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2 FRONT MATTER

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4 Title

5 Shape-Morphing Living Composites

6 Authors

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18 **One Sentence Summary:** Baker's yeast embedded in a hydrogel enables the engineering of
19 smart materials that change shape in response to specific cues.

20
21 **Abstract**

22
23 This work establishes a means to exploit genetic networks to create living-synthetic composites that
24 change shape in response to specific biochemical or physical stimuli. Baker's yeast embedded in a
25 hydrogel forms a responsive material where cellular proliferation leads to a controllable increase in
26 the composite volume of up to 400%. Genetic manipulation of the yeast enables composites where
27 volume change on exposure to L-histidine is 14 \times higher than volume change when exposed to D-
28 histidine or other amino acids. By encoding an optogenetic switch into yeast, spatiotemporally-
29 controlled shape change is induced with pulses of dim blue light (2.7 mW/cm²). These living, shape-
30 changing materials may enable sensors or medical devices that respond to highly specific cues
31 found within a biological milieu.

32
33 **MAIN TEXT**

34
35 **Introduction**

36
37 Materials that change shape enable mechanical activity in devices, such as smart garments, sensors,
38 microfluidics, or drug-delivery platforms (1-4). In these devices, traditional actuators, like
39 solenoids, are too large, heavy, or power intensive to be used. Shape change in synthetic polymers
40 and gels can be triggered using temperature, electric fields, or chemicals (5-8). The specificity of
41 the response is dictated, and limited, by the physical characteristics of the material (9). One
42 approach to induce specificity in the physical characteristics of a hydrogel is to build polymer
networks from biomacromolecules, such as DNA, allowing for detection of analytes that directly
bind to these constituents (10). Binding of designed DNA sequences can induce 100-fold
volumetric hydrogel expansion by successive extension of cross-links, using a DNA hybridization
cascade. In the design of chemically-responsive hydrogels, this mechanism is limited to detection
of analytes capable of highly specific binding motifs. In living organisms, direct DNA binding is
not the typical mechanism by which sensing occurs. Genetic information in cells encodes

43 components that enable appropriate responses to a wide range of specific chemical and physical
44 cues.

45 Composites that combine the tunable properties of synthetic materials and the responsive nature of
46 living organisms represent a powerful strategy to imbue multifunctionality in a single material.
47 Several living composites have been previously reported including self-healing concrete (11),
48 ethanol-producing 3D-printed hydrogels (12), gels that self-heal using photosynthesis (13), and
49 wearable fluorescent biosensors (14). However, these living composites lack the ability to respond
50 mechanically to environmental cues. One example of a mechanically-active living composite is a
51 bilayer of an elastomer and mammalian muscles that bends through contraction and relaxation of
52 the muscle cells (15). However, muscle cells only thrive over a very narrow set of conditions,
53 limiting the range of applications where these materials can be used. Notably, shape change in
54 living organisms is not limited to contraction of muscles. Tissue morphogenesis in animals and
55 plants is controlled in part by cellular proliferation (16, 17). However, a strategy that harnesses
56 proliferation of living cells to control the shape change of synthetic materials has yet to be reported.

57 Here we describe hybrid materials where living *Saccharomyces cerevisiae* (i.e., Baker's Yeast or
58 Brewer's Yeast) embedded within a polyacrylamide hydrogel proliferates in response to a
59 combination of environmental cues, which induces shape change in the composite (Fig. 1A). By
60 controlling cell loading or hydrogel stiffness, we control the magnitude of volume change in the
61 composites. This shape change is further controlled by patterning proliferation within a monolith.
62 Critically, yeast provide a versatile platform for genetic engineering of the conditions required for
63 proliferation. Using this control, we design composites that respond only in the presence of a single
64 chirality of a single amino acid or to brief pulses of dim visible light. We harness this shape change
65 to create microfluidic channels that respond selectively to fluids flowing through the channel.
66

67 Results and discussion

68 *S. cerevisiae* is an ideal model organism to realize responsive, living composites. These unicellular
69 organisms thrive within solid matrices (12), are much stiffer (1-10 MPa) (18) than many hydrogels
70 (10-100 kPa) (19), and are known to survive over a wide range of conditions (20). Our key
71 observation is that as these stiff cells proliferate within a solid hydrogel matrix, a global increase in
72 volume is observed. We hypothesize that this volume increase is not due to ordinary swelling of a
73 hydrogel, but instead is attributable to local displacement of the hydrogel by the newly formed cells.
74 After the composite is exposed to the appropriate conditions for cell growth, a dramatic increase in
75 cell count can be observed (Fig. 1B). To quantify the effect of proliferation on macroscopic volume
76 change, living composites were polymerized with 0.9-1.1 billion cells/mL of pre-gel solution (6
77 wt% of dry yeast). Composites were incubated in YPD (yeast extract, peptone, D-glucose) media
78 at 30 °C for 48 h. YPD contains the necessary nutrients for the yeast, and as such, cell proliferation-
79 induced shape change occurs, resulting in a change in area of $124.2\% \pm 10.3\%$ and a volume change
80 of $200.9\% \pm 2.4\%$ (Fig. 1C, movie S1). By incubating composites in media without a fermentable
81 carbon source (D-glucose), the area of the disk only increased by $6.3\% \pm 0.4\%$ as the yeast are
82 incapable of proliferation (Fig. 1D, fig. S1). Similarly, hydrogels without encapsulated yeast
83 incubated in YPD only undergo a volume change of $1.2\% \pm 0.5\%$ (fig. S2). The shape change of
84 the living composites in rich media is also not attributable to passive cell size changes; cell viability
85 is required for shape change to occur. We pattern cell viability, using UV light (254 nm) exposed
86 through a mask, in living composites covalently bound to glass (Fig. 1E). Only the regions of the
87 hydrogel not exposed to UV, the letters "ALIVE," contain viable cells and undergo a volume
88 increase on exposure to YPD. This expansion is greater than 110% of the initial film thickness after
89 36 h (Fig. 1F, fig. S3). Shape change is accompanied by a change in topography, from smooth to
90 rough, as the growing colonies deform the surface in a heterogeneous manner at the sub-mm scale.

91 The described experiments show that yeast proliferation is the primary mechanism associated with
92 volume change in these hybrid living materials.

