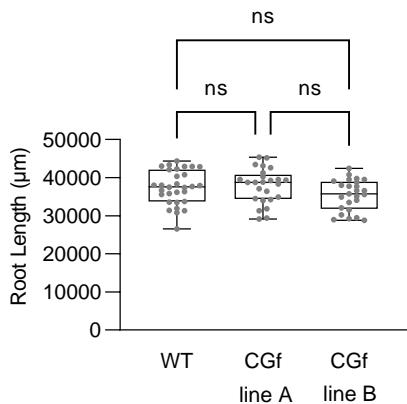
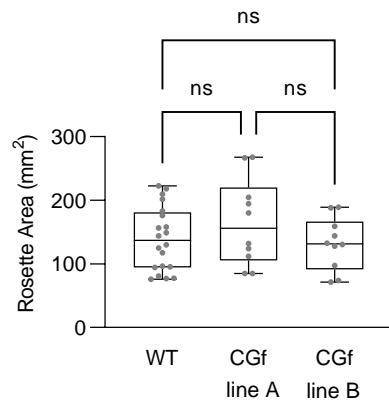
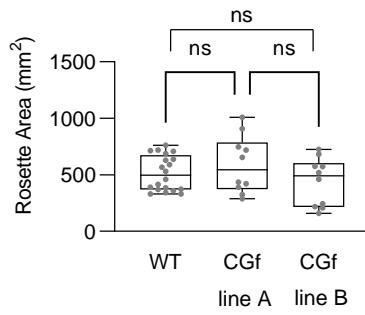
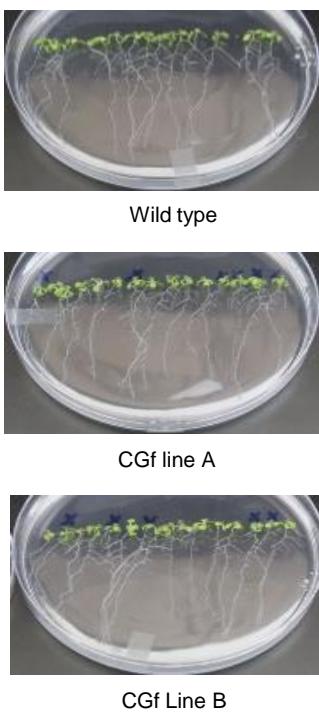
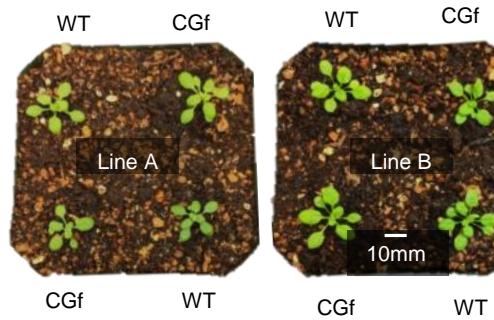
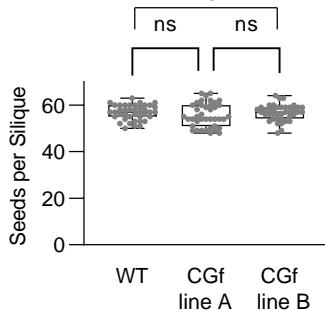
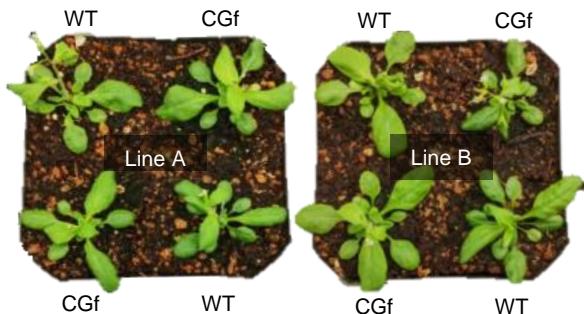
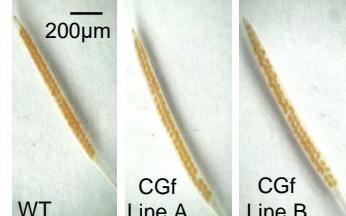
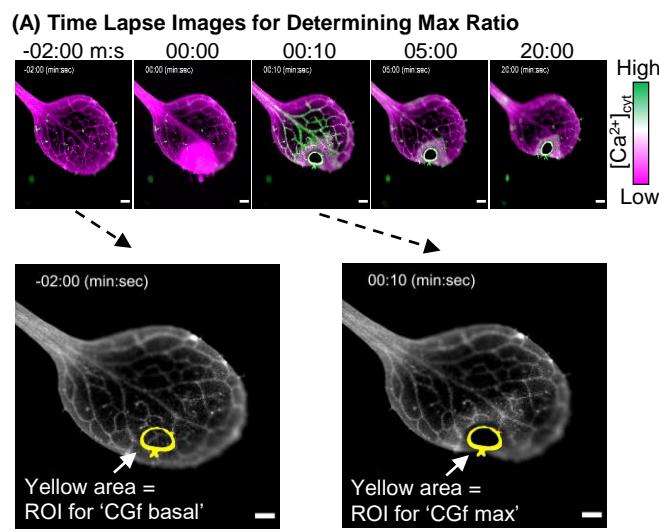


