

1 **Title: Multiplexed dynamic control of temperature to probe and observe mammalian cells.**

2

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13 **Abstract**

14 Temperature is a critical parameter for biological function, yet there is a lack of approaches to  
15 modulate the temperature of biological specimens in a dynamic and high-throughput manner.  
16 We present the thermoPlate, a device for programmable control of temperature in each well of a  
17 96-well plate, in a manner compatible with mammalian cell culture and live cell imaging. The  
18 thermoPlate maintains precise feedback control of temperature patterns independently in each  
19 well, with minutes-scale heating and cooling through  $\Delta T \sim 15-20^\circ\text{C}$ . A computational model that  
20 predicts thermal diffusion guides optimal design of heating protocols. The thermoPlate allowed  
21 systematic characterization of both synthetic and natural thermo-responsive systems. We first  
22 used the thermoPlate in conjunction with live-cell microscopy to characterize the rapid  
23 temperature-dependent phase separation of a synthetic elastin-like polypeptide (ELP<sub>53</sub>). We  
24 then measured stress granule (SG) formation in response to heat stress, observing differences  
25 in SG dynamics with each increasing degree of stress. We observed adaptive formation of SGs,  
26 whereby SGs formed but then dissolved in response to persistent heat stress ( $\geq 42^\circ\text{C}$ ). SG  
27 adaptation revealed a biochemical memory of stress that depended on both the time and  
28 temperature of heat shock. Stress memories continued to form even after the removal of heat  
29 and persisted for 6-9 hours before dissipating. The capabilities and open-source nature of the  
30 thermoPlate will empower the study and engineering of a wide range of thermoresponsive  
31 phenomena.

32 **Introduction**

33 Temperature plays important roles in the function and control of biological systems. In humans,  
34 core temperature is narrowly maintained around 37°C and, although temperature can rise  
35 during infection, the febrile response is also tightly controlled to remain below ~41°C<sup>1</sup>. In the  
36 biosciences, temperature can be harnessed as a non-invasive perturbation, for example with  
37 temperature-sensitive protein mutants that provide conditional knockdown of proteins<sup>2-4</sup>.  
38 Separately, heat-shock promoters can allow expression of custom transgenes in response to a  
39 brief increase in temperature. Such technologies can control gene expression in model

46 organisms<sup>5</sup> and, because heat effectively penetrates opaque tissues, also provide remote  
47 control of engineered cells within mammals<sup>6-8</sup>. Exposing mammalian cells to temperatures  
48 above ~42°C induces heat shock, which can be toxic over extended periods. This response has  
49 been harnessed for cancer therapy, where thermal ablation of tumors is a common, minimally-  
50 invasive treatment strategy<sup>9</sup>.

51  
52 Despite its importance, there are relatively few experimental methods to systematically  
53 manipulate temperature. Incubators can be set to desired temperatures, but changes in  
54 temperatures are slow, challenging studies of phenomena that occur over fast (~minutes)  
55 timescales. Incubators are also expensive, have a large footprint, and allow only one  
56 temperature to be tested at a time. Commercial devices such as the CherryTemp or VAHEAT  
57 are similarly limited in throughput. Thermocyclers can be used to specify multiple temperatures  
58 simultaneously<sup>10</sup>, but these are not compatible with long term cell survival or simultaneous  
59 imaging. The current lack of methods for microscope-compatible temperature modulation in  
60 microwell plates limits our understanding of biological responses to temperature changes and  
61 hinders the engineering of thermally-controllable biological systems.

62  
63 We address this gap with the thermoPlate, a device for programmable heating and thermometry  
64 of samples in 96-well plates. The thermoPlate allows heating of individual wells with arbitrary  
65 temperature profiles and is compatible with both live-cell microscopy and standard cell culture  
66 incubators. We characterize thermoPlate performance, finding accurate heating across the plate  
67 to < 0.1°C of the desired set point. We also present a model that predicts heat diffusion across  
68 the sample plate, allowing the user to rapidly test the feasibility of arbitrary desired heating  
69 programs *in silico*. We then harnessed the utility of the thermoPlate to characterize the rapid  
70 temperature-dependent phase-separation of an elastin-like polypeptide (ELP). Finally, we  
71 examined the dynamics of stress granule (SG) formation in response to graded levels of heat  
72 stress. We found that SGs form but spontaneously dissolve due to the formation of biochemical  
73 memories of stress, which depended on both the timing and intensity of heat stress.

74

## 75 **Results**

### 76 *thermoPlate Design, Calibration, and Dynamics*

77 The thermoPlate maintains a desired temperature by continuous deposition and  
78 measurement of heat in each well of a 96-well plate (**Figure 1A**). This is achieved with a  
79 custom-designed circuit board that accommodates a pair of thermistors in each of 96 positions  
80 in the format of a 96-well plate. A thermistor is a resistor whose resistance changes as a  
81 function of temperature, thus enabling its common use as a thermometer (“reader”). However,  
82 thermistors can also deposit heat when sufficient current is passed through them (“heater”). By  
83 pairing one reader and one heater thermistor in each well, the thermoPlate achieves  
84 independent measurement and heating of all 96 wells (**Figure 1A, B**). To operate, the user  
85 uploads the desired heating protocols to the on-board Arduino microcontroller. The Arduino  
86 regulates the current of each heater through one of 12 shift registers linked to 96 transistors,  
87 and it receives the temperature readings from each reader connected to a voltage divider  
88 through one of 6 multiplexers (**Figure S1**). With the set point and temperature of each well, the  
89 Arduino implements proportional-integral-derivative (PID) feedback control, adjusting the current

90 of each heater to reach and maintain the sample at the desired set point for every well (**Figure**  
91 **1C**. See **Methods** for more details on feedback control). The thermoPlate sits atop a 96-well  
92 culture plate to which it is mated with a 3D printed adapter (**Figure 1D**). The thermoPlate can  
93 operate in standard cell culture incubators. Alternatively, when the culture plate has a  
94 transparent bottom, this assembly is also compatible with live cell imaging using an inverted  
95 microscope. The thermoPlate can run in a standalone format but can also be connected to a  
96 computer to view real-time readouts of the current well temperatures.

97 With all components and 3D printed parts in hand, the thermoPlate can be assembled in  
98 8 hours, for a total cost of ~\$300. Design files, a parts list, assembly instructions, and operation  
99 protocols can be found in the **Methods** section and our online repository (link in **Methods**).

100 Multiplexed temperature control requires consistent, calibrated readings between each  
101 of the 96 readers. We characterized the variability in temperature readings in a fully-assembled  
102 thermoPlate in a cell culture incubator that was set to one of 5 temperatures between 26-45°C,  
103 with the true temperature verified by two digital thermometers. While the average error between  
104 thermoPlate and true temperature increased slightly at higher temperatures, we observed a  
105 consistent pattern of variability between wells (SD ~ 0.26°C) (**Figure 1E**). From these results,  
106 we derived adjustment factors for each well that corrected readings to within 0.1°C of the true  
107 temperature across all temperatures, with a standard deviation of 0.06°C between wells in  
108 subsequent experiments (**Figure 1E**). Once calculated, calibration values maintained accurate  
109 readings when tested after weeks of use (**Figure S2**).

110 We next tested the thermoPlate's ability to control prespecified temperatures, with  
111 temperature measured by the calibrated reader thermistors. 15 wells were set to simultaneously  
112 maintain temperatures ranging from 28°C to 42°C (3-17°C above ambient) in 1°C increments  
113 (**Figure 1F**). After a rapid approach and equilibration at the designated set point, each well  
114 maintained its temperature with high accuracy (average error of 0.04°C from set point) and  
115 precision (average SD of 0.05°C within individual trace) over 15 hrs of operation (**Figure 1F**).  
116 Moreover, temperatures could be dynamically toggled between multiple setpoints, returning to  
117 the desired temperature with high fidelity (at steady state: average error of 0.07°C, average  
118 within-trace SD = 0.06°C) (**Figure 1G**).

## 119 *Characterizing speed and accuracy of thermoPlate heating*

120 We next characterized key operational features of thermoPlate heating, including its  
121 range, accuracy, heating/cooling speed, and overshoot. We devised an experiment where the  
122 thermoPlate would sequentially heat each well from 3°C above ambient temperature to a  
123 prespecified higher temperature for 20 minutes, followed by cooling back to 3°C above ambient.  
124 Note that because the thermoPlate has no active cooling, in most experiments the baseline  
125 (lowest) experimental temperature should be set to higher than ambient to allow rapid cooling  
126 to—and maintenance of—this desired low temperature. **Figure 2A** demonstrates the above  
127 described experiment for heating through  $\Delta T = 10^\circ\text{C}$ . All wells experienced this heat pulse with  
128 high consistency between wells (**Figures 2A,B**). From this type of experiment, we could  
129 systematically quantify error (difference between the temperature and the set point after 20  
130 minutes), rise/fall time (the time until temperature reached 99% of the target temperature), and  
131 overshoot/undershoot (difference between the maximum/minimum and setpoint).

133 To quantify performance as a function of heating magnitude, we chose 11 wells  
134 distributed across the plate and repeated the above experiment with heating of either 5, 10, 15,  
135 or 20°C above ambient, with each well heated sequentially (**Figure 2C**). While all wells reached  
136 their target temperature for up to  $\Delta T = 15^\circ\text{C}$ , only some wells could reach  $\Delta T = 20^\circ\text{C}$  within 20  
137 minutes. The rate at which a well approached  $\Delta T = 20^\circ\text{C}$  was highly dependent on its position  
138 on the plate, with corner wells approaching 20°C more rapidly than center wells (**Figure S3**).  
139 This variability in heating speed likely arises because of differences in heat diffusion in different  
140 parts of the plate, with higher numbers of neighbor wells acting as a larger heatsink that limits  
141 heating. Rise/fall times were 2 min for  $\Delta T = 5^\circ\text{C}$  and 4 min for  $\Delta T = 10^\circ\text{C}$ , with increasing  
142 variability between wells with increasing  $\Delta T$  (**Figure 2D**). However, error (0.05-0.15°C) and  
143 overshoot/undershoot (0.1-0.3°C) were minimal and did not correlate with  $\Delta T$ .