93 Proliferation-driven shape change can be tuned by controlling the initial composition of the living
94 composite (Fig. 2A, 2B). On varying the initial concentration of yeast from 1 wt% to 18 wt%, the
95 volume change after 48 h increases from $123.8\% \pm 3.9\%$ to $337.2\% \pm 17.4\%$. This volume change
96 is accompanied by a concomitant increase in dry mass, which varies from $177\% \pm 11\%$ to $320\% \pm$
97 35% (Fig. 2A). While we observe that some cells escape from the composite and proliferate in the
98 media, this increase in dry mass suggests that most of the cells are retained in the hydrogel matrix.
99 We also note that the shape of the grown composites is largely stable for over 128 days in deionized
100 water at room temperature (fig. S4). The increase in dry mass and shape stability further supports
101 our hypothesis that CO_2 production or passive swelling from the hydrogel matrix is not the
102 mechanism responsible for shape change. This mass change represents new material that can be
103 produced on demand with only as much external intervention or equipment as is needed to ferment
104 grape juice. In the case of materials with 18 wt% yeast, the solid components of the as-synthesized
105 composites are 35.9% polymer and 64.1% yeast. After growth, the yeast content increases to 85.6%.
106 These growing composites may provide opportunities to produce materials directly from renewable
107 feedstocks or even waste streams (21).

108 The mechanical properties of the hydrogel matrix also control the proliferation-induced shape
109 change. By altering the feed ratio of crosslinker from 0.05 wt/v% to 0.6 wt/v%, at constant yeast
110 loading (6 wt% dry yeast) and acrylamide concentration (10 wt/v%), the Young's modulus of the
111 composites after synthesis increases from $8 \text{ kPa} \pm 1 \text{ kPa}$ to $204 \text{ kPa} \pm 16 \text{ kPa}$. As stiffness increases,
112 the volume change during cell proliferation decreases from $255.8\% \pm 7.3\%$ to $107.9\% \pm 1.2\%$ (Fig.
113 2B, fig. S5). We attribute this decrease to increased elastic resistance to the expanding colonies,
114 perhaps resulting in limited cell proliferation. Given the tradeoffs between composite stiffness,
115 yeast loading, and volume change, we selected composites with 0.1 wt/v% crosslinker and 6 wt%
116 yeast for further studies as these composites have relatively high initial elastic modulus and large
117 stimulus-response (fig. S6).

118 Spatial control of volume change can be programmed to yield composites that morph controllably
119 from 2D to 3D. Informed by prior work where spatially-controlled swelling is used to guide shape
120 selection in hydrogels (22-26), we fabricated composite disks (12 mm diameter and 0.5 mm
121 thickness) and used UV light (254 nm) to kill cells in programmed areas (insets of Fig. 2C, 2D).
122 After irradiation, no shape change is observed when the hydrogel is equilibrated in water, indicating
123 that passive swelling of the gel is not significantly altered. After incubation in YPD, spatially-
124 controlled proliferation induces a 2D to 3D transformation. The flat disk shown in Figure 2C grows
125 in area in the center of the disk, while being constrained around the perimeter, resulting in a
126 hemispherical cap (+ Gaussian curvature). By contrast, the disk depicted in Figure 2D grows along
127 the perimeter, while being constrained in the center, resulting in a saddle-like geometry (- Gaussian
128 curvature).

129 Programming of the stimulus that induces shape change of living composites can be achieved by
130 genetic manipulation of the yeast. *S. cerevisiae* is a model eukaryote commonly used for
131 heterologous protein expression (27-29). The yeast strain we employ (L40) is deficient in L-
132 histidine metabolism. This metabolic feature, termed auxotrophy, prevents proliferation in the
133 absence of L-histidine in the growth environment (Fig. 3A). This strain was used to fabricate
134 composites that morph into 3D helical shapes only in the presence of L-histidine. Rectangular free-
135 standing films were patterned with UV light to cause cell death in the areas indicated in Fig. 3B.
136 Incubation for 48 h in selective media lacking L-histidine did not affect the shape of the composite.
137 When these composites were incubated in media containing L-histidine, the flat films morph into a
138 helix (Fig. 3C). Over 48 h in media lacking L-histidine, disks of these composites only increase in

139 volume by $20.7\% \pm 6.1\%$, despite the other 20 amino acids, D-glucose, and nitrogen base present
140 in the media. The same composites were then incubated for another 48 h in otherwise identical
141 media containing L-histidine, resulting in a volume change of $278.3\% \pm 12.9\%$ (Fig. 3D, fig. S7).
142 Critically, these composites remain dormant during periods of unfavorable conditions and then
143 respond when conditions match those programmed by the genetics of the yeast. The ability to
144 withstand unfavorable conditions stands in stark contrast to the fragile nature of cells from
145 multicellular organisms. To further demonstrate the biochemical specificity of these composites,
146 volume change of living composites was measured for samples incubated in media containing L-
147 histidine and compared to the volume change of samples grown in media with D-histidine and
148 without any histidine. Much like the composites exposed to media lacking L-histidine, the inclusion
149 of D-histidine into the media did not lead to significant volume change at the end of an incubation
150 period of 72 h, as D-histidine is not naturally incorporated into proteins (Fig. 3E, fig. S7).
151 Previously, hydrogels that swell by recognition of a single enantiomer of a chiral molecule have
152 been achieved through molecular imprinting; however, the volume change in these materials is
153 often less than 20% (30). Notably, the reported living composite exhibits a volumetric change up
154 to $20\times$ larger than the volume change observed in imprinted hydrogels.

155 Microfluidic devices fabricated with living composites respond with specificity to the fluid flowing
156 through the channels. We fabricated responsive microfluidics by using replica molding to form
157 channels within a living composite. All cells were rendered inviable with UV light except for the
158 cells within two of the microfluidic channels (Fig. 3F). Flow of a fluorescent fluid is then used to
159 visualize the performance of these devices. Before flowing media containing L-histidine, the
160 channels are all open, and the intensity of the fluorescence is similar across the device (Fig. 3G).
161 After flowing media with L-histidine through the inlet, the two microchannels with viable cells
162 grow in volume resulting in the channels becoming blocked, while the other channels remained
163 open. By contrast, similar devices exposed to otherwise identical media lacking L-histidine have
164 all channels open (fig. S8). These smart microfluidic devices could enable new strategies for
165 biosensors that directly manipulate the flow of fluid without external intervention, traditional
166 sensors, or actuators. While materials that respond to biochemical cues are ideal for devices that
167 change autonomously with their environment, diffusion limits the ability of these cues to generate
168 on-demand shape change with control in space and time.

169 Optogenetic switches can be engineered into yeast to enable photoresponsive composites, where
170 shape change can be spatiotemporally controlled. We generated a yeast strain to express a
171 photoresponsive transcriptional switch that induces gene expression after illumination with blue
172 light (455 nm). This strain has two *Arabidopsis thaliana* proteins in a yeast two-hybrid system (31).
173 Blue light stimulation induces binding of CRY2 fused to the LexA DNA-binding domain and CIB1
174 Gal4 activation domain chimera. In the presence of light, the HIS3 gene is activated enabling
175 cellular proliferation in the absence of L-histidine (Fig. 4A). We also generated two additional
176 strains: a positive control that does not require light for activation of HIS3 (32) and a negative
177 control that lacks the CIB1 protein and is auxotrophic for L-histidine with or without blue light.
178 Metabolic activity of the experimental strain is over $100\times$ higher when exposed to light than when
179 kept in the dark, as measured by a β -galactosidase assay which probes lacZ, a reporter gene, activity
180 (Fig. 4B, fig. S9). In the negative control strain, metabolic activity is low in blue light and in the
181 dark; while the positive control presented high metabolic activity in both conditions.