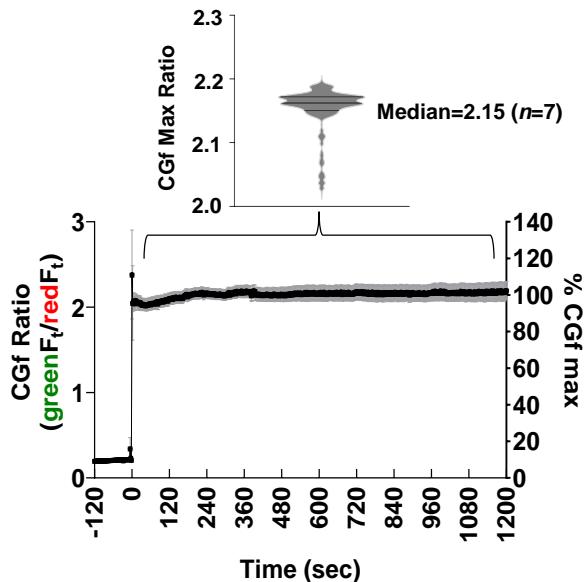
Supplemental Figure S1. Western blot detection of full length mCherry-GCaMP6 proteins (77kDa) *in planta* using primary RFP monoclonal antibody and secondary F(ab')2-Goat anti-Mouse IgG (H+L) (HRP) antibody. Left to right, Lane 1-8: COL-0 negative control, CGf, CGm, CGu, followed by confirmation with second transgenic line in lanes 5-8. CGf (fast), CGm (medium), and CGu (ultra) are identical mCherry fusions to three GCaMP6 variants GCaMP6f (K_D 220-375nM), GCaMP6m (K_D 167nM), and GCaMP6u (K_D 890nM) ([Chen et al., 2013; Helassa et al., 2016](#)).

(A) Root Length Analysis**(B) Rosette Area at 4 Weeks****(C) Rosette Area at 5 Weeks****(D) Root Length Images****(E) Picture of 4-Week-Old Plants****(G) Seed Set Analysis****(F) Picture of 5-Week-Old Plants****(H) Destained Siliques**

Supplemental Figure S2. Growth comparisons between wild type and CGf plants provide evidence that CGf expression did not alter plant development and reproduction. (A) Comparison of root lengths of 8-day old seedlings grown on plant media described in methods. n=30 (WT) or n=25 (CGf line A and B) from 5 biological replicates. (B,C) Measurement of total rosette area of the (B) 4-week-old and (C) 5-week-old plants. n=20 (WT) or n=10 (CGf line A and B) from five biological replicates. (D- F) Representative pictures of root length or plant growth shown in above data (A-C, respectively). (G) Seed set comparison between WT and two independent CGf lines. n=40 of siliques for all 3 groups from 10 independent plants in replicate. (H) Representative photos of decolorized siliques used to measure seed set. Scale bar = 200 μm . (A-C, G) Statistical analyses were done using One-way ANOVA multiple comparisons with the Turkey post-hock test. Error bars in each box plot show the min to max values of representing data sets (grey dots). “ns” = not significant. All figures are comparisons between WT and two independent lines: CGf line A (ss2543) and line B (ss2544).