144 Although heat diffusion limited the heating range of individual wells, it could also be  
145 harnessed to boost heating within groups of wells. We tested this concept by heating either  
146 individual wells or groups of 4 or 9 wells to  $\Delta T = 20^\circ\text{C}$ . Although, as before, individual wells  
147 struggled to reach this level of heating, wells in a 3x3 matrix rapidly equilibrated to this high set  
148 point (**Figure 2E**). Grouped wells achieved a faster approach to high temperatures at the  
149 expense of small increases in error (0.15-0.2°C) and overshoot (~1°C) (**Figure 2F**). In sum, the  
150 thermoPlate can reliably and rapidly achieve heating of up to  $\Delta T = 15^\circ\text{C}$  in individual wells and  
151 can also be programmed to achieve  $\Delta T > 20^\circ\text{C}$  with appropriate arrangement of the heated  
152 wells.

153

#### 154 *Modeling heat diffusion in the thermoPlate*

155 Heat diffusion constrains the heating protocols that can be run simultaneously because it  
156 places limits on the temperature differential between two neighboring wells. For example, a well  
157 heated to 15°C above ambient will diffusively heat its neighboring wells by ~10°C, placing a  
158 lower limit on the possible temperatures of those neighbors. As the number of wells in use  
159 increases, it becomes increasingly challenging to predict whether a desired protocol is  
160 achievable, particularly when dynamic heating protocols are desired.

161 We thus created a model of heat diffusion in 96-well plates to allow users to rapidly  
162 assess the feasibility of particular heating programs. We implemented an explicit finite  
163 difference approximation of a modified heat equation. This model simulates dynamic heat  
164 stimuli and calculates the rate of heat spread through each well of the plate over time. Unlike  
165 the classical heat equation, our model has 6 (vs 1) parameters that describe the rate of heat  
166 spread. These additional parameters account for observed differences in heat spread across  
167 different contexts, for example in edge wells vs center wells (**Figure 3A**). These parameters  
168 also accounted for small asymmetries around heated wells (left vs right, top vs bottom) likely  
169 due to asymmetric positioning of the electrical components.

170 To fit the model, we generated eight sets of training data where different sets of wells (1-  
171 12 wells) underwent dynamic heating/cooling programs, with temperature changes of 1-15°C  
172 over durations of up to 90 min (**Figure 3B**). These training experiments were designed with  
173 increasing levels of complexity, from simple heating of one well (Dataset 1) to more complex,  
174 dynamic heating of 12 wells arranged both individually and in groups (Dataset 8) (**Figure S4**).  
175 The model was parameterized for each training set by minimizing the root mean squared error  
176 (RMSE). These 8 parameter sets were then averaged to create a final set of parameters that

177 describe thermoPlate behavior. We validated the model using Leave-One-Out-Cross-Validation,  
178 wherein fitted parameters from 7 training sets were averaged and used to predict the 8th  
179 dataset, resulting in a mean average error of 0.23°C across all wells and times. Comparing the  
180 predictions of the full model vs the data across all data sets, the average error over an  
181 experimental run was 0.22°C (**Figure 3C-E**), with error falling within 1°C for > 99% of timepoints  
182 and within 0.5°C for > 75% of timepoints (**Figure 3E**), showing close agreement between model  
183 and data. Transient spikes in error (1-2°C) could on occasion be observed near wells that  
184 experienced a sudden large temperature change (**Figure 3D**).

185 The predictive power of this model allows users to rapidly test whether particular heating  
186 patterns and spatial arrangements are permissible. For this purpose, we created an online  
187 interface where users can upload the desired heating protocol and easily simulate and visualize  
188 heat generation and spread (**Figure 3F**). The visualization also highlights impermissible  
189 temperature set points, allowing the user to modify their experimental design accordingly.  
190

#### 191 *Controlling and characterizing thermogenetic phase separation*

192 We applied the thermoPlate to probe and observe thermally-controlled biochemistry in  
193 mammalian cells. We first confirmed that the thermoPlate was fully compatible with mammalian  
194 cell culture, with no observed impact on proliferation or survival of HEK 293T cells after even 48  
195 hr of direct contact with the thermoPlate (**Figure S5**). Measurement of evaporation revealed  
196 that, while significant evaporation could be observed in edge wells under prolonged heating (24  
197 hr at >10°C), evaporation could be prevented by adding a thin layer of paraffin oil over each well  
198 (**Figure S6**).

199 We first leveraged rapid and dynamic thermal control to characterize phase separation  
200 of an elastin-like polypeptide (ELP<sub>53</sub>), which forms protein condensates in mammalian cells  
201 above a critical temperature (**Figure 4A,B**)<sup>11</sup>. ELP condensates form rapidly (~minutes) and  
202 over a small temperature range (~1-2°C)<sup>12,13</sup>. We used the thermoPlate coupled with live-cell  
203 confocal microscopy to observe ELP<sub>53</sub> condensation in response to increased temperature  
204 between 28-39°C, at 1°C resolution (**Figure 4C**). ELPs condensed rapidly and as a function of  
205 temperature, with measurable condensation above 31°C and a ~linear increase in magnitude  
206 between 33-38°C (**Figure 4C,D**). We next quantified the speed of ELP formation kinetics,  
207 finding that ELP condensation increased within 1 min of heating and reached steady-state within  
208 10 min (**Figure 4E**). Notably, the thermoPlate achieved its target temperatures in <2 minutes,  
209 sufficiently fast to observe the slightly slower ELP condensation. The fast and dynamic  
210 responses of both the thermoPlate and ELPs allowed us to reversibly toggle ELP phase  
211 separation over 45 cycles of heating and cooling (4 min 35°C, 4 min 29°C). ELP condensation  
212 and dissolution remained consistent over all cycles, with no decrease in dynamic range,  
213 suggesting that such stimulation could be maintained indefinitely (**Figure 4F, Movie S1**).  
214

#### 215 *Characterizing dynamics and memory of heat stress in mammalian cells*

216 We next applied the thermoPlate to examine native thermal responses of mammalian  
217 cells. We focused on the dynamics of stress granule formation in response to heat shock.  
218 Stress granules (SGs) are condensates of protein and RNA that form in the cytosol in response  
219 to stressors including heat, hypoxia, and oxidation<sup>14</sup>. While the composition and function of SGs  
220 in response to heat stress have been studied extensively<sup>15-20</sup>, their dynamics are relatively less

221 explored. To observe stress granules, we stably expressed a fluorescent fusion of the SG  
222 scaffold G3BP1-mCherry in HeLa cells, and we imaged G3BP1 condensation as a function of  
223 time and temperature.

224 We first asked at what temperature SGs could be observed by exposing cells to constant  
225 heat between 41 to 47°C (**Figure 5A**). We found a highly non-linear relationship between  
226 temperature and the magnitude of stress-granule formation. Although no SGs were observed at  
227 41°C, SG rapidly formed at 42°C and increased in magnitude at 43°C and 44°C. Surprisingly,  
228 maximal SG intensity decreased with progressively further increases in temperature (**Figure**  
229 **5B-D**). Although cells remained morphologically unperturbed with SGs at 42°C and 43°C over  
230 at least 8 hrs, cells heated with 44°C lost cytoplasmic volume and became immobile, with higher  
231 temperatures leading to faster onset of this phenotype (**Figure 5C, Movie S2**). In all heating  
232 conditions, SGs formed rapidly within the first 5 min.

233 Notably, SG formation was transient across all conditions despite constant heating, and  
234 in most cells G3BP1 returned to a diffuse state within several hours of the onset of heat shock  
235 (**Figure 5B**). We asked if SG adaptation required constant heat stress or alternatively could  
236 form in response to short heat pulses. We examined SG formation in response to periodic 1-hr  
237 pulses of heat stress (43.5°C), a pulse duration shorter than the time scale of SG adaptation  
238 (**Figure 5B**). Although the SGs formed and dissolved in sync with the first two heating and  
239 cooling pulses, SGs largely did not form during the 3<sup>rd</sup> and subsequent pulses, suggesting that  
240 cells continued adapting to heat stress despite heat not being applied for the entire duration of  
241 adaptation (**Figure 5E,F, Movie S3**). Indeed, even though the second heat pulse yielded strong  
242 SG formation, SG magnitude began to decrease during this second window, indicating that the  
243 first heat pulse influenced SG behavior during the second heat pulse (**Figure 5E**). Collectively,  
244 these results show that cells store memories of prior heat exposure that shape future responses  
245 to repeated stress.

246 We next characterized the timescales of this memory formation. We programmed the  
247 thermoPlate to apply two pulses of heat stress (43.5°C) separated by varying periods of  
248 relaxation at 37°C (**Figure 5G**), and we quantified SG formation during the second pulse. SG  
249 formation depended on the interval between pulses in a biphasic manner. Although SGs were  
250 prominent during the second pulse after a 1 or 2 hr interval, they were not observed with  
251 intervals of 3 and 4.5 hrs. At still longer intervals, SG formation began to reappear on the  
252 second pulse. With a 9 hr interval, SG formation during the 2nd pulse resembled that of the first  
253 pulse (**Figure 5G,H**). Thus, cellular memory of prior stress was maximal for 3-6 hrs after which  
254 the memory began to fade and cells again responded to heat shock as on the first exposure.

255 Finally, we wondered whether memories of prior heat shock depended on the duration  
256 or the temperature of heat shock, or both. We designed a “pump-probe experiment” where we  
257 delivered a “pump” of heat stress with variable duration (15, 30, 60 min) and temperature (43-  
258 44.5°C), returned to 37°C for 3 hr, and then “probed” SG formation with a subsequent 1 hr of  
259 heat shock (43.5°C) (**Figure 5I**). Indeed, memory (suppression of SG formation on the second  
260 pulse) depended on both time and temperature of the initial heat shock pump. Suppression of  
261 SGs was stronger with increasing temperatures, and the magnitude of suppression increased  
262 with increasing pump duration. Notably, even a short 15 min pump was sufficient for  
263 measurable SG suppression at higher temperatures (**Figure 5I,J**). Additionally, the rate of  
264 dissolution of SGs after removal of the pump pulse depended on the intensity and duration of

265 heat stress, with slower dissociation in response to longer and more intense heating (**Figure 5I**),  
266 indicating a second form of stress memory. This effect could only be observed at or above 44°C  
267 heating and could be observed even after a brief 15 min pump duration (**Figure 5K**). In sum, our  
268 results indicate multiple mechanisms by which cells can store memories of prior heat stress and  
269 demonstrate the utility of the thermoPlate to explore these stress responses efficiently with high  
270 precision, throughput, and temporal resolution.

