissues.

11 The use of ssDNA provides programmability beyond complementarity, enabling systematic
12 control over strand length and surface density on the synthetic cells. This mirrors natural cell
13 collectives, where cells dynamically regulate the expression level and type of adhesion
14 molecules to control processes such as epithelial-mesenchymal transition [35]. Consistent with
15 this concept, our results show that mechanical compressibility and therefore emergent
16 mechanical construct properties, can be tuned through ssDNA surface density, in a minimalist
17 manner. Additionally, we could characterize the relevance of DNA binding strength (i.e. linker
18 length) and density, specifically for the 20bp linker length system with respect to lowest and
19 highest ssDNA concentration of 2 and 20 mol%. Constructs formed with 2 mol% ssDNA tend
20 to be mechanically more fragile, which was indicated by the lower indentation force needed to
21 compress 10% of the total construct height. In contrast, we did not observe decreased
22 structural stability of constructs formed with 20 mol% ssDNA while at the same time having
23 low stiffness. We hypothesize that this either originates from incomplete incorporation of
24 ssDNA in the dsLB membrane or steric hinderance by too many ssDNA strand in the inter-
25 dsLB space. Additionally, DNA strand-displacement could be employed to restructure
26 assembloids *in situ* and releasing the hosted natural cells without subjecting them to excessive
27 mechanical stress.



1 Moreover, using ssDNA facilitates construct assembly under cell culture conditions, such as
2 standard cell culture medium supplemented with serum, which commonly contains free biotin
3 and therefore interferes with the biotin-streptavidin linkage. Also, the structural integrity of
4 constructs and synthetic tissues in fully supplemented medium remains over the total
5 incubation time of 4 days. By further implementing a 96-well plate compatible formation and
6 culture format, we developed a faster, more standardized, and scalable platform capable of
7 hosting cells within a 3D environment.

8 The hybridization and stability of ssDNA is primarily influenced by two major factors, cation
9 concentration and the presence of digestive nuclease present in fetal bovine serum (FBS),
10 thus making cell culture medium a harsh environment for DNA and especially ssDNA
11 nanotechnology. Previous studies have shown that reducing FBS concentrations to 1% and
12 increasing Mg^{2+} levels can prolong DNA stability [36]. Additionally, encapsulating DNA
13 nanostructures within lipid bilayers has been reported to protect against nuclease-mediated
14 digestion [37]. However, we observed stable ssDNA interactions, indicated by preserved
15 synthetic tissue and zone integrity, for several days in fully supplemented cell culture medium
16 and even for weeks under long term storing conditions (PBS, 4 °C). We hypothesize that the
17 unexpectedly high stability originates from a synergistic combination of 1) enhanced stability
18 by the lipid bilayer integration 2) respectively short ssDNA length, 3) locally enhanced Mg^{2+}
19 concentration within the dsLB membrane, and 4) close proximity between the dsLBs
20 membranes creating a DNase excluded zone.

21 We demonstrated biological applicability of these new 3D culture environments by using them
22 to activate and expand primary human $CD8^+$ T cells. Their biological performance was
23 benchmarked against commercially available activation beads (Dynabeads) as they are the
24 established standard in both clinical and academic applications due to their strong and rapid
25 activation and broad commercial availability. Thus, we successfully applied a self-assembling
26 synthetic tissue which microanatomic organization is directed by partially complementary
27 ssDNA hybridization. Particularly, the parallelized well plate formation approach, in which the
28 single synthetic cells can function as “synthetic cell ink”, emphasizes the substantial potential



1 for future industrial and academic applications, including high-throughput drug or toxicity
2 screenings. By tuning the biological functionality of individual synesthetic cells, synthetic
3 tissues can function as custom-tailored 3D cell culture environments, not just for immune cells
4 such as natural killer cells, neutrophils or lymphocytes but also for fibroblasts or stem cells.

5 Since construct formation *via* ssDNA hybridization enables the formation of assembloids in a
6 well-defined and controlled manner, this approach builds a foundation for introducing synthetic
7 cell morphogen gradients within a 3D environment [38]. To achieve this and to mimic structural
8 and functional heterogeneity of biological organoids, additional synthetic cell types such as
9 DNA-conjugated GUVs or coacervates can be incorporated, each leveraging their individual
10 advantages to form emergent multicellular systems.