182 Optogenetic control of histidine auxotrophy enables photocontrolled cellular proliferation and
183 therefore shape change in living composites. We fabricated composites from each of the three
184 different strains described above and exposed each composite to brief, dim pulses of blue light (2.7
185 mW/cm^2 , 2 s on, 2 min off) or darkness (Fig. 4C). Over 72 h, living composites made with the
186 experimental strain undergo a volume change of $315.9\% \pm 2.1\%$ when exposed to light and a
187 volume change of $86.2\% \pm 5.7\%$ when kept in the dark. (Fig. 4D, 4E). By comparison, the positive

control composites undergo a volume change of over 260% in both light and dark, and negative control composites grow only $103.1\% \pm 20.1\%$ in the light and $10.2\% \pm 5.3\%$ in the dark (Fig. 4E, fig. S9). We note that traditional photoresponsive polymers, which use light to power shape change, typically require irradiation intensities of more than 100 mW/cm^2 (33, 34). By comparison, these living composites rely on light to trigger an optogenetic switch that has been optimized by evolution. The activation of this switch subsequently enables a metabolically-powered change in volume. As a result of this pathway, the time-averaged intensity required is at least $2250\times$ smaller than traditional photoresponsive polymers.

The combination of patterned cell viability and patterned light illumination can be used to provide spatiotemporal control of complex shape change in living composites comprised of transformed yeast with the optogenetic switch. The cell viability within a film of the living composite was patterned using UV light to leave only two circular regions on the film with viable cells (fig. S10). As shown in Figure 2C, proliferation should lead to the formation of a hemispherical cap. By exposing these two regions sequentially, each region sequentially actuates from flat to hemispherical (Fig. 4F).

Living composites undergo cell proliferation-induced shape change controlled by the initial composition of the composite or by patterning regions of viable cells. These materials capitalize on genetic control of biological mechanisms, namely cellular proliferation, to enable responsiveness in topography or shape in response to specific cues. A host of devices from drug delivery platforms to environmental sensors could be enabled by these findings.

Materials and Methods

Materials

Acrylamide, n,n'-methylenebisacrylamide (MBAA), ammonium persulfate (APS), n,n,n',n'-tetramethylethylenediamine (TEMED), 3-amino-1,2,4-triazole (3-AT), L-histidine, adenine sulfate, sulforhodamine B, bisphenol A ethoxylate diacrylate (BPA), and poly(ethylene glycol) diacrylate (PEG-DA) (700 g/mol) were purchased from Sigma-Aldrich. The photoinitiator Irgacure 369 (I-369) was donated by BASF corporation. Methacryloxyethyl thiocarbonyl Rhodamine B (PolyFluor 570) was purchased from Polysciences. Commercial yeast (*Saccharomyces cerevisiae*, Active Dry yeast, Fleischmann's) was purchased from Tom Thumb (Richardson, TX). Yeast extract, yeast nitrogen base without amino acids, peptone, D-(+)-glucose, D-histidine, and trypan Blue were purchased from Fisher Scientific. 3-(Trimethoxysilyl) propyl methacrylate (TPM) was purchased from Acros Organics. Rain-X® was purchased from Wal-Mart (Richardson, TX). All chemicals were used as received without further purification.

Genetically-engineered yeast strains and plasmids

The genotype of the L40 yeast strain is *MATA ade2 his3 leu2 trp1 LYS::lexA-HIS3 URA3::lexA-LacZ* (ATCC, MYA-3332) (35). L40 yeast were transformed with experimental constructs CRY2 LexA DNA-binding fusion in the expression vector pDBTrp (pDBTrp-LexABD-CRY2FL) (Plasmid #78210, Addgene) and a separate CIB1 Gal4 activation domain fusion vector pGADT7 (pGal4AD-CIB1) (Plasmid #28245, Addgene). pDBTrp-LexABD-CRY2FL along with pGADT7 empty vector was used as negative control. In the positive control, we made use of a previously described interaction between *Caenorhabditis elegans* FBF2 (residues 121-C-terminus fused to the Gal4 activation domain present in pGADT7) and CPB1 (residues 1-80 fused to the LexA DNA-binding domain encoded by the pBTM116) (32,36).

237 Mold construction

238
239 For volume change and mechanical testing experiments, molds were made of two glass slides
240 (75x51mm) previously cleaned with Rain-X® to avoid gel adhesion. Slides were separated with 1
241 mm or 500 μ m rectangular spacers, wrapped with parafilm closing one of the open sides, and fixed
242 using binder clips.

243
244 For living composite coatings, molds (75x25mm) with one glass slide cleaned with Rain-X® and
245 one treated with a methacrylate-functionalized silane were assembled. The two glass slides were
246 separated with two 250 μ m polystyrene spacers on each side and fixed with binder clips. For the
247 silane treatment, glass slides were cleaned following a similar process described in the literature
248 (37). Briefly, glass slides were sonicated for 5 min in acetone and isopropanol mixtures and rinsed
249 three times in dH₂O. Afterwards, substrates were sonicated for 30 min in a mixture of water and
250 Alconox cleaner (Alconox Inc. USA), rinsed, and stored in dH₂O overnight. For silanization, glass
251 slides were modified for 30 min with a 5 v/v% mixture of TPM in toluene at 65°C. Then, the slides
252 were rinsed with toluene, dried with N₂ gas, and baked on a hotplate at 120°C for 5 min.

253
254 Determination of cell density

255
256 To determine cell concentrations in active dried yeast, cell density was measured with a UV/Vis
257 spectrophotometer by observing optical density at 660 nm. Briefly, 50 mL mixture with 0.6 g yeast
258 in dH₂O were prepared. Then, a 1:10 dilution was made by mixing 0.1 mL of the mixture with 0.9
259 mL of dH₂O. Diluted samples were pipetted into a 1 mL cuvette for spectrophotometer
260 measurements. Optical densities between 1 and 1.1 were measured, which correspond to numbers
261 of cells of 1.89 x10⁷ and 2.25 x10⁷ cells respectively. These results indicate that the active dried
262 yeast contained between 15 and 18 billion cells per gram.