271

## 272 Discussion

273 The thermoPlate allows modulation of the temperature of biological samples in a  
274 multiplexed and small format that is compatible with both long term cell culture and live-cell  
275 microscopy. After calibration, the thermoPlate achieved its desired set point temperatures  
276 rapidly and accurately, with minimal variation between wells, allowing profiling of the response  
277 to thermal dynamics with high resolution in temperature and in time. The device is open source  
278 and can be assembled by non-specialists in several hours by following the assembly protocols  
279 associated with this manuscript, for a total cost of ~\$300 in parts. The design files can also be  
280 modified to adapt the device for user-specific applications as needed, for example for use in  
281 larger well-plate formats.

282 Heat diffusion constrains the types of heating protocols that can be run simultaneously.  
283 Nevertheless, these constraints need not limit throughput. For example, if a sustained  
284 temperature sweep is desired, a temperature gradient can be specified across the plate, and all  
285 96-wells can be used simultaneously. For more complex and dynamic patterns, samples must  
286 be placed more strategically to allow the samples to heat and cool appropriately. While this is  
287 straightforward for a few samples, it becomes more challenging with each additional heated  
288 well. For this reason, we developed a model of thermal diffusion that accurately predicts heating  
289 dynamics of each well for arbitrary protocols. The user can rapidly simulate a desired protocol  
290 *in silico* and observe whether the intended thermal patterns are permitted within the constraints  
291 of thermal diffusion. If they are not, the user adjusts the inputs, simulates, and iterates until a  
292 suitable arrangement is reached. We also introduce a web-based GUI for easy access to this  
293 model.

294 The thermoPlate also harnesses thermal diffusion to passively cool the samples. The  
295 simplicity of passive cooling is balanced by two additional constraints: 1) potentially slow cooling  
296 kinetics and 2) inability to maintain certain wells at ambient temperatures while heating other  
297 wells, since over time all wells will increase temperature by a few degrees as heat spreads  
298 throughout the plate faster than it diffuses out to the environment. To address both concerns,  
299 we maintained the environment at least ~3°C cooler than the lowest experimental temperature.  
300 This ensures that low experimental temperatures can be maintained and also results in faster  
301 cooling to this low level. With this approach, heating and cooling occur on comparable ~minutes  
302 timescales (**Figure 2D**).

303 The thermoPlate enables rapid characterization of new thermogenetic tools, where  
304 temperature acts as an inducer for genetic or protein activity<sup>6–8</sup>. We demonstrated this ability by  
305 systematically measuring the steady-state, kinetics, and dynamics of ELP<sub>53</sub>, an elastin-like  
306 peptide that forms condensates above a critical temperature. Although ELP phase separation is  
307 rapid, the fast heating of the thermoPlate was able to capture rapid ELP condensation and its  
308 slower approach to equilibrium, showing kinetics that are consistent with prior reports of ELP

309 condensation in cells<sup>11,12</sup>. ELP condensation could also be toggled indefinitely with fast minutes-  
310 scale ON- and OFF-kinetics. Such rapid clustering matches the timescales of even the fastest  
311 optogenetic clustering approaches<sup>21</sup>. In a separate report, the thermoPlate was used to  
312 systematically characterize the thermogenetic Melt protein, further demonstrating the potential  
313 of the thermoPlate for thorough characterization of temperature-sensitive proteins<sup>22</sup>.

314 The thermoPlate also empowers studies of native cellular responses to temperature,  
315 which we demonstrated by exploring the dynamics of the stress granule (SG) formation in  
316 response to time-varying heat-shock temperatures. We found a rich set of dynamic features of  
317 SG formation that respond to both the time and temperature of heat stress in a non-linear  
318 manner. Rather than a binary response, SG formation differed with each additional degree of  
319 heating above 41°C, with a maximal peak amplitude at 44°C. Notably, SGs showed adaptation  
320 to heat stress, such that G3BP1 puncta spontaneously dissolved despite persistent heating.  
321 Adaptation of SGs has been previously reported in the context of oxidative stress<sup>23</sup> and viral  
322 infection<sup>24</sup>. Such adaptation is consistent with negative feedback, including through transcription  
323 of heat shock proteins (HSPs)<sup>25-29</sup> and of suppressors of the integrated stress response  
324 including GADD34<sup>24,30,31</sup>. Nevertheless, to our knowledge the adaptation of SGs to chronic heat  
325 stress has not been previously reported, demonstrating the utility of simultaneous thermal  
326 control and live-cell imaging.

327 Heat stress resulted in the formation of at least two types of biochemical memories. A 1  
328 hr pulse of stress was sufficient to suppress SG formation upon subsequent stress, with  
329 maximal suppression observed 3-6 hrs after the initial stress pulse. These kinetics are in line  
330 with a recent finding that 1 hr of heat stress induced suppressed activation of the integrated  
331 stress response after a second heat pulse<sup>24</sup>. However, this prior work did not observe the status  
332 of SG formation during this second pulse. Notably, our measurements also revealed SG  
333 suppression after only 15 and 30 min of initial heat shock. With these shorter durations, the  
334 strength of SG suppression increased with increasing temperature of prior heat shock. A second  
335 form of biochemical memory appeared in the kinetics of SG dissolution after stress removal.  
336 This memory was observed immediately after removal of heat stress but only above 44°C,  
337 consistent with fast post-translational mechanisms like protein clustering distinct from SG  
338 formation<sup>32</sup>. While further work will be required to identify the specific mechanisms that underlie  
339 our observations, our results highlight the discovery potential of instrumentation for easy,  
340 precise, and dynamic control of temperature under a microscope.

341

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348

349

#### 350 **Author Contributions:**

351 W.B. and L.J.B conceived study. W.B. P.I., L.J.B. designed hardware structure and component  
352 layout. W.B. and P.I. constructed hardware, performed experiments, and analyzed data. T.M.

353 and P.I. developed the model of thermal diffusion. Z.H. generated the ELP<sub>53</sub> construct. L.J.B.  
354 supervised the work. W.B., T.M., and L.J.B. drafted manuscript and made figures. All authors  
355 contributed to editing.  
356

357 **METHODS**

358 *PID Implementation*

359 PID (Proportional, Integral, Derivative) control was implemented using a discrete form of the  
360 standard PID algorithm. PID constants were generated empirically by testing several  
361 combinations. In brief, the Arduino collects the temperature of every well every ~2 seconds.  
362 These values are used to calculate the current error (proportional), change in error from the  
363 previous reading per unit time (derivative), and integral (time weighted sum of all past error).  
364 The algorithm applies the PID constants to each error value and calculates a new duty ratio to  
365 apply to the corresponding heater. For more detailed information on programming architecture,  
366 see the **Code** section of the **Repository**.

367 *thermoPlate Calibration*

368 Calibration was performed by placing a thermoPlate on a 96 well plate containing PBS. This  
369 assembly was equilibrated in a standard mammalian cell culture incubator at ~25°C. Two digital  
370 thermometers (RC-4 Elitech Digital Temperature Data Logger) were placed inside two 15mL  
371 conical tubes filled with water and sealed with parafilm (to prevent evaporation). Both conical  
372 tube/thermometer were placed next to the thermoPlate, and a plastic box was placed over the  
373 thermoPlate and thermometers to avoid variation due to airflow. The average of the two digital  
374 thermometers was recorded as the true temperature. 5 temperature readings were recorded  
375 from each well. The average of those 5 readings was assigned as the final temperature reading  
376 of that well. The incubator temperature was then increased to 30°C and allowed to equilibrate  
377 for 2 hours. Readings from the thermoPlate and the digital thermometers were recorded, and  
378 this procedure was repeated in 5°C steps until 45°C. For each well, the temperature readings of  
379 the thermoPlate vs digital thermometers was then plotted, and the slope and y-intercept of the  
380 linear fit were extracted. These values were used to calibrate each raw temperature reading  
381 during thermoPlate operation. See **Manual** for more details on thermoPlate operation.

382 *thermoPlate Modeling*

383 A model of heat transfer within a thermoPlate-heated 96-well plate was made in MATLAB to  
384 simulate the temperature of each well given a desired heating input. Well temperatures were  
385 simulated through time. At each timestep, the instantaneous heat derivative of each well was  
386 calculated and multiplied by the simulation timestep to obtain the well's temperature at the next  
387 timestep. The model uses 8 parameters.  $H_p$  describes heat transfer between neighboring  
388 unheated wells. Around heated wells,  $H_{left}$ ,  $H_{right}$ ,  $H_{top}$ , and  $H_{bottom}$  describe heat transfer  
389 between the heated well and its neighbors to the left, right, top, and bottom, respectively. This  
390 was done to capture the consistent asymmetries seen in temperature around wells that are  
391 actively heated.  $H_e$  describes heat transfer from an edge well to the environment, **loss**  
392 describes a consistent loss of heat from each well to the environment. Finally, **heat** is a constant  
393 temperature increment that is added to each heated well during a simulation step where the well  
394 is below its set temperature. When a heated well is at or above its setpoint, this **heat** is not  
395 added for that step. Example calculations of the change in temperature of wells over a single  
396 time step are shown in **Figure S7**. Parameters were obtained by fitting the model to eight real  
397 datasets and averaging resultant 8 parameter sets to obtain generalizable parameters. Fitting  
398 was done by minimizing an error function, defined as the difference between the simulation and

399 the real data summed across every well and timepoint. Minimization was done using the  
400 MATLAB function fminsearch.

401 *Cell Culture*

402 HeLa cells were maintained in 10% fetal bovine serum (FBS) and 1% penicillin/streptomycin  
403 (P/S) in DMEM. Cell lines were not verified after purchase. Cells were not cultured in proximity  
404 to commonly misidentified cell lines.