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1 Experimental section

2

3 DsLB production

4 DsLBs were produced using the established and reported protocols [8, 21]. Briefly, 100 mg of
5 PDMS (Sylgard 184, Dow Corning USA) were pre-emulsified in 900 μL of PBS (adjusted to pH
6 7 with HCl) and 4 mM of sodium dodecyl sulfate (SDS, Sigma–Aldrich, Germany). The oil in
7 water pre-emulsion was further emulsified in a sonification bath for 2 min. By adding 22 mM of
8 MgCl_2 (Sigma–Aldrich, Germany) a cationic layer was formed around the oil droplets which
9 binds anionic small unilamellar vesicles (SUVs) of which 100 μL (6 mM) were added, mixed
10 and incubated for 2 min protected from light. Excess SUVs were removed by three washing
11 steps repeating centrifugation at 10000 $\times g$ for 30 sec, discharging of the supernatant and
12 resuspension in PBS adjusted to pH 7.

13 The SUVs were produced using extrusion as described previously [8]. They consist of 20 mol%
14 EggPG, 5 mol% 18:1 MPB PE (maleimide), 1 mol% LissRhodamine B-PE, Atto488-PE or 18:0
15 Cy5 PE and EggPC acting as a filling lipid (all Avanti Polar Lipids, USA).

16

17 DsLB immune functionalization

18 A maleimide functionalized lipid (Avanti Polar Lipids, USA) was integrated with 5 mol% in the
19 SUVs and furthermore in the dsLB membrane enable the attachment of antibodies and
20 proteins *via* thiol connections. To define the total accessible amount of maleimide in the dsLB
21 membrane, a SUV 1:1 dilution series ranging from 60 μM to 0.47 μM was prepared in a 96-
22 well plate and used for calibration. The dsLB suspension was diluted 1:10 with PBS into the
23 well plate [8, 21].

24 The dsLB fluorescent intensity was measured with a TECAN Spark plate reader (Tecan Group,
25 Switzerland) controlled by TECAN SparkControl software with in-built gain optimization. The



1 excitation/emission settings were adjusted to Rhodamine B (537/582 nm), Atto488-PE
 2 (495/545 nm) or CY5-PE (630/675 nm) labelled dsLBs [21].

3 Based on the total lipid membrane concentration and therefore accessible maleimide
 4 headgroups, the immune-stimulating antibodies against human CD3 (UCHT1, Invitrogen) and
 5 CD28 (CD28.2, Invitrogen) at a ratio (maleimide/antibody) of 0.5 high (~1020 molecules/ μm^2)
 6 and a 1:3 ratio between $\alpha\text{CD3}/\alpha\text{CD28}$ were attached to the dsLB membrane. In order to
 7 remove sodium azide which, impacts the pH and therefore the efficiency of the thiol –
 8 maleimide reaction, the antibodies were washed using the MircoSpin™ G-25 Columns (Cytiva,
 9 US) according to manufacturer instructions. The washing leads to a reduction of antibody
 10 concentration of 9% (+/-1%) which was considered for all calculations. The dsLBs were
 11 incubated with the antibodies in PBS adjusted to pH 7 at RT for 1 h resuspending every 15
 12 min. To get rid of un-bound antibodies the immune- dsLB were centrifuged at 10,000 xg for 30
 13 sec and resuspended in PBS. The dsLB's membrane concentration was adjusted to 150 – 300
 14 μM with the help of the total lipid concentration measured using the plate reader [8, 21].

16 DsLB ssDNA functionalization and construct formation

17 By integrating single stranded DNA (ssDNA) *via* cholesterol residues into the antibody-
 18 functionalized dsLB membrane and subsequent addition of partly complementary linker DNA,
 19 single dsLBs can be connected into constructs in a very controlled manner. Following ssDNA
 20 strands, inspired by [39], were used driving the construct formation and are accordingly named
 21 in the main text.