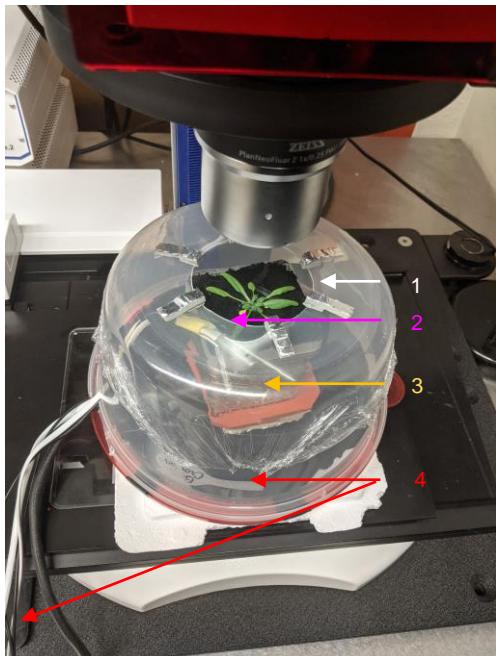


(B) CGf Ratio and % CGf Max in Leaves

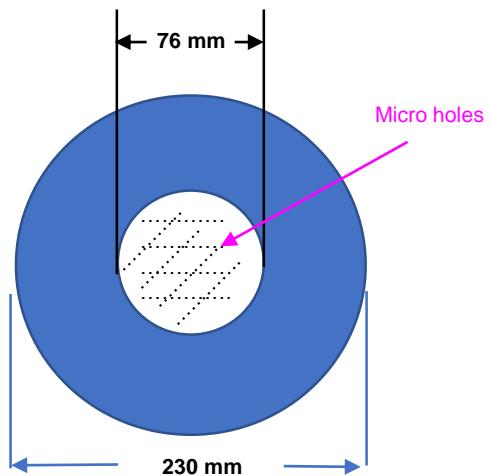


Supplemental Figure S3. Determining maximum CGf ratio in leaf. **(A)** Representative time lapse images of a true leaf expressing CGf. Time indicates minutes:seconds (m:s). Negative time points represent before stimulation, while positive time points are after burning a small leaf area for 2 seconds with a high power, red laser (Laser 301, 1000 MW 650nm Red Beam Light Single-point Laser Pointer; WDLaser Kowloon, Hong Kong) for 2 seconds. Color scale indicates $[Ca^{2+}]_{cyt}$ levels. Scale bar = 500 μ m. An enlarged representative black & white frame images show areas selected for detecting CGf basal (yellow ROI on the left side -2 min time point image) or CGf max (yellow ROI on the right side 10 sec image). **(B)** Fluorescent intensity and % CGf max of CGf signals in leaves at the laser-induced wound site. Fluorescent signal changes were calculated as described in methods. Images were obtained at 2 second intervals for 22min (2min pre-stimulus period followed by 20min post-stimulus) using the Zeiss AxioZoom V16 microscope described in methods. Error bars represent SEM of n=7 independent plants. % CGf max were calculated from the fluorescent intensity data using CGf max ratio (2.15). $greenF_t$ and $redF_t$ indicate GCaMP6f fluorescent intensity ($greenF_t$) and mCherry fluorescent intensity ($redF_t$) at the time of event. Violin graph (inset) shows median of maximum CGf ratio. CGf max ratio was calculated using data points from 10 to 1200 second duration (n=4172 data points from n=7 independent plants). Solid line = median value of data. Dotted lines = quartile of data.