405 *Plasmid design and assembly*

406 Constructs for stable transduction and transient transfection were cloned into the pHR lentiviral  
407 backbone with an SFFV promoter driving the gene of interest. The pHR backbone was  
408 linearized using MluI and NotI restriction sites. G3BP1 (Ophir Shalem Lab) and mCherry inserts  
409 were generated via PCR and cloned into the linearized viral backbone as a single fusion protein  
410 using HiFi cloning mix (NEB). H2B-iRFP nuclear marker was (Addgene Plasmid #90237). ELP<sub>48</sub>  
411 was obtained from Addgene (Addgene Plasmid #68395). Sequencing of ELP<sub>48</sub> showed it  
412 contained 44 ELP repeats. ELP<sub>48</sub> was digested using NdeI and BamHI and ligated into a small  
413 backbone pTA for the ease of cloning. ELP<sub>9</sub> was ordered as primers and inserted via HiFi into  
414 the pTA-ELP<sub>48</sub> linearized with AgeI and BsmBI, generating pTA-ELP<sub>53</sub>. EGFP was then  
415 amplified and inserted via HiFi to pTA-ELP<sub>53</sub>, which was linearized using MluI and XcmI. ELP<sub>53</sub>-  
416 EGFP was then subcloned via HiFi to the viral backbone pHR digested with MluI and NotI.

417 *Plasmid transfection.*

418 HEK 293T cells were transfected using the calcium phosphate method, as follows: Per 1 mL of  
419 media of the cell culture to be transfected, 50  $\mu$ L of 2x HeBS buffer, 1  $\mu$ g of each DNA  
420 construct, and H<sub>2</sub>O up to 94  $\mu$ L was mixed. 6  $\mu$ L of 2.5mM CaCl<sub>2</sub> was added after mixing of  
421 initial components, incubated for 1:45 minutes at room temperature, and added directly to cell  
422 culture.

423 *Lentiviral packaging and cell line generation*

424 Lentivirus was packaged by cotransfected the pHR transfer vector, pCMV-dR8.91 (Addgene,  
425 catalog number 12263), and pMD2.G (Addgene, catalog number 12259) into Lenti-X HEK293T.  
426 Briefly, cells were seeded one day prior to transfection at a concentration of 350,000 cells/mL in  
427 a 6-well plate. Plasmids were transfected using the calcium phosphate method. Media was  
428 removed one day post-transfection and replaced with fresh media. Two days post-transfection,  
429 media containing virus was collected and centrifuged at 800 x g for 3 minutes. The supernatant  
430 was passed through a 0.45  $\mu$ m filter. 500  $\mu$ L of filtered virus solution was added to 700,000  
431 HeLa cells seeded in a 6-well plate. Cells were expanded over multiple passages, and  
432 successfully transduced cells were enriched through fluorescence activated cell sorting (BD  
433 FACS Aria Fusion).

434 *Preparation of cells for plate-based experiments*

435 All experiments were carried out in Falcon™ 353219 Black 96 well plates. Briefly, wells were  
436 coated with 50  $\mu$ L of MilliporeSigma™ Chemicon™ Human Plasma Fibronectin Purified Protein  
437 fibronectin solution diluted 100x in PBS and were incubated at 37 °C for 30 min. HeLa or HEK  
438 293T cells were seeded in wells at a density of 60,000 cells/well in 100  $\mu$ L and were spun down

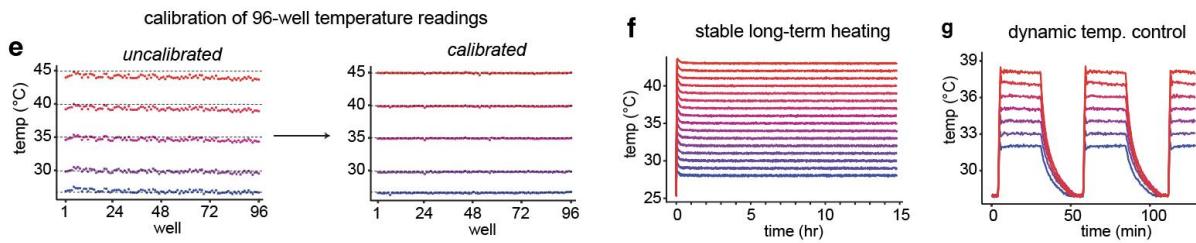
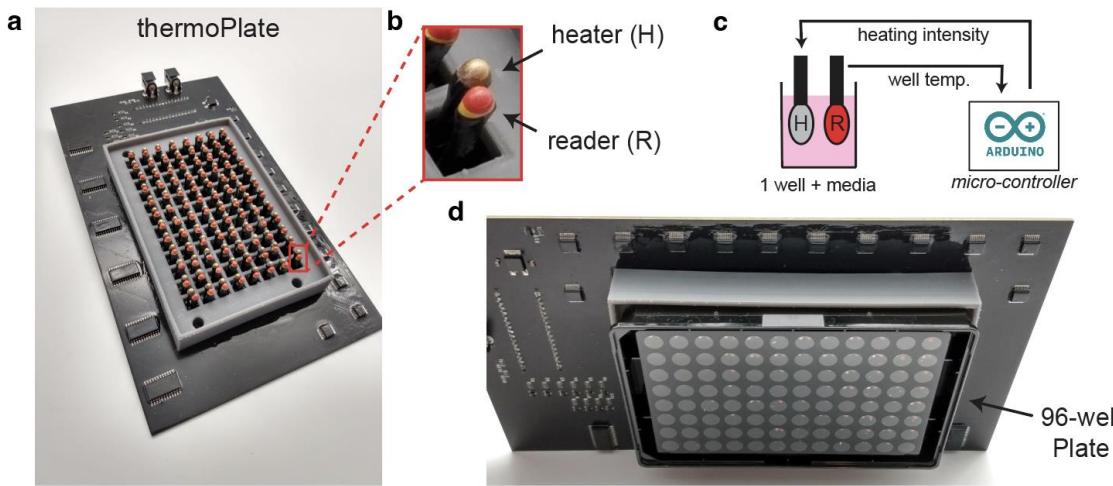
439 at 20 x g for 1 minute to ensure an even distribution of cells across the wells. For imaging ELPs,  
440 HEK 293T cells were transfected with GFP-ELP<sub>53</sub> and H2B-iRFP 12-24 hours post-seeding in  
441 96 well plates using the calcium phosphate method. Transfection mix/media was removed 24  
442 hours post transfection and replaced with fresh media. Cells were imaged within 4-8 hours of  
443 this media change.

444 *Imaging*

445 Live-cell imaging. Live-cell imaging was performed using a Nikon Ti2-E microscope equipped  
446 with a Yokagawa CSU-W1 spinning disk, 405/488/561/640 nm laser lines, an sCMOS camera  
447 (Photometrics), a motorized stage, and an environmental chamber (Okolabs). Cells expressing  
448 the construct of interest were imaged with a 20X air objective at variable temperatures and 5%  
449 CO<sub>2</sub>.

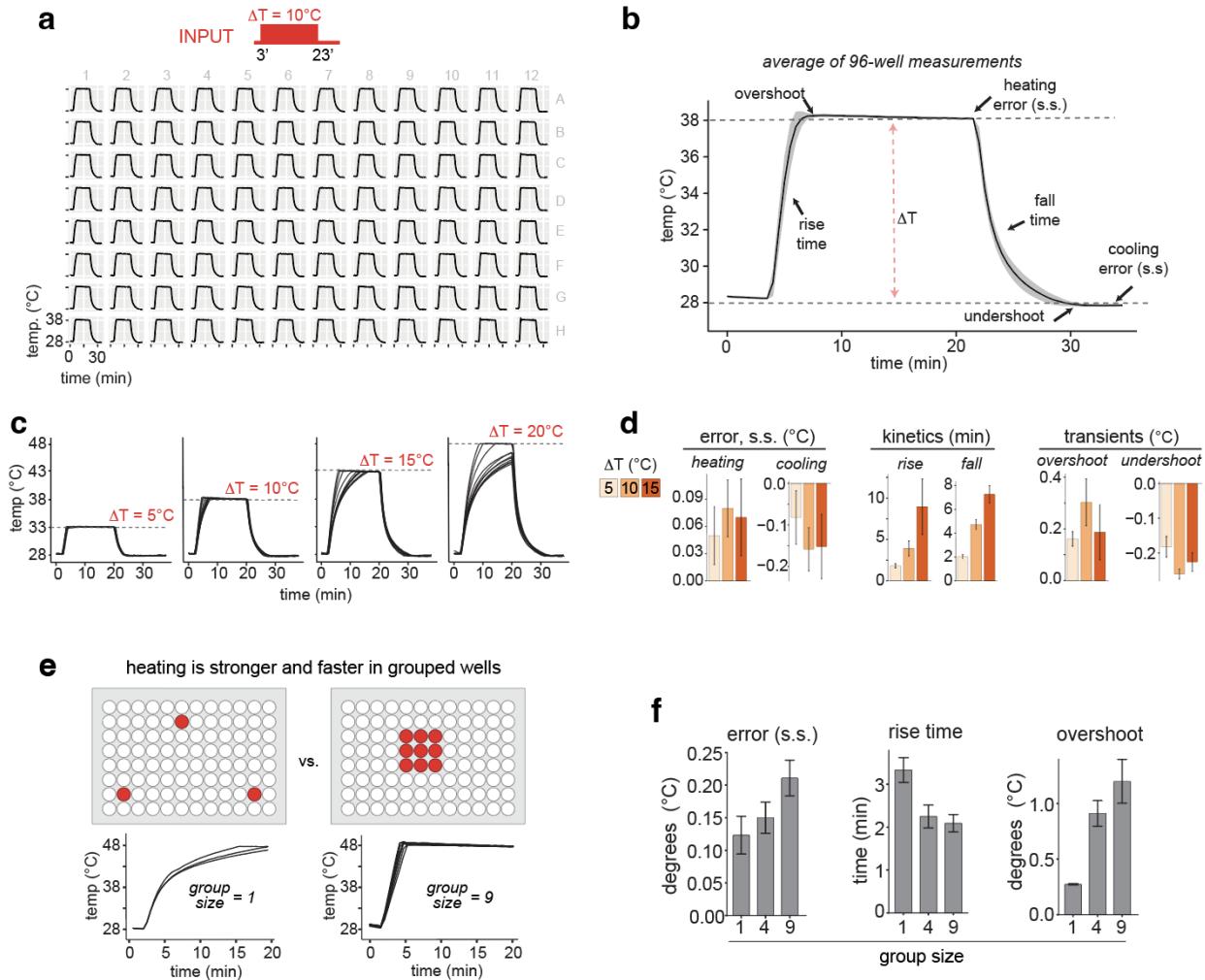
450 Imaging quantification. Images were processed using Cell Profiler <sup>33</sup>. Cells were segmented  
451 using the H2B-iRFP imaging channel, and cytoplasm was identified using a 20 pixel ring around  
452 the nucleus. Nuclear and cytoplasmic fluorescence values were then exported and analyzed  
453 using R (<https://cran.r-project.org/>) and R-Studio (<https://rstudio.com/>). Data was processed  
454 and visualized using the tidyR <sup>46</sup> and ggplot2 <sup>47</sup> packages. Stress granules and ELP  
455 condensates were quantified by averaging the maximum pixel intensity of each cell within a  
456 frame and then averaging those values across multiple replicates.