23 Table 1: SsDNA sequences (5' → 3') with the corresponding name all purchased from Biomers GmbH, Germany

Name	Sequence 5' → 3'
Linker 1a 11bp	5'/TTCTTGTGAACACCTAGCTCTTTCAGTATCTG/3'
Linker 1b 11bp	5'/TTCTTGTGAACCAGATAGTGAAAGAGCTAGGA/3'



Linker 1a 13bp mm	5'/CTTACTTGTGAACACCTAGCTCTTTCACATCTG/3'	View Article Online DOI: 10.1039/D6SM00119J
Linker 1b 13bp mm	5'/CTTACTTGTGAACCAGATAGTGAAAGAGCTAGGA/3'	
Anchor 15bp-Cy5	Cy5/5'/GTTCAACAAGAAAGCGTTTT/3'/Chol	
Anchor 15bp	5'/GTTCAACAAGAAAGCGTTTT/3'/Chol	
Linker 1a 15bp	5'/CGCTTTCTTGTGAACACTCTTTCACATCT/3'	
Linker 1b 15bp	5'/CGCTTTCTTGTGAACAGATAGTGAAAGAGA/3'	
Anchor 20bp	5'/GTTCAACAAGAAAGCGGAAGCTTTTT/3'/Chol	
Anchor 20bp-Cy5	Cy5/5'/GTTCAACAAGAAAGCGGAAGCTTTTT/3'/Chol	
Linker 1a 20bp-Cy3	Cy3/5'/GCTTCCGCTTTCTTGTGAACACCTAGCTCTTTCACATCTG/3'	
Linker1a 20bp	5'/GCTTCCGCTTTCTTGTGAACACCTAGCTCTTTCACATCTG/3'	
Linker 1b 20bp	5'/GCTTCCGCTTTCTTGTGAACCAGATAGTGAAAGAGCTAGGA/3'	
Linker 2a 20bp-Cy5	Cy5/5'/GCTTCCGCTTTCTTGTGAACACTGATCTAATGGTGCACGTT/3'	
Linker 2a 20bp	5'/GCTTCCGCTTTCTTGTGAACACTGATCTAATGGTGCACGTT/3'	
Linker 2b 20bp	5'/GCTTCCGCTTTCTTGTGAACAACGTGCACCATTAGATCAGA/3'	
Anchor 25bp-Cy5	Cy5/5'/GTTCACAGATGAAGAAAGCGGAAGCTTTTT/3'/Chol	
Anchor 25bp	5'/GTTCACAGATGAAGAAAGCGGAAGCTTTTT/3'/Chol	
Linker 1a 25bp	5'/GCTTCCGCTTTCTTCATCTGTGAACACCTAGCTCTTACGCATCACTATCTG/3'	
Linker 1b 25bp	5'/GCTTCCGCTTTCTTCATCTGTGAACCAGATAGTGATGCGTAAGAGCTAGGA/3'	
Dis strand	5'/AAAGCTTCCGCTTTCTTGTGAAC/3'	

1

2 Single stranded DNA (ssDNA) was functionalized with cholesterol in order to integrate the

3 anchor ssDNA into the dsLB membrane *via* self-assembly in concentrations ranging from 2 –

4 20 mol% based on the total lipid membrane concentration determined as previously described.

5 To achieve lymphBUT formation, the highly concentrated dsLBs (150 – 300 μ M) were

6 incubated with the anchor ssDNA as well as ssDNA linker a and ssDNA linker b using a ratio

7 equimolar ratio of 1:1:1 for 1 h at RT within a 0.5 mL microtube. The total reaction volume was

8 kept between 20 – 50 μ L. During the incubation period the microtube was placed upside down

9 to prevent dsLB loss due to sticking to the tube walls, facilitating construct formation at the



1 PBS/air interface. To produce consistently sized constructs 40 μL of a 150 μM dsLBs solution
2 was constant over the experiments. The total lipid concentration was determined as previously
3 described. The formed constructs were gently flushed with 200 μL of PBS or cell culture
4 medium and allowed to rest for another 1 h at RT.

5

6 **Multi-zone construct formation**

7 Multi-zone constructs were assembled through a sequential process in which multiple dsLB or
8 construct subunits were connected to yield a single, continuous, functional construct. Each
9 subunit can be formed from differently functionalized dsLBs carrying immune-stimulating
10 antibodies or different membrane fluorophores, as well as labelled ssDNA strands and ssDNA
11 strands length of 15bp, 20bp or 25bp. Assembly was performed in either PBS adjusted to pH
12 7 or in cell culture medium, as specified below.