263
264 Preparation of living-composite materials with active dried yeast

265
266 Polyacrylamide hydrogels with embedded yeast were prepared at room temperature by free radical
267 polymerization of acrylamide monomer and MBAA crosslinker. Stock solutions of 0.4 g/mL
268 acrylamide and 0.02 g/mL MBAA were prepared in dH₂O to create polyacrylamide gel precursor
269 solutions. All pre-gel solutions were prepared with a final concentration of 10 wt/v% acrylamide.
270 Pre-gel solutions were prepared with a final concentration of 0.1 wt/v% MBAA and ~ 1 billion
271 cells/mL of pre-gel solution, unless otherwise noted. To polymerize these solutions, a 10 wt/v%
272 APS stock solution was added at 1% of the total solution volume, and TEMED was added at a ratio
273 of 0.1% of total solution volume. Polymerizing solutions were then vortexed for 3 s and quickly
274 pipetted into molds. Filled molds were flipped every 45 s while polymerization occurred to avoid
275 yeast settling. After 10 min, polymerized living composites were de-molded and rinsed three times
276 with dH₂O to remove unpolymerized acrylamide residues. Living composites were stored in dH₂O
277 for 24 h prior to mechanical testing and volume change experiments. For mechanical testing and
278 volume change experiments, pre-gel solutions were prepared with 0.05, 0.1, 0.3 and 0.6 wt/v%
279 MBAA and 6 wt/v% yeast (~ 1 billion cells/mL of pre-gel solution). To test volume change and
280 Young's modulus with varying yeast content, composites with final concentrations of 0, 1, 6, 12,
281 and 18 wt% yeast were prepared.

282
283 Area, volume and mass change quantification of living-composites embedding active dried yeast

284
285 Living composites embedded with commercial active dried yeast were cut into 10 mm diameter
286 disks. The dimensions of each disk were measured prior to incubation. Samples with varying dry

287 yeast, and crosslinker concentration were incubated at 30°C in YPD rich media without agitation
288 (1% yeast extract, 2% peptone, and 2% D-glucose). To measure area and volume changes, three
289 samples for each composition were incubated in 7 mL of rich media that was changed every 6 h.
290 Area change was measured every hour for 48 h using a MightyScope 5M digital microscope, and
291 volume changes were measured every 24 h for 48 h using a Canon Rebel T5i camera.

292 Dry mass change was obtained by weighing samples with varying yeast content before and after
293 cell proliferation. Briefly, one set of living composites that was not exposed to media was dried at
294 30°C under vacuum to allow water evaporation. An identical set of living composites was incubated
295 in YPD media for 48 h with media change every 6 h and then dried under the same conditions.
296 Upon drying, samples were weighed, and mass was measured. Data presented are an average of 3
297 samples per composition.

299 300 Material characterization

302 Samples (3 mm x 3 mm x 1 mm) were cut from polymerized living composites with varying yeast
303 and crosslinker content after equilibration in dH₂O. Compression testing was performed using a
304 MicroSquisher (CellScale biomaterials testing). Briefly, a tungsten beam 1.016 mm in diameter
305 was glued to a 6 mm x 6 mm platen on one end. This compliant beam was attached to an actuator
306 with a cantilever beam grip at the opposite end from the platen. Samples were loaded to the test
307 chamber filled with dH₂O at room temperature. The beam was brought into contact with the sample
308 and then moved at a rate of 0.5 mm/min. Force as a function of displacement was measured along
309 the height (1 mm dimension) of the samples and calculated by the Microsquisher software using
310 the beam's stiffness, displacement, and length. Strains from 1 to 10% were used to calculate
311 Young's modulus, as the stress-strain response in this region was linear.

312 313 Optical images of living composites

315 Microscopic imaging was carried out using an Olympus FV3000RS confocal laser scanning
316 microscope. To visualize embedded yeast cell budding, living composites with 10 wt/v%
317 acrylamide and 0.1 wt/v% MBAA were synthesized mixing approximately 1x10⁶ cells/mL of the
318 pre-gel solution. Before polymerization, a 0.05 wt/v% aqueous solution of PolyFluor 570 was added
319 at 1% of the total pre-gel solution volume. Cells, after incubation for 48 h in YPD media, were
320 further stained by submerging samples in a 0.05 wt/v% aqueous solution of Trypan Blue for 3 min
321 and then washed two times in dH₂O. Budding of embedded cells and new colonies were observed
322 throughout the thickness and area of the imaged samples (n=3).

323 324 Macroscopic fluorescence images

326 For fluorescence imaging, living composites were dyed with a 0.05 wt/v% solution of
327 sulforhodamine B in water. By shining light at a wavelength of 455 nm, fluorescent images of the
328 UV patterned coatings and free-standing structures were obtained using a DSLR camera (Canon
329 Rebel T5i) fitted with a red filter (Hoya HMC R25A). This filter blocks light below 600 nm, thus
330 allowing visualization of the emitted light.

331 332 UV photopatterning of composites with active dried yeast

334 Living composites covalently bound to methacrylate-functionalized glass molds with 500 µm
335 thickness were prepared as described above. Composites were allowed to equilibrate in water before
336 UV exposure. A shadow mask of the word "ALIVE" was designed in AutoCAD and laser cut from

337 black polymer sheets. Irradiation with 254 nm UV light with an intensity of 2 mW/cm² was
338 performed from one side for 35 min using an UVP UVLink 1000 crosslinker chamber. Samples
339 were placed on a dark background during irradiation.

340
341 For UV patterning of free-standing composites, 12 mm diameter disks were cut from films 500 μm
342 thick and patterned to induce cell death in a 6 mm diameter inner circle or a 3 mm wide ring pattern,
343 using aluminum foil as a mask. Irradiation with 254 nm UV light with an intensity of 2 mW/cm²
344 was performed from one side for 35 min using the UV chamber.

345
346 For cylindrical helix patterning, living composite samples were synthesized with the same
347 composition of monomers as described above, and 0.9-1.1 billion cells/mL of pre-gel solution.
348 Samples were cut into rectangular shapes (40 mm length, 5 mm height and 0.5 mm thickness) and
349 patterned to induce cell death in 2.3 mm wide rectangles separated by 1.1 mm and positioned at 56°
350 angle along the length of the samples (Fig. 3A). Irradiation was performed with the same
351 wavelength, intensity, and time as described above.

352
353 After irradiation, living composites bound to glass were incubated for 36 h at 30 °C with a media
354 change every 12 h. Free-standing disks were incubated for 48 h at 30 °C with media change every
355 12 h. These samples were then imaged using a fluorescent dye as described above. Cylindrical helix
356 films were incubated for 48 h at room temperature with media change every 6 h and then imaged
357 using a MightyScope 5M digital microscope every 5 min.