457 **Figures**

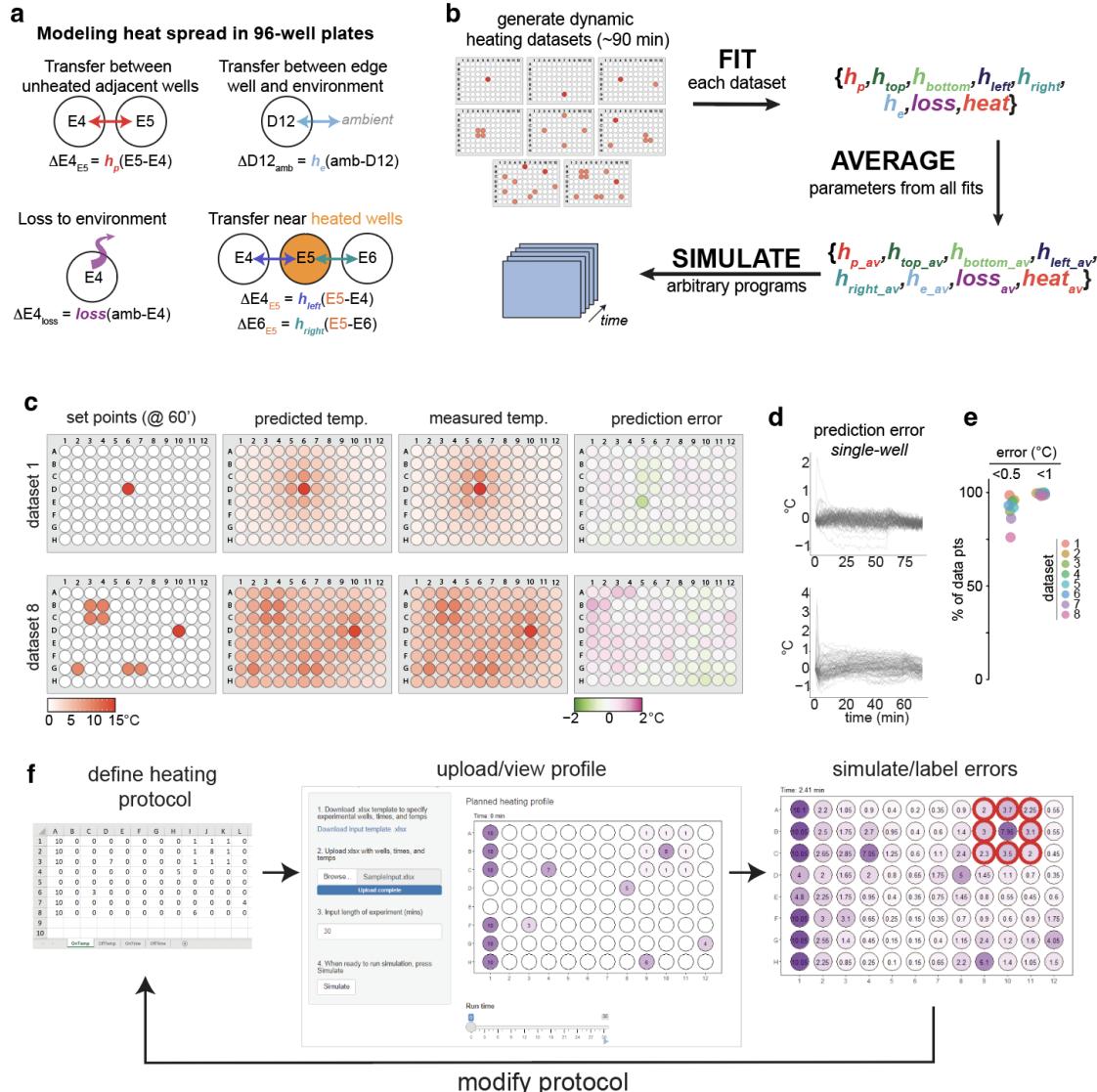


458

459 **Figure 1. Design and use of the thermoPlate for simultaneous, independent temperature**  
460 **control in 96 well plates.** a) Image of a fully assembled thermoPlate device. b) Enlarged image  
461 of a single heater/reader pair, which regulates the temperature of a single well of a 96 well plate.  
462 c) An Arduino takes readings from a reader and dynamically adjusts the duty ratio of the heater  
463 using PID feedback control. e) Raw readings taken from the reader of each well when placed in  
464 a cell culture incubator set to various temperatures before and after calibration. Color and  
465 dashed lines represent the incubator temperature. See Methods for details on calibration. f)  
466 Traces showing 16 wells that were set to 16 different temperatures in 1°C increments (28-43°C)  
467 with an ambient temperature of 25°C for 15 hrs. g) Dynamic temperature control in multiple  
468 wells over multiple temperatures. 7 wells were heated to different temperatures for repeated  
469 cycles of 30 minutes of heating followed by 10 minutes of cooling to 28°C, with an ambient  
470 temperature of 25°C.



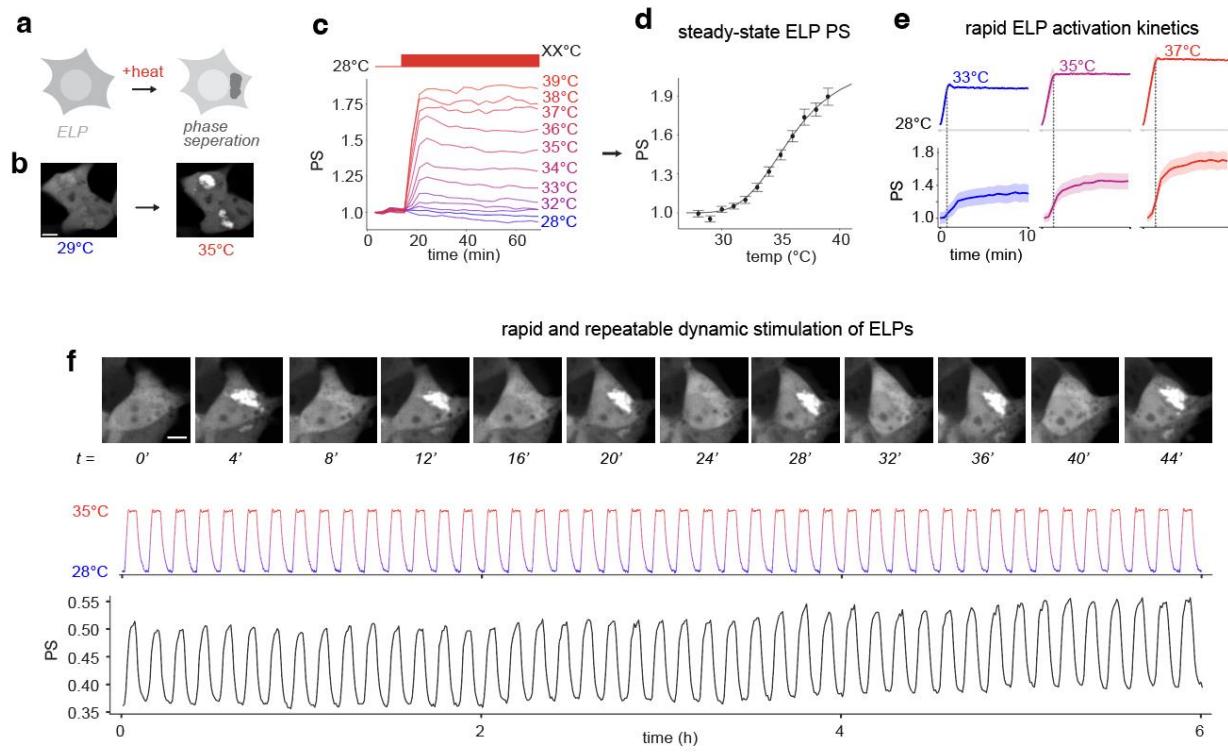
471  
472 **Figure 2. Characterization of thermoPlate heating and cooling** a) Temperature profile of  
473 each well heated with a 20 min pulse of 10°C. Wells were heated sequentially. b) The mean of  
474 all 96 traces from (a) labeled with key metrics of thermoPlate performance. Heating error =  
475 difference between the measured and set point temperature at 20 minutes. Rise time = duration  
476 at which temperature reached 0.5°C of the desired temperature. Overshoot = difference  
477 between maximum measured and set point temperature during heating. Cooling error, fall time,  
478 and undershoot represent the same parameters during the cooling phase. Error ribbons = 1 SD.  
479 s.s. = steady state. c) Heating each of 12 wells through 5, 10, 15, or 20°C for 20 minutes and  
480 then cooling to 28°C (ambient = 25°C). d) Mean of each performance metric from (c). Error bars  
481 represent = SD. e) Heating amplitude and speed can be increased by heating groups of wells.  
482 Wells were programmed to heat by  $\Delta T = 20^\circ\text{C}$  as either single wells or a group of 9 wells. f)  
483 Performance metrics of heating as a function of well grouping. Wells were heated by  $\Delta T = 10^\circ\text{C}$ .  
484 Data represents mean +/- SD from 3-9 wells.