13 **Microtube formation method:** For the multi-zone construct formation subunits were either
14 formed with (activating zone) or without (scaffolding zone) immune-stimulating antibodies in
15 separate 0.5 mL Eppendorf tubes. Briefly, we applied a sequential addition approach where
16 partially adding linker a and linker b to dsLBs already leads to the formation of sub-zonal
17 constructs. After combining those sub-zonal constructs the rest of the linker a and b is added
18 to reach the aimed ratio of anchor: linker a and linker b. This formation happens in four steps
19 and results in rounded sphere-like geometries.

20 Step 1: The dsLB membrane was functionalized with anchor ssDNA (5 mol% of the total lipid
21 membrane) and anchor ssDNA was partially saturated (50%) with linker a and b ssDNA. This
22 mixture was incubated for 30 min in a total reaction volume of 20 -50 μL .

23 Step 2: Subsequently, the formed constructs were flushed with either 200 μL of sterile PBS or
24 cell culture medium and incubated at RT for 1h.

25 Step 3: The activating and the scaffolding subunits were combined in one 0.5 mL Eppendorf
26 tube and excess PBS or cell culture medium was removed to obtain a final volume of $\sim 50 \mu\text{L}$.



1 To initiate the inter-subunit connection, the second 50% of linker a and b of ssDNA was added
2 to achieve a 1:1:1 ratio of anchor:linker a:linker b.

3 Step 4: The Eppendorf tube was incubated upside down, protected from light, for 1 h at RT.
4 The resulting multi-zone constructs were subsequently washed by flushing with 200 μ L of PBS
5 or cell culture medium and stored at 4 $^{\circ}$ C until further use.

6 **Well plate formation method:** For the combination of multiple dsLB subtypes into a multi-
7 zone dsLB agglomerate a flat-bottom 96-well plate served as a physical template that
8 promoted planar pancake-like construct formation. Immuno-functionalized dsLBs (75% of the
9 synthetic tissue) and plain dsLBs (25% of the synthetic tissue) were mixed with anchor and
10 both linker ssDNA in individual microtubes before being transferred dropwise into the 96 well.
11 Each drop later presents a zone. The ssDNA concentration for cell culture experiments was
12 always 5 mol% and a ssDNA length of 20bp. The dsLBs mixtures formed into synthetic tissues
13 after 1 h incubation at RT protected from light. The synthetic tissues can then be stored at 4
14 $^{\circ}$ C until further use.

16 **Laser scanning confocal microscopy imaging of constructs**

17 The fluorescently labelled dsLB membrane and labelled ssDNA as well as dsLB agglomerate
18 referred to as constructs or assembloids were visualized using the confocal laser scanning
19 microscope LSM 880 (Carl Zeiss AB) equipped with a 10x objective (EC "Plan-Neofluar"
20 10x/0.30 M27, Carl Zeiss AG, Germany) and 63x immersion oil objective (Plan-Apochromat
21 63x/1.4 oil DIC M27, Carl Zeiss AG, Germany) with the 405, 488 and 633 laser lines. The
22 primary human CD8⁺ T cells were fixed with 2% PFA for 30 min and the T cell nucleus was
23 stained with Hoechst 33342 trihydrochloride (Thermo Fisher, Germany). For this the fixed T
24 cells were incubated with 1 μ M Hoechst 33342 for 30 min and washed by replacing half of the
25 volume three times. The images were analysed with the ImageJ software (NIH, USA) by snaps
26 or z-projection of stacks and background subtraction [8, 21].



1

2 **Construct disassembly using ssDNA higher affinity displacement strands**

3 Constructs were assembled following previously described protocols using anchor 20bp and
4 linker 11bp or 13bp mm ssDNA strands. To perform strand displacement the environmental
5 ionic strength from PBS or cell culture medium needs to be reduced. For this the constructs
6 were washed by two times sequential exchange of PBS/medium by ddH₂O and subsequently
7 incubated for 30 min at RT. For construct disassembly, a higher affinity ssDNA strand
8 complementary to the anchor strand (disassembly strand, “Dis strand”) was added in 200% of
9 the anchor. The samples were monitored over 45 min at 37 °C in a 1 min interval *via* confocal
10 laser scanning microscope LSM 880 (Carl Zeiss AB).