358
359 Topography measurements of UV photopatterned living composite coatings

360
361 Topography (Fig. 1F) of living composite coatings was imaged using a digital microscope (Keyence
362 VHX-1000). To characterize the change in film thickness after cell proliferation, measurements
363 along the depth profile of the letter “A” were taken. The camera limit points were set by focusing
364 on the highest point of the grown letter and on the coated UV-killed surface. Between these limits,
365 pictures were taken at 100 \times magnification along the surface of the letter. Images were then stitched
366 using the Keyence software.

367
368 Quantifying shape change in living composites with auxotrophic yeast strain

369
370 Prior to composite synthesis, the auxotrophic yeast strain (CRY with empty vector, denoted as
371 negative control strain) was grown overnight in selective media (0.7% yeast nitrogen base w/o
372 amino acids, 2% D-glucose, and appropriate amino acid supplements (38) lacking tryptophan,
373 leucine and histidine) containing L-histidine. Subsequently, overgrowths were made in 50 mL of
374 YPAD (1% yeast extract, 2% peptone, 0.004% adenine sulfate, 2% D-glucose) media at 30°C for
375 15 h. Growth was followed by measuring optical density at 660 nm until desired yeast concentration
376 was reached ($\text{OD}_{660} = 1-1.1$). Cells were then centrifuged in 50 mL conical tubes and washed twice
377 in distilled water before encapsulation. Composites with encapsulated auxotrophic yeast were
378 synthesized by using 10 wt/v% acrylamide and 0.1 wt/v% MBAA with 0.9-1.1 billion cells/mL of
379 pre-gel solution, as described above. For volume change experiments, living composites were first
380 equilibrated in water for 24 h and then cut into 10 mm diameter disks with a thickness of 500 μm .
381 Disks were incubated in selective media lacking L-histidine with 10 mM 3AT (HIS3 gene
382 competitive inhibitor) for 48 h at 30°C, with a media change every 12 h. After this time, composites
383 were incubated in selective media containing L-histidine and 10mM 3AT for another 48 h, with a
384 media change every 12 h. Disks were measured after growth for quantification of volume change.

386 Identical disks were also exposed to selective media containing L-histidine, selective media
387 containing D-histidine, a stereoisomer of the natural amino acid L-Histidine, and selective media
388 without L-histidine. These, composites were incubated for 72 h at 30°C with a media change every
389 12 h. Data presented are an average of 3 samples per experiment.

390

391 Controlled blockage of microfluidic device

392

393 For shape change experiments using a microfluidic device, composites were cast into microfluidic
394 polymer micromolds. These molds were built with 75x51mm glass slides. One of the two slides
395 was functionalized with methacrylate groups by the processes described above. The other slide was
396 coated in Rain-X®. Slides were separated with one 250 μ m polystyrene spacer on each side and
397 BPA mixture with 1 wt% I-369 photoinitiator was pipetted into the mold. Using a Vivitek D912HD
398 (B9Creator) projector with the UV filter removed and the optics modified to decrease the focal
399 length, a positive mold of a microfluidic device with an inlet channel 600 μ m wide and four outlet
400 channels 400 μ m wide was polymerized onto the methacrylate-functionalized glass slide. To
401 remove all unpolymerized BPA, slides were cleaned by multiple washes between acetone and
402 isopropanol. This micromold was then used to create a cell for the polymerization of the living
403 composite. After polymerization, microfluidic devices were allowed to equilibrate in dH₂O for 24
404 h. Using UV-patterning (254 nm wavelength, 35 min, 2 mW/cm² intensity) most of the composite
405 was rendered inviable. Two of the four outlet channels, as indicated in Fig. 3F, were kept alive by
406 preventing UV exposure on the channel areas with a shadow mask. Selective media containing L-
407 histidine was then flowed at a rate of 34 μ L/min through one set of microfluidic devices (n=3) and
408 selective media without L-histidine was flowed through another set of devices (n=3) at the same
409 rate. Before and after media flow, channels were injected with a PEG-DA solution mixed with
410 sulforhodamine B aqueous solution at 1% of the total PEG-DA volume. Flowing this solution
411 allowed for visualization of the flow through the microchannels before and after growth.
412 Topographical quantification of these samples was performed following the same process described
413 for living composite coatings.

414

415 Yeast two-hybrid assays

416

417 L40 cells were co-transformed with the appropriate plasmids as described above and, in the
418 literature (39). Cells were plated on selective media agar plates with 10 mM 3AT. Two replicate
419 plates were grown in the dark and two were irradiated with blue light at 455 nm for 2 s every 2 min.
420 Plates were incubated at 30°C for 3 days

421 Transformants were grown to saturation in selective media lacking L-histidine overnight at 30°C.
422 Afterward, aliquots (100 μ L) of the saturated cultures were outgrown in 1 mL of fresh minimal
423 media and incubated at 30°C for 4 h. Optical density at 660 nm was recorded using a Spark® 20M
424 multimode reader. β -Galactosidase expression was quantified using the Beta-Glo® Assay System
425 (Promega). Briefly, 50 μ L of cell cultures were added to an equal volume of reagent and allowed
426 to incubate for 45 min prior to quantification. The luminescence values were normalized to cell
427 densities for each culture.

428

429 Optogenetic control of shape change

430

431 Control of proliferation by exposure to blue light is achieved by encapsulating experimental yeast
432 strains that express photosensory proteins. Pre-gel solutions with a composition of 10 wt/v%
433 acrylamide and 0.1 wt/v% crosslinker were mixed with 0.9-1.1 billion transformed cells/mL of

434 solution. Positive and negative control strains were encapsulated using the same pre-gel
435 composition and cell concentration. For volume change experiments, composites with each of the
436 three strains were cut into 10 mm diameter disks with 500 μm thickness. Samples were incubated
437 in selective media lacking L-histidine with 10 mM 3AT for 72 h with media change every 12 h.
438 Then, samples were irradiated with blue light (455 nm) (Fig. 4C) or kept in the dark. The intensity
439 of irradiation was measured with a solar power meter (Amprobe Solar-100) and set at 2.7 mW/cm².
440 For blue light experiments, samples were irradiated for 2 s every 2 min. Volume changes were
441 measured every 24 h for all samples before and after incubation. Data presented are an average of
442 3 samples per experiment.

443
444 For spatiotemporal control of proliferation, free-standing films with encapsulated experimental
445 yeast strain were UV patterned as shown in fig. S10. Films were kept in dH₂O for 24 h in the dark
446 before blue light exposure. Films were then incubated in selective media with 10 mM 3AT at 30°C
447 in the dark for 72 h. For the first 48 h, only the left half of the film is exposed to blue light to induce
448 proliferation of cells. After this time, the right half was irradiated for 24 h. Images are representative
449 from 3 trials.