486 **Figure 3. Mathematical modeling predicts the temperature of all wells in a sample plate.**

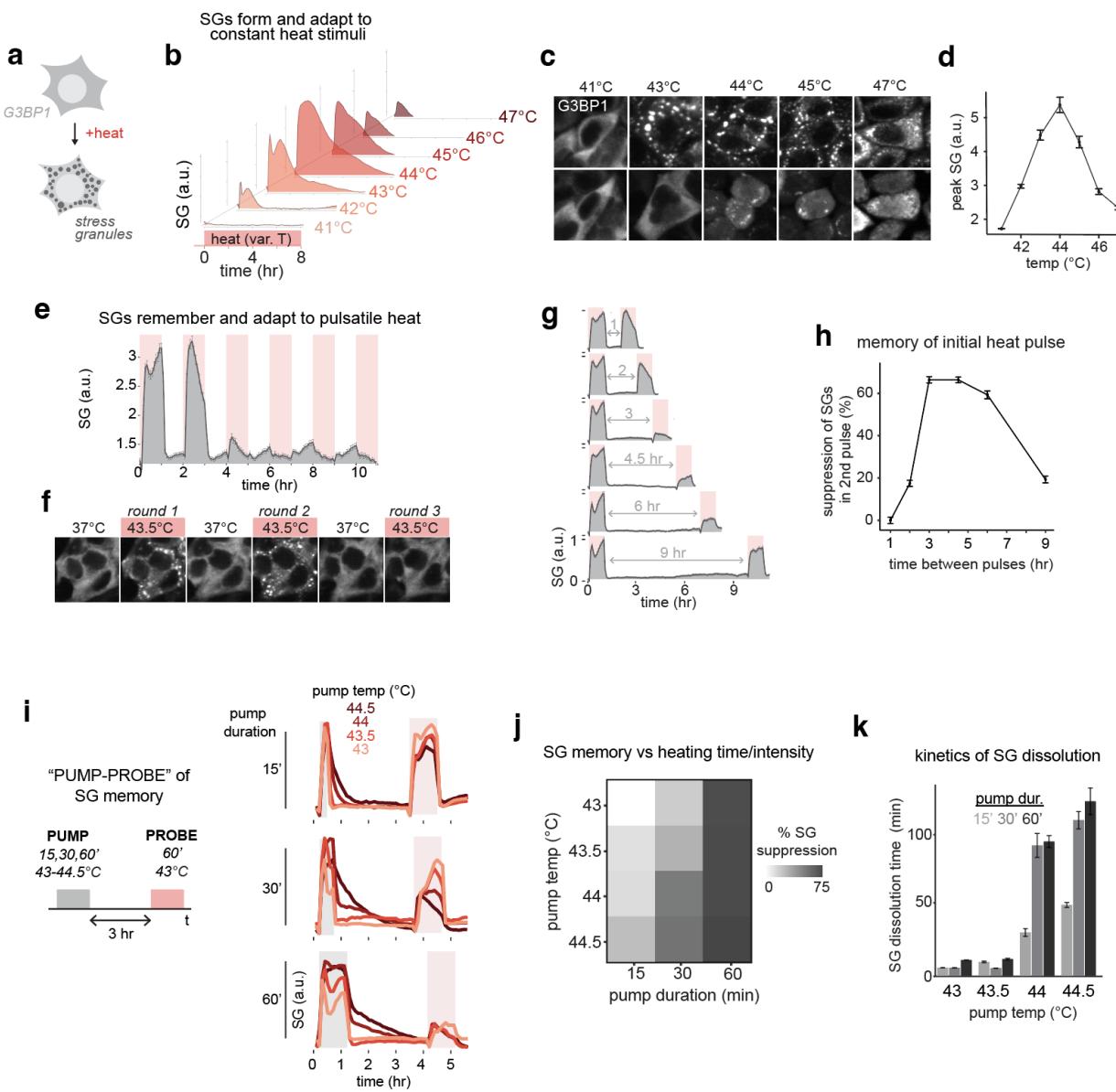
487 a) Diagram of parameters used to model heat diffusion in a 96-well plate that is heated by a  
 488 thermoPlate. b) Strategy for parameter fitting. 8 parameter sets were fit to 8 separate  
 489 experiments that spanned a range of wells, temperature, and heating dynamics. Parameters  
 490 were then averaged across the parameter sets. c) Comparisons of set points, model  
 491 predictions, actual temperatures, and error between predictions and model for the 60 min time  
 492 points of two heating datasets. d) Traces of error (predicted-measured) within each well over the  
 493 entire duration of two datasets. Each trace represents the error within a single well. e)  
 494 Summaries of error for each well and time point of each data set. f) A Shiny App GUI allows  
 495 users to upload heating profiles, review the profiles visually, and simulate heating and heat  
 496 diffusion during a run. The simulation shows the predicted temperature of all wells and indicates  
 497 whether any prespecified temperature profiles are unachievable due to thermal diffusion from  
 498 neighboring wells (red circles in right panel). The user can then adjust and simulate the adjusted  
 499 protocol until no errors are found.

500



501

502 **Figure 4. The thermoPlate allows rapid characterization of the phase separation of an**  
503 **elastin-like polypeptide (ELP<sub>53</sub>)**. a) Diagram of ELP phase separation (PS) upon elevated  
504 temperature. b) Representative image of ELP phase separation when transition from 29°C to  
505 35°C in HEK 293T cell lines. c) Cells expressing ELP-GFP were exposed to various  
506 temperatures for 1 hour while undergoing simultaneous confocal microscopy. Each trace  
507 represents the mean of ~500 cells. PS values are normalized to t = 0. See methods for PS  
508 quantification. d) The equilibrium PS at each temperature from (c) plotted vs temperature. Each  
509 point represents the mean of ~500 cells +/- 1 SEM. PS is normalized to values at 28°C. e)  
510 Quantification of the kinetics of ELP PS at 3 temperatures. Traces represent the mean of ~500  
511 cells +/- 1 SEM. PS values are normalized to t = 0. f) Rapid and dynamic control of ELP PS can  
512 be maintained indefinitely. Data show 45 cycles of 4 minutes of heating to 35°C followed by 4  
513 minutes at 28°C. The temperature trace represents the mean of 3 wells. PS traces represent the  
514 mean of ~1000 cells.

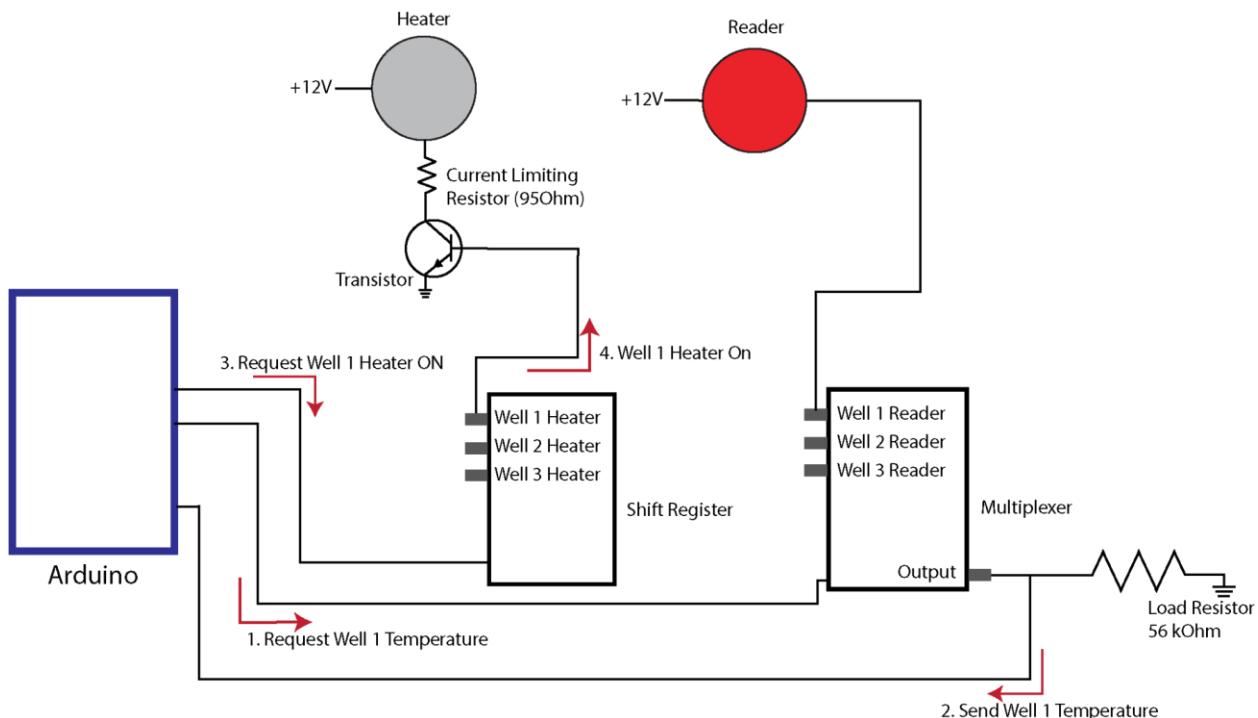


515

516 **Figure 5. The thermoPlate allows visualization of mammalian heat shock response with**  
 517 **high resolution in temperature and time.** a) Heat induced stress granule (SG) formation was  
 518 observed by measuring condensation of G3BP1-mCh in response to time varying temperature  
 519 changes in HeLa cells. b) SG formation in response to constant heating. Each trace represents  
 520 mean SG formation from two wells. c) Representative images of experiment in (b). d) Maxima of  
 521 SG formation across all temperatures tested. e) SG response dynamics in response to pulsatile  
 522 stimuli. Red bars indicate heating intervals (1 hr at 43.5°C), which were separated by 1 hr  
 523 intervals at 37°C. f) Interrogating temporal properties of heat shock memories that suppress  
 524 stress granule formation. Cells were heated for 1 hr intervals (43.5°C) separated by variable  
 525 intervals at 37°C, and SG formation on the second pulse was measured. h) The degree of