(A) Picture of Whole Plant Heat Chamber

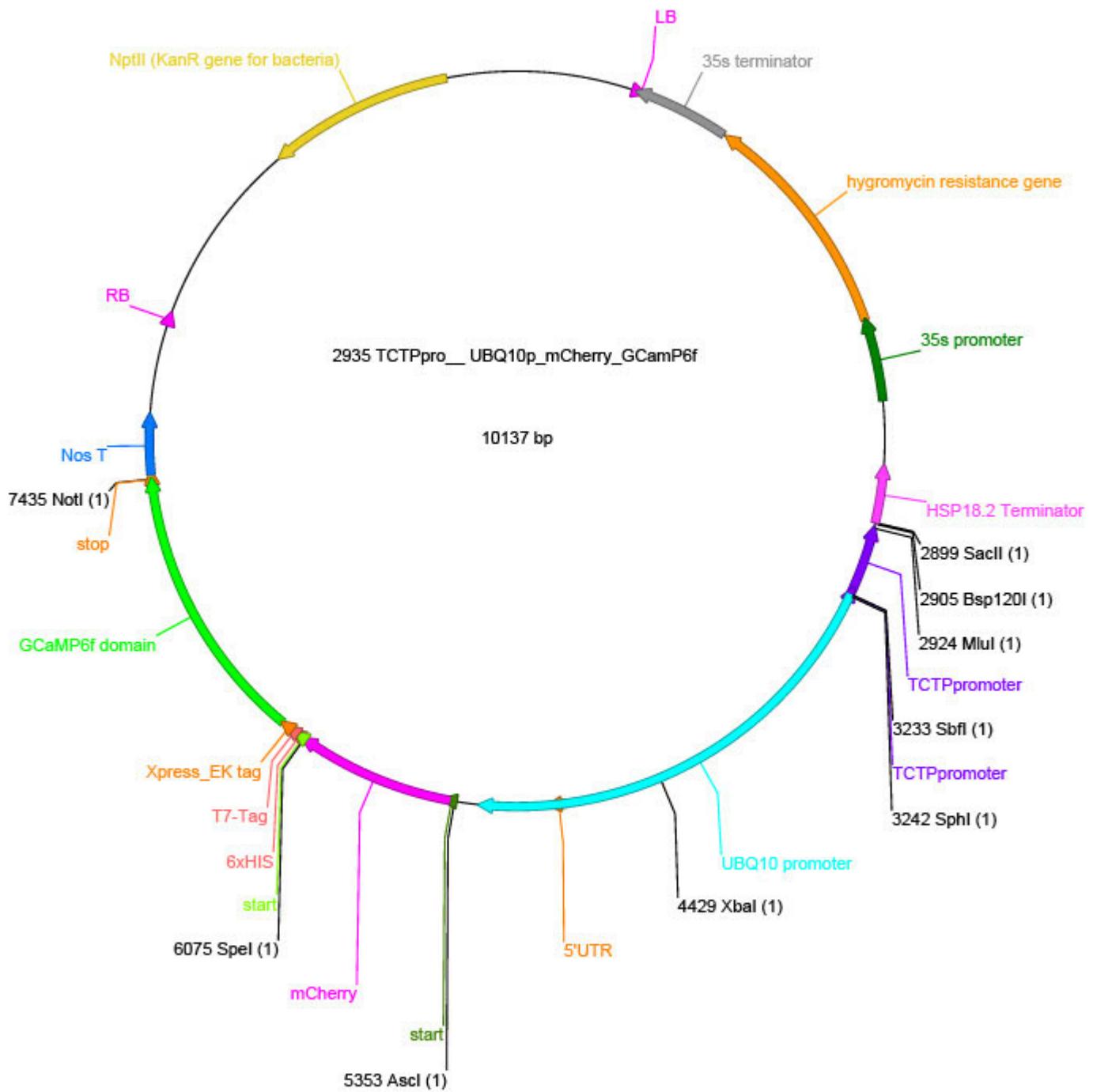


(B) Dome Lid Diagram



Supplemental Figure S4. Custom heat chamber used for whole plant heat stress experiments. (A)