450
451 **References and Notes**

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532

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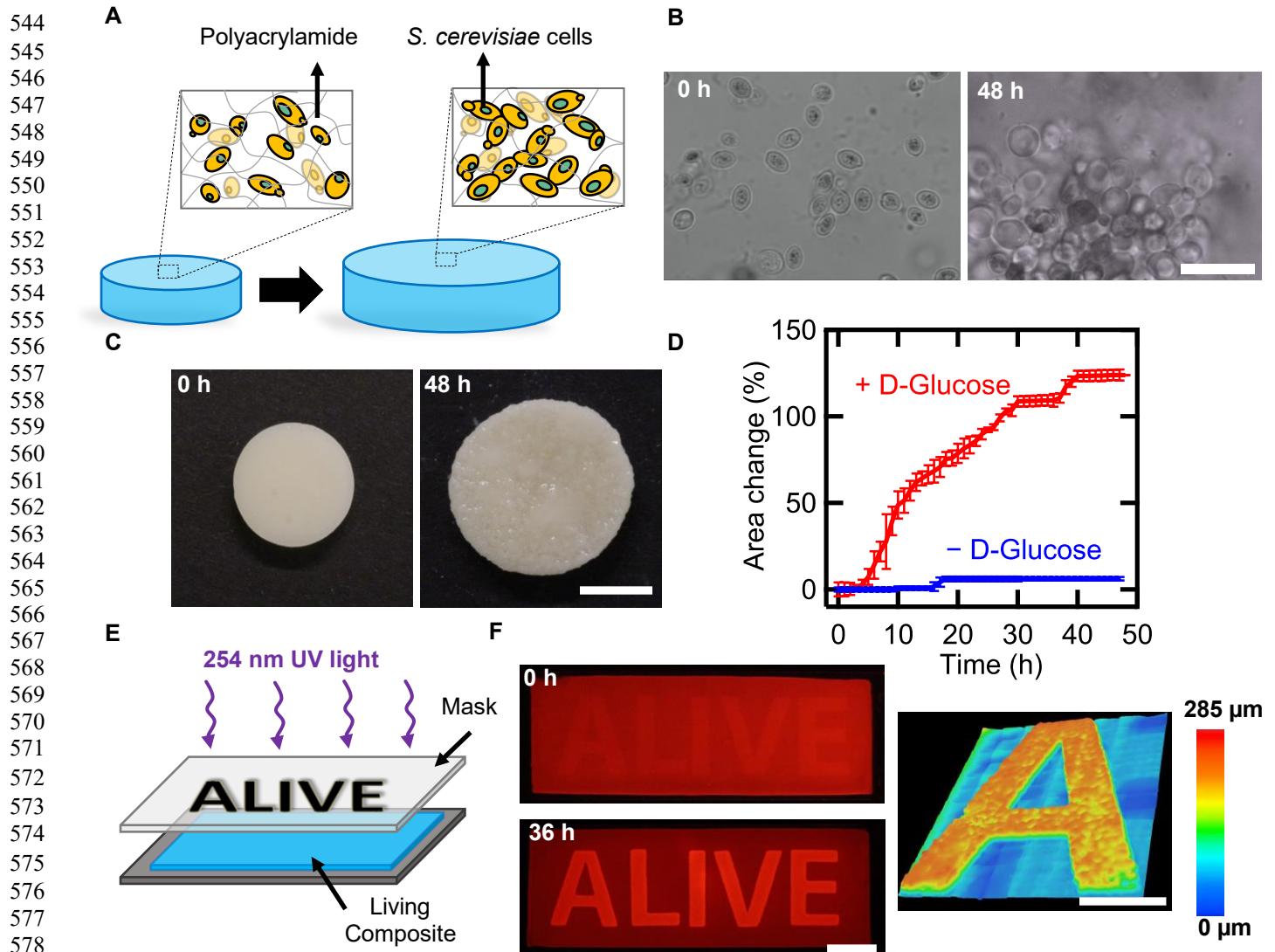


Fig. 1. Controlled expansion of polyacrylamide gels by proliferation of yeast. (A) Schematic of shape-change in living composites. In YPD, yeast proliferate and cause expansion in the polymer matrix. (B) Optical micrographs of a living composite before and after growth in media (Scale bar: 30 μ m). (C) Macroscopic expansion of a living composite gel with 6 wt% yeast (Scale bar: 7 mm). (D) Area change over time of a sample with 6 wt% yeast in the presence of media with and without glucose. (E) Photopatterning process of a living composite. (F) Fluorescence images of a living composite after UV patterning (top) and after incubation in YPD (bottom) (Scale bar: 10 mm). Topography of an initially flat living composite after exposure to YPD (right) (Scale bar: 5 mm). Each data point represents the mean ($n=3$), and error bars represent standard deviation. Trend lines are only intended to guide the eye.