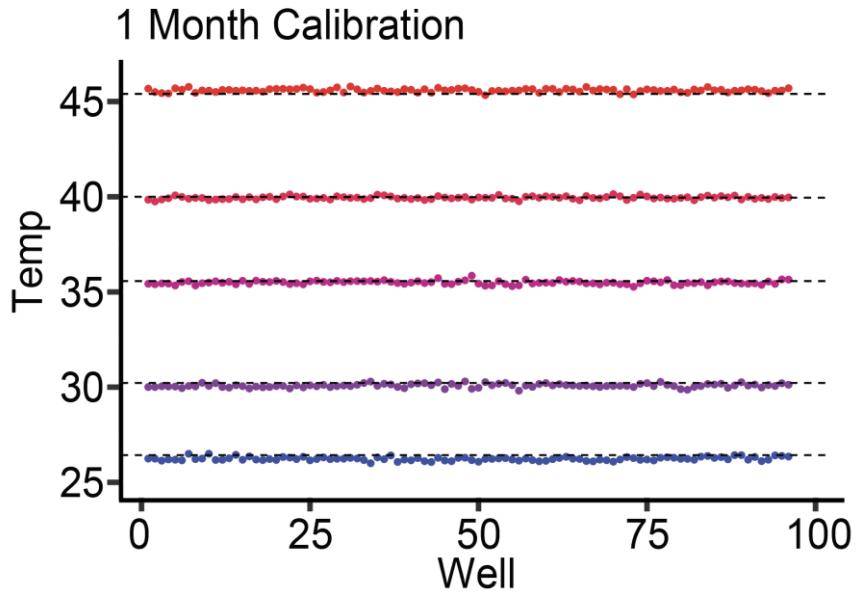
526 suppression of SGs on the second pulse from (g) is quantified as a function of the interval  
527 between heat pulses. i) A “pump-probe” experiment to interrogate the effects of temperature  
528 and duration of heat shock on SG memory. The temperature and duration of the “pump” heat  
529 pulse were varied, and the magnitude of SG formation on a second heat pulse (1 hr, 43.5°C)  
530 was measured. j) Quantification of SG memory (suppression during the probe pulse) as a  
531 function of pump temperature and duration from experiment in (i). k) Quantification of kinetics of  
532 SG dissolution after the pump pulse. Dissociation time was calculated as the time at which SG  
533 magnitude dropped 90% from its peak value before adaptation. Data represents mean +/- SD of  
534 four replicates.

535 **Supplementary Figures**



536

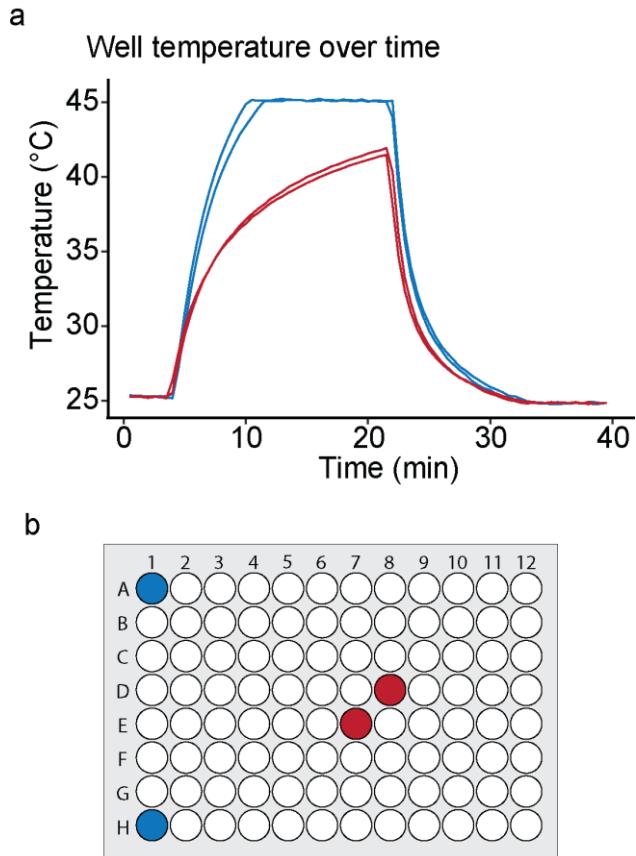
537 **Figure S1. Wiring diagram of a single Heater/Reader thermistor pair.** Each Heater/Reader  
538 pair is controlled via an Arduino microcontroller which coordinates the following actions: 1) The  
539 temperature of a well is requested by the Arduino from the multiplexer. 2) The multiplexer  
540 retrieves and returns the temperature of the well via the Reader. The Arduino calculates the  
541 necessary heating intensity (in the form of a duty cycle) via a PID algorithm. 3) The Arduino  
542 requests that the shift register accordingly adjusts the on/off state of the Heater. 4) The shift  
543 register will adjust the Heater accordingly. This process is repeated 96 times for each well of a  
544 96-well plate.



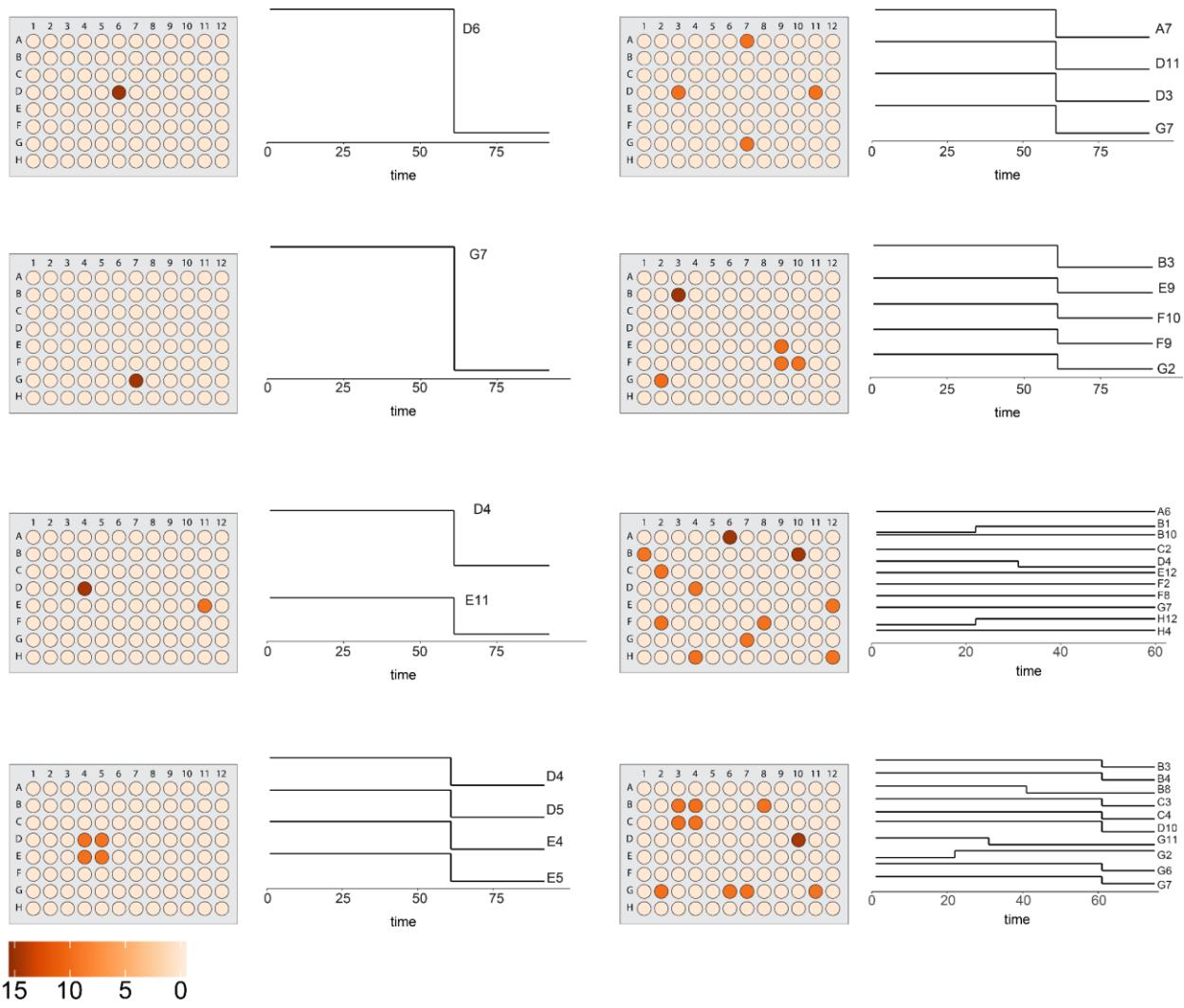
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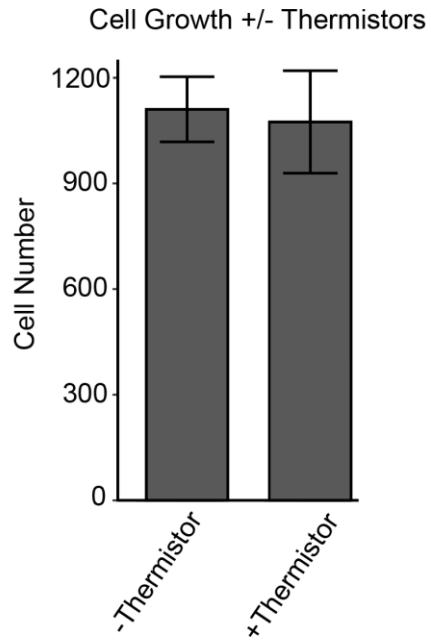
546 **Figure S2. Calibration stability after 1 month of use.** a) The same procedure performed in  
547 Figure 1e (raising the ambient temperature of an incubator in 5°C increments and recording  
548 thermoPlate readings at those temperatures) was repeated on the same thermoPlate 1 month  
549 after initial calibration, with temperature readings adjusted using the same calibration values  
550 produced in Figure 1e. Average reading error remained unchanged compared to prior  
551 calibration (<0.1°C) while average SD between wells increased marginally to 0.086°C, indicating  
552 calibration values are stable over time. Color and dashed lines represent the incubator  
553 temperature. See Methods for details on calibration.