Photo of the custom heat chamber with a representative 5-week-old plants under a Zeiss AxioZoom V16 microscope. Picture legend as follows. 1) Dome lid of the heating chamber (white). 2) Representative 5-week-old WT plants grown in GA-7-3 magenta vessel. 3) 5mm thick silicon pad with heat insulation tape on the bottom (orange). A thick silicone pad with heat insulation tape is used to prevent direct heating of soil vessels during heat stress experiments. 4) A digital heating pad with a power cable (red). **(B)** Dome lid diagram showing open 76mm diameter window opening for imaging plants during heat stress experiments. To prevent drought during heat stress experiment, the 76mm diameter window was covered in clear plastic wrap with hundreds of micro-holes. Micro-holes were made using a 27-gauge needle to prevent condensation during heat stress experiment.



Supplemental Figure S5. CGf Plasmid Map and DNA Sequence

TCTPpro __ UBQ10p_mCherry_GCamP6f

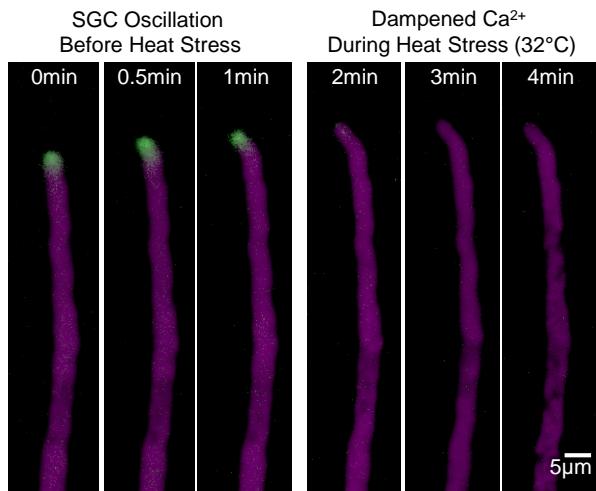
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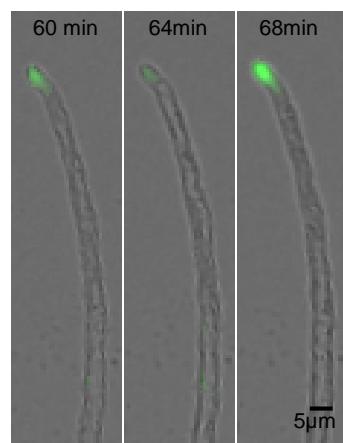
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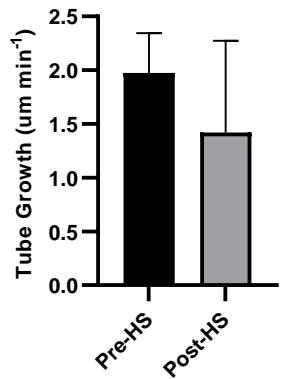
A) Time Lapse Images Before and During Heat Stress



B) Recovered Ca^{2+} Signal



C) Growth Rate Before and After Heat Stress



Supplemental Figure S6. Heat stress triggered dampened calcium oscillations do not indicate pollen cell death. (A) Representative time lapse images showing steady growth tip-focused Ca^{2+} (SGC) oscillations shifting to a dampened Ca^{2+} (DC) oscillation during heat stress. Time course: 22°C for 1min, $22\text{-}32^\circ\text{C}$ in 1min, followed by 2min at 32°C . (B) Tip-focused Ca^{2+} oscillations (green) are restored within 1hour post-heat stress, as shown by representative GFP/brightfield overlay images of the same pollen tube shown in pane A. (C) Post-heat stress (post-HS) growth rates are similar to growth rates observed prior to heat stress (pre-HS). Heat treated pollen tubes recovered in the dark at 22°C for ~1hour prior to measuring pollen tube growth over a 10min time course. Error bars are standard deviation of $n=6$ pollen tubes.