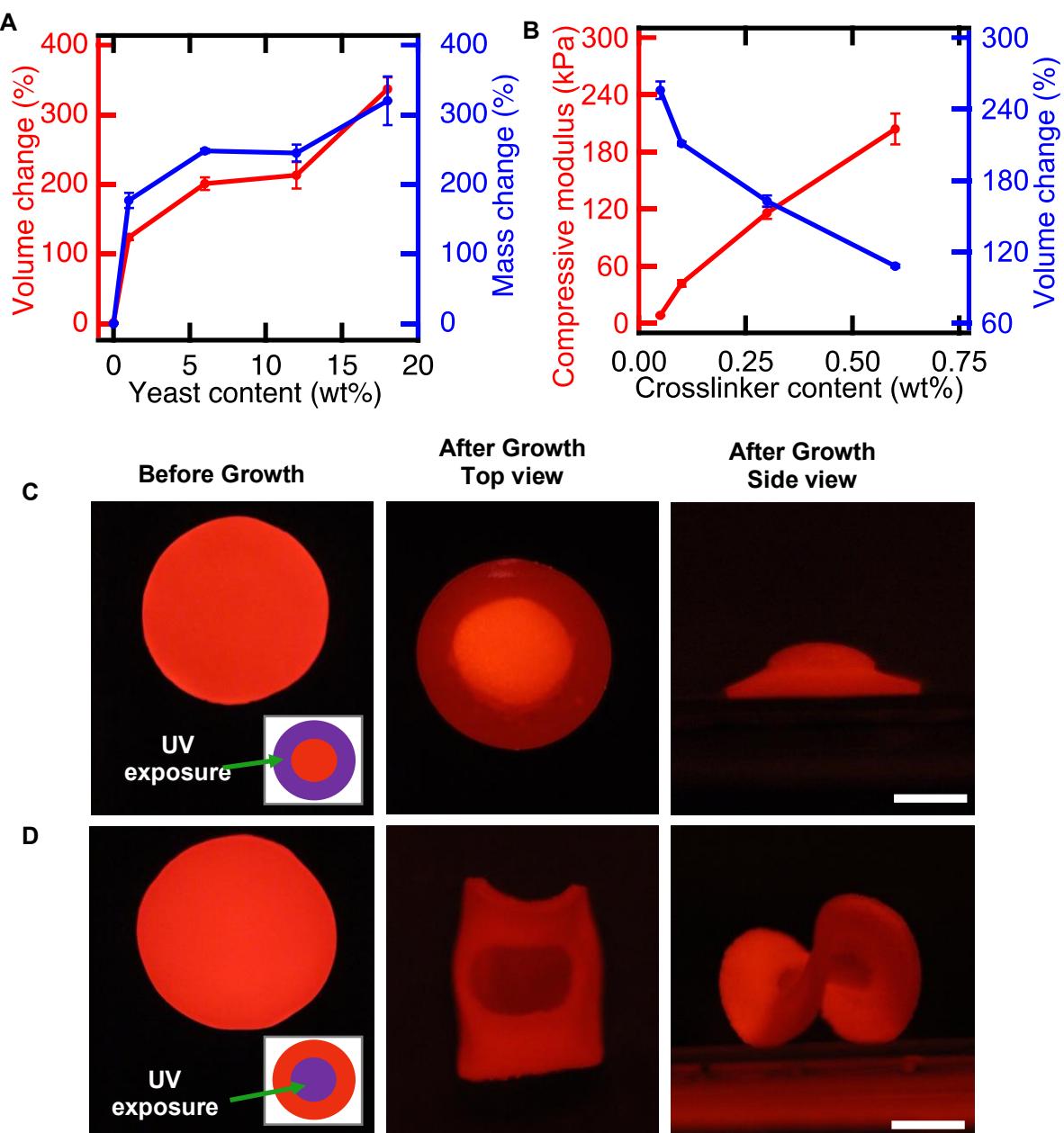
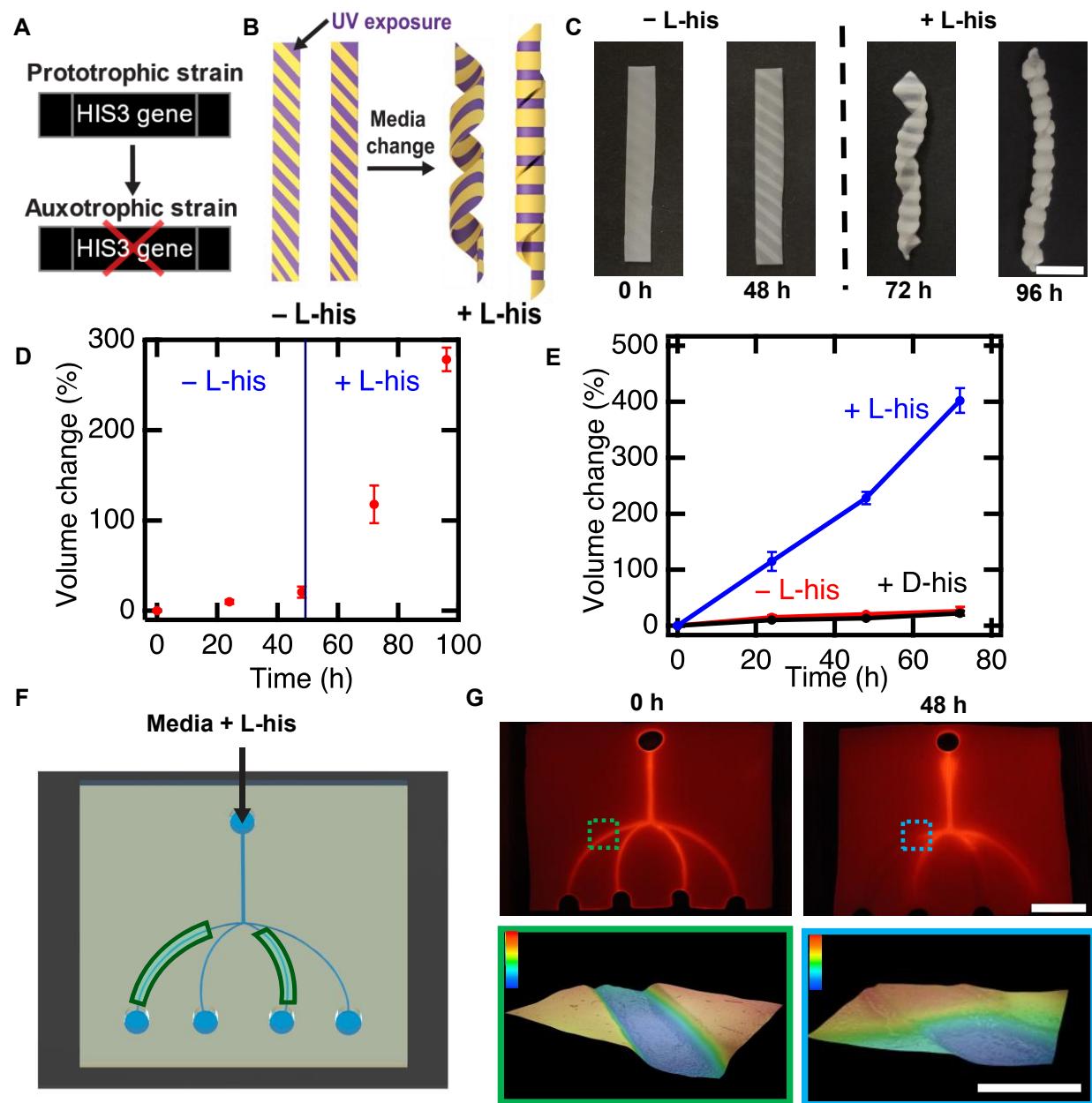
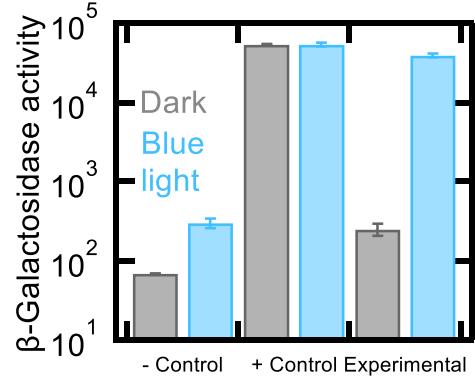
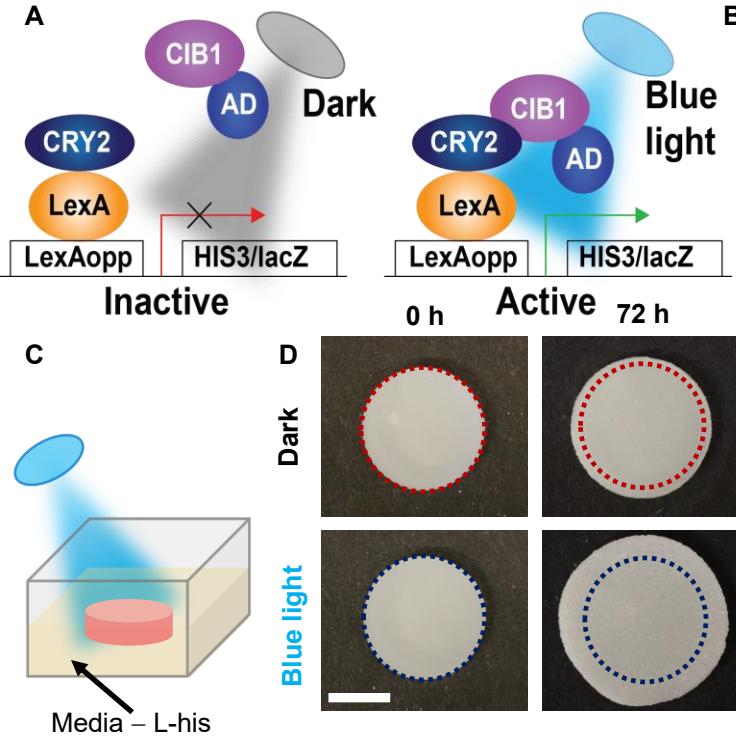