554

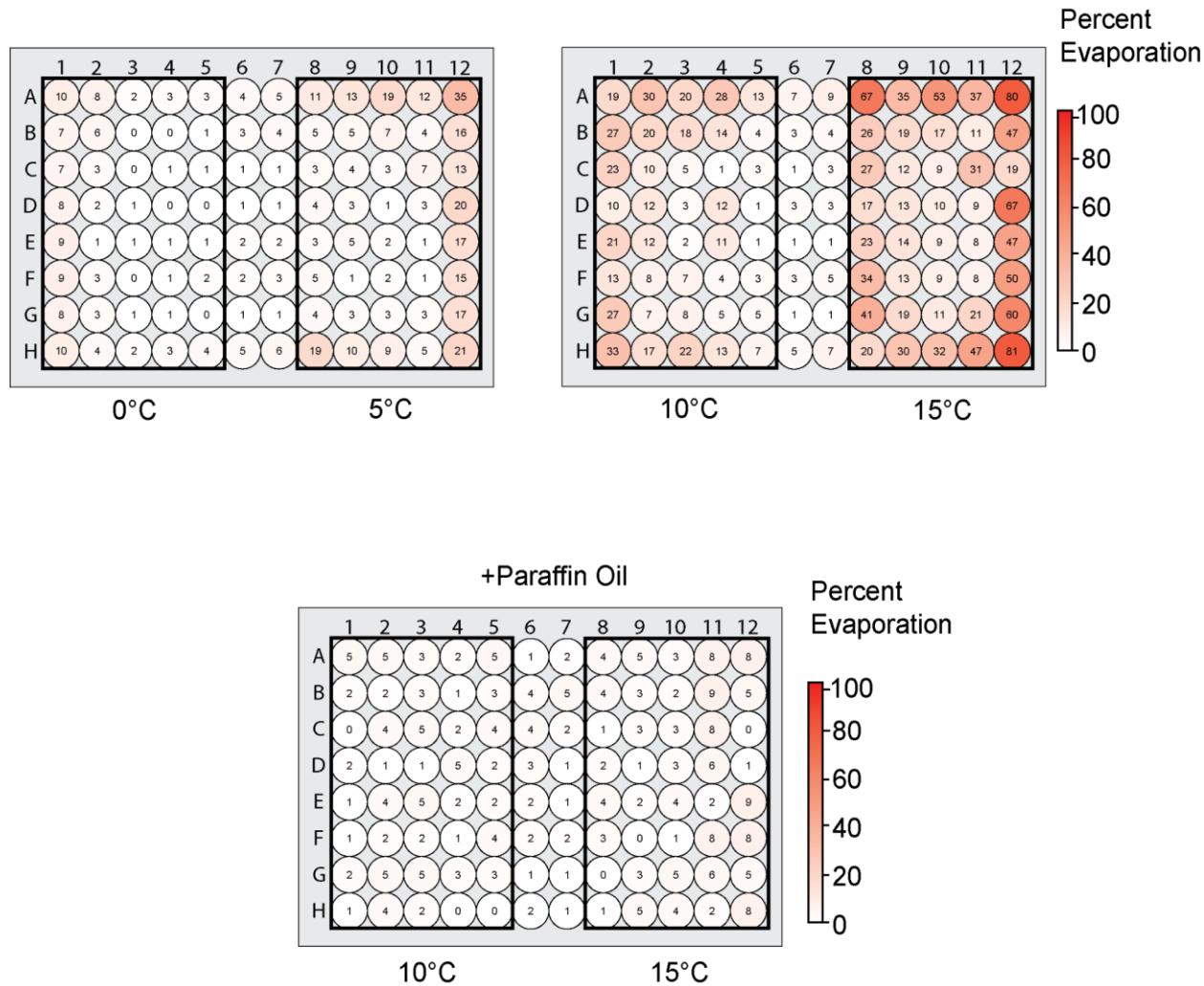


555 **Figure S3. Edge wells heat faster and reach higher steady states.** a) Traces are taken from  
556 Figure 2c, showing kinetics and steady state temperature from wells undergoing constant  
557 heating. Color represents the position on the plate (center wells labeled in red and edge wells  
558 labeled in blue) showing that edge wells are able to achieve faster and higher steady state  
559 heating than center wells. b) Diagram of the position on the plate of each trace found in (a).

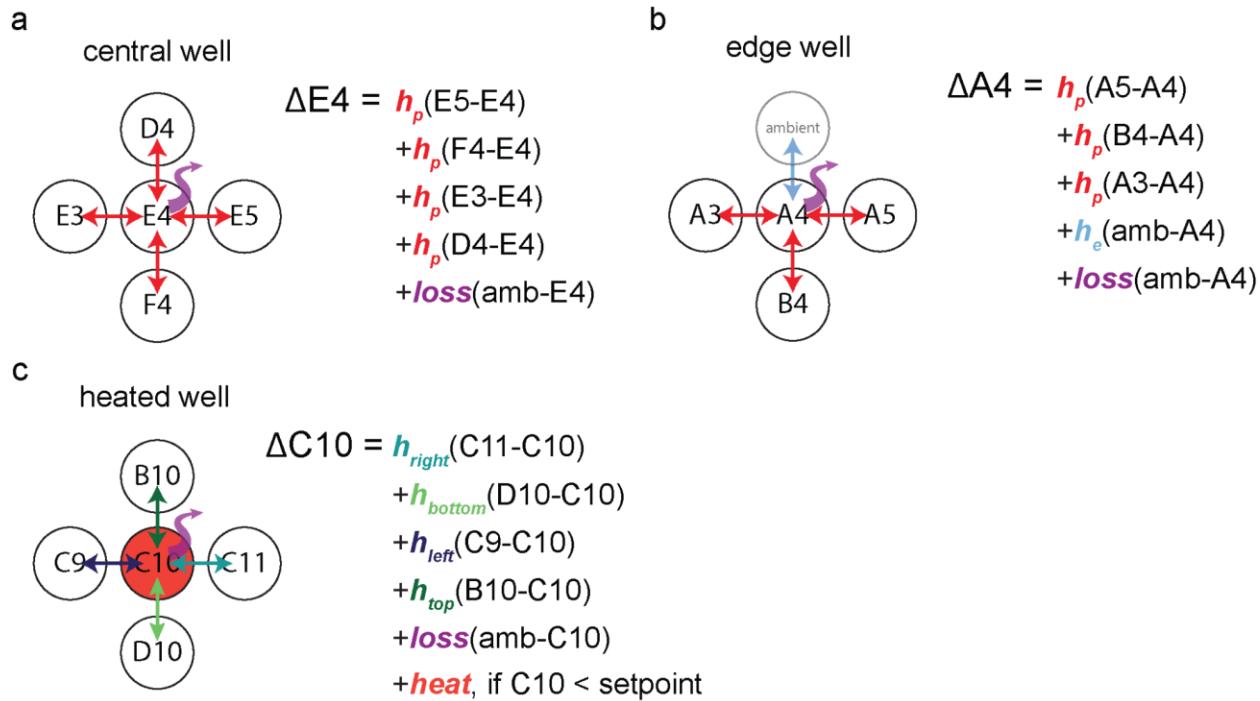




566 **Figure S5. Cell growth quantification with and without exposure to the thermoPlate. a)**  
567 Every well of a 96 well plate was seeded with HeLa cells and kept for two days in a standard  
568 cell culture incubator, with half of the wells exposed to thermoPlate heater/readers. Cells were  
569 then fixed, stained with DAPI, imaged, and nuclei were counted to quantify population size in  
570 each well. No significant difference in cell numbers was observed between the two conditions,  
571 indicating that cell growth and survival are not significantly altered by thermoPlate exposure.  
572 Each bar represents the mean +/- SD of N = 48 wells.



573  
574 **Figure S6. Evaporation during thermoPlate usage.** a) A 96 well plate filled with 150uL of  
575 PBS was placed in a cell culture incubator and wells were set to heat by either 5 or 10°C for 24  
576 hours. The volume of PBS remaining in the well was measured via a micropipette. Color  
577 represents the percent evaporation in each well. b) The same procedure as (a) was repeated  
578 heating wells by either 15 or 20°C. Major evaporation (>50%) was limited to wells set to 20°C  
579 above ambient. c) The same procedure as (b) was performed with a thin layer of 50uL of  
580 paraffin oil coating each well to prevent evaporation. Evaporation in all wells was <10%  
581 regardless of heating demonstrating that paraffin oil can be used to extend the duration of long-  
582 term heating.



583  
584

585 **Figure S7. Diagram of all parameters used for model fitting.** To fit heat spread across a 96  
586 well plate during thermoPlate operation, wells were categorized into 1 of 3 categories: a)  
587 a central well that transfers heat only to neighbor wells and vertically to the environment, b)  
588 edge wells that transfer heat to neighbors, vertically to the environment, and horizontally to the  
589 environment, and c) heated wells that can transfer heat in any of the ways shown in (a, b) but  
590 also receive heat deposition from a thermoPlate heater. For all scenarios, the temperature  
591 difference between the well of interest and its neighbors/environment are weighted by fitted  
592 parameters to calculate heat transfer.

593 **Supplementary Movie Captions**

594

595 **Movie S1. Rapid cycling of ELP phase separation.** HEK293T cells expressing GFP- ELP<sub>53</sub>  
596 exposed to 4 minutes of 35°C followed by 4 minutes of 29°C repeatedly over 45 cycles. ELP  
597 condensates rapidly formed and deformed with temperature over all cycles. Time is hh:mm:ss.  
598 Scale bar = 10  $\mu$ M

599

600 **Movie S2. Stress granule formation during constant heat stress.** HeLa cells expressing  
601 G3BP1-mCherry were imaged during exposure to various temperatures (41°C-47°C) over  
602 ~8hrs. Stress granules form in response to 42°C but are transient in all cases, decreasing over  
603 time after initial activation. Time is hh:mm. Scale bar = 30  $\mu$ M

604

605 **Movie S3. Stress granule adaptation during pulsatile heat stress.** HeLa cells expressing  
606 G3BP1-mCherry were exposed to continuous rounds of 1 hr at 43.5°C followed by 1hr at 37°C.  
607 Stress granules form in response to the first two pulses of heat but are significantly attenuated  
608 upon the third pulse, indicating a memory of previous stress. Time is hh:mm. Scale bar = 15  $\mu$ M.

609

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