11

12 **Mechanical construct properties**

13 Micro-compression was used to evaluate the mechanical properties of lymphBUT formed with
14 different ssDNA concentrations (2 mol%, 3 mol%, 5 mol%, 10 mol% and 20 mol%) as well as
15 different ssDNA length (15bp, 20bp, 25bp). For this, the constructs were transferred onto a
16 testing anvil within a CellScale fluid bath filled with 45 mL. The cantilever was assembly
17 manually by glueing a 1x1 mm square stainless-steel platen to a tungsten microbeam with a
18 diameter of 0.0762 mm and a modulus of 411000 MPa which was mounted into the
19 Microindenter G2 CellScale (CellScale biomaterials testing, Canada). The constructs were
20 compressed with a compression magnitude of 10% of the total construct height measuring the
21 indentation force. The z-compression was controlled in a ramp setting with a loading duration
22 of 30 sec, a holding duration of 5 sec and a recovery duration of 30 sec. Real time imaging
23 was performed with a USB digital camera with a zoom lens and XYZ automated stage tracked
24 at a frequency of 5 Hz [21].

25

26 **T cell isolation and cultivation**



1 Functional assays were performed with primary human CD8⁺ T cells. The T cells were isolated
2 from leukapheresis reduction system chambers from healthy voluntary blood donors using
3 negative isolation. The isolation was performed with commercial selection kits (RosetteSep
4 Human CD8⁺ T cell Enrichment Cocktail, STEMCELL technologies, Germany) following the
5 manufacturers instruction. The blood was provided by the Institute for Clinical Hemostaseology
6 and Transfusion Medicine, Saarland University Medical Center provided the blood according
7 to the ethics agreement number 34/23 (Ethikkommission Ärztekammer des Saarlandes). The
8 isolated CD8⁺ T cells were stored in RPMI 1640 w/ L-Glutamine (VWR, Germany) medium
9 supplemented with 40% fetal bovine serum (Gibco, Germany), 1% penicillin/streptomycin
10 (Gibco, Germany), 1% non-essential amino acids (Biowest) and 50 mM HEPES (Sigma-
11 Aldrich, Germany) and additional 10% dimethyl sulfoxide (DMSO) (Sigma Aldrich, Germany)
12 and stored at -80 °C until further use [8, 21].

13 The primary human CD8⁺ were thawed and cultivated overnight in RPMI 1640 w/L-Glutamine
14 (VWR, Germany) medium supplemented with 10% fetal bovine serum (Gibco, Germany), 1%
15 penicillin/streptomycin (Gibco, Germany), 1% non-essential amino acids (Biowest, France)
16 and 50 mM HEPES (Sigma-Aldrich, Germany) with 100 U mL⁻¹ of the growth factor Interleukin
17 2 (IL-2) in T 25 cell culture flasks at 37 °C and 5% CO₂ [8, 21].

19 **T cell *ex vivo* activation and expansion within synthetic tissues**

20 100,000 – 120,000 isolated CD8⁺ T cells were added to the synthetic tissues produced
21 according to the well plate formation method and filled with fully supplemented medium to a
22 total volume of 200 µL and a total IL-2 concentration of 100 U mL⁻¹. The cells were then co-
23 cultivated at 37 °C and 5% CO₂ for 4 days with the synthetic tissues. The wells at the outer
24 border were filled with PBS in order to avoid evaporation and therefore biased results.
25 Dynabeads human T-Activator CD3/CD28 beads (Gibco, Germany) used according to
26 manufacturer suggestions served as controls [21].