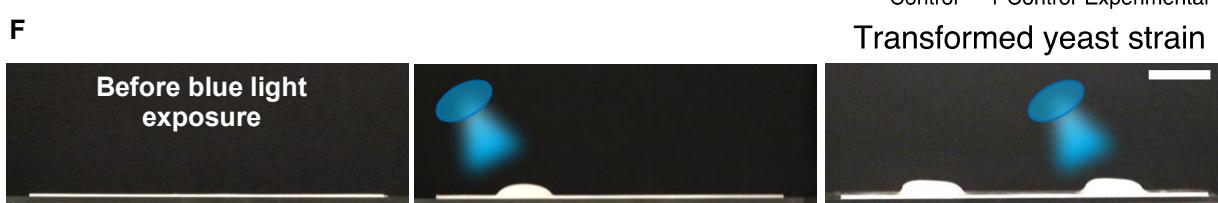
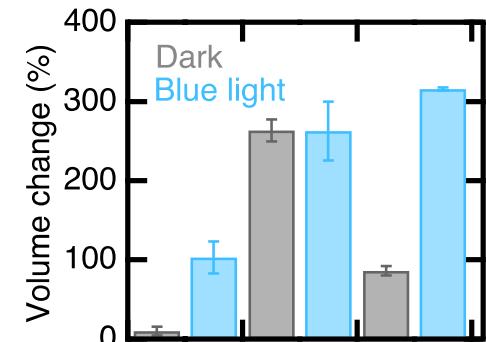
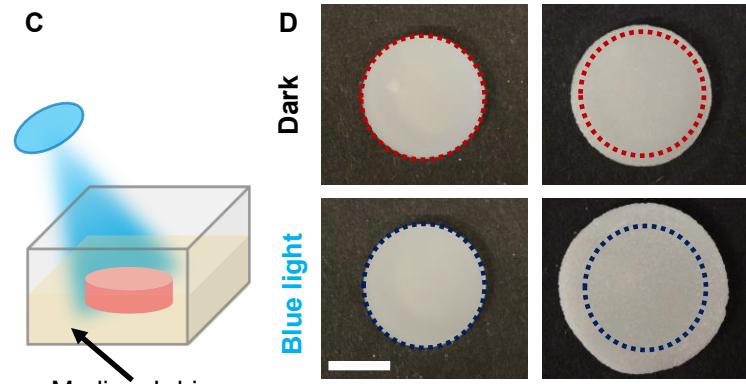
Fig. 2. Shape change of living composites can be controlled. (A) Volume and mass change of living composites as a function of yeast content. (B) Compressive modulus and volume change as a function of crosslinker content. (C) A flat disk exposed to spatially-pattered UV light (left) in a 3 mm wide ring pattern (inset). After incubation in media, a hat-like structure with positive Gaussian curvature is observed (center, right). (Scale bar: 5 mm). (D) A flat disk exposed to spatially-pattered UV light (left) in a 6 mm inner circle (inset). Upon incubation in media, a saddle-like structure with negative Gaussian curvature is observed (center, right). (Scale bar: 5 mm). Each data point represents the mean ($n=3$), and error bars represent standard deviation. Trend lines are only intended to guide the eye.



683 **Fig. 3. Genetic engineering enables controlled composite response to specific cues.** (A) Deletion of the HIS3 gene
684 results in failure to proliferate in media lacking histidine. (B) Schematic of a UV-patterned living composite with
685 growth triggered by the amino acid L-histidine. (C) UV-patterned living composites with auxotrophic yeast do not
686 substantially change in shape in media lacking L-histidine. Shape change into a helical structure after incubation in
687 media containing L-histidine (Scale bar: 10 mm) (D) Volume change over time for auxotrophic living composites
688 before and after L-histidine exposure. (E) Volume change over time for auxotrophic living composites incubated in
689 media lacking histidine, with D-histidine, or with L-histidine. (F) Schematic of a living microfluidic device where the
690 composites forming the channels indicated in green contain living auxotrophic yeast. (G) Fluorescence image of fluid
691 traversing the microfluidic device before exposure to media (top left) (Scale bar 10 mm). Fluorescence image of fluid
692 traversing the microfluidic device after media containing L-histidine flows for 48 h through the channels. Topography
693 of a living channel before and after (color scale: 0-0.3 mm) growth (bottom) (Scale bar: 1 mm). Each data point
694 represents the mean (n=3), and error bars represent standard deviation. Trend lines are only intended to guide the eye.
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Transformed yeast strain



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Fig. 4. Genetic engineering enables optogenetic control of shape change. (A) Schematic of a light sensitive yeast two-hybrid. Blue light induces expression of HIS3 and lacZ reporters by inducing conformational changes in CRY2 to favor interaction with CIB1. Reporter genes are transcribed by recruitment of the Gal4 activation domain (AD). (B) β -Galactosidase assays of an auxotrophic strain lacking CIB1 (negative control), a strain not auxotrophic for L-histidine in the dark (positive control), and the auxotrophic strain depicted in A (experimental). (C) Schematic of a living composite irradiated with blue light in growth media lacking L-histidine. (D) Volume change of living composites with experimental yeast irradiated with blue light or kept in the dark (Scale bars: 5 mm). (E) Volume change of living composites with each yeast strain when exposed to blue light or kept in the dark. (F) Patterned photoresponsive living composite with the experimental yeast strain in media lacking L-histidine where blue light is first targeted on the left side and then the right side (Scale bar: 10 mm). Each data point represents the mean (n=3), and error bars represent standard deviation.