27



1 **CD8⁺ T cell activation studies using Flow Cytometry**

2 To further study primary human CD8⁺ T cell activation and immunosuppression signal the
3 assembloids were resuspended in order to separate dsLBs and T cells and centrifuged at 300
4 xg for 5 min. The supernatant was disposed and the cells were resuspended in PBS + 1%
5 Albumin fraction V (BSA) (Sigma–Aldrich, Germany) containing staining antibodies (1:400)
6 against the surface marker CD25 (BC96, BioLegend, UK) and PD-1 (NAT105, BioLegend, UK)
7 conjugated to Alexa Fluor 488 or Alexa Fluor 647. The cells were stained with the antibodies
8 for 1 h at RT protected from light. Subsequently the cells were washed once with PBS,
9 resuspended in PBS containing 2% PFA (Sigma–Aldrich, Germany) and incubated for 30 min
10 at RT protected from light followed by washing step with PBS. After disposing the remaining
11 PFA the cells were resuspended in PBS containing 1 nM Hoechst 33342 trihydrochloride
12 (Thermo Fisher, Germany) for staining the T cell nucleus for 30 min protected from light.
13 Finally, the cells were washed once more and resuspended in PBS + 1 BSA and resuspended
14 in a final volume of 200 μ L PBS + 1% BSA and stored at 4 °C until further flow cytometer
15 measurements. For the quantification of the surface marker the Attune NxT Flow Cytometer
16 and with the Attune™ Software (Thermo Fisher, Germany) and equipped with 405, 488, 561
17 and 637 nm laser lines were used. For the analysis a minimum of 10,000 events were
18 considered and analysed with the FlowJo V.10 software (FlowJo LLC, USA) [8, 21].

20 **Shelf-life stability of multi-zone constructs**

21 Multi-zone construct formation is based on surface interactions induced by complementary
22 ssDNA between the activating and scaffolding zones. The time dependent structural and
23 mechanical stability of multi-zone constructs was evaluated by confocal microscopy. For this,
24 constructs made from two zones visualized with either Cy5 or Cy3 labelled linker ssDNA were
25 formed as previously described and transferred to a 48-well suspension plate with 200 μ L of
26 PBS. The constructs were evaluated according to their mechanical stability and zone integrity
27 over 2 weeks (n=2).



1

2 **Metabolic degradation resistance of ssDNA constructs and synthetic tissues in cell** 3 **culture medium**

4 SsDNA used for construct formation could potentially be targeted for degradation by e.g.
5 nucleases in the cell culture medium, thus leading to structural disassembly of the constructs
6 and synthetic tissues. To evaluate this, constructs and synthetic tissues were formed in
7 Eppendorf tubes with 5 mol% and 20bp ssDNA, transferred to 48-well plates and stored in fully
8 supplemented RPMI 1640 medium over 4 days at 37 °C or in PBS at 4 °C. Structural stability
9 was analysed *via* confocal microscopy for 4 days ever 24 h (n=4) and the LEICA DFC450
10 stereo microscope (Leica Microsystems, Germany) and a 2x (PLANAPO 2.0x/39, Leica
11 Microsystems, Germany) or a 5x objective (PLANAPO 5.0x/0.5, Leica Microsystems,
12 Germany). Stereo microscopic images were analysed as mentioned below.

13 Furthermore, constructs produced with the well-plate strategy and incubated in fully
14 supplemented RPMI 1640 medium with additional 100 µg/mL and 500 µg/mL DNase I (Roche
15 Diagnostics GmbH, Germany) at 37 °C for 1h and 96 h. The constructs structural integrity was
16 evaluated by the total construct area imaged with a LEICA DFC450 stereo microscope (Leica
17 Microsystems, Germany) and a 2x (PLANAPO 2.0x/39, Leica Microsystems, Germany) or a
18 5x objective (PLANAPO 5.0x/0.5, Leica Microsystems, Germany). The images were analysed
19 with the ImageJ software (NIH, USA) by global-threshold segmentation and automated particle
20 detection. Low structural integrity was identified by increasing total construct area.

21

22 **Data processing and statistical analysis**

23 Graphs were plotted with GraphPad Prism 7 as mean +/- SD of technical and biological
24 replicates. Statistical analyses were performed with the in-build GraphPad Prism 7 software
25 function. The applied statistical analysis was noted for the individual figures in the figure
26 legends. Schematic illustrations were created with BioRender.com [8, 21].

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Competing interests

The authors declare no competing interests.

Authors contributions

A.B. performed and designed experiments, analysed the data, supervised experimental implementation and wrote the manuscript. E.A.L.L. performed experiments and analysed data. G.M.H. performed experiments and analysed data. K.J. designed the ssDNA and helped in protocol development. O.S. designed the study, supervised the experimental implementation and wrote the manuscript.



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Data availability statement

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All data underlying this study is available in the figures or supplementary information. No specific analysis code or software has been developed for this study and no publicly available data has been used.

