# The Role of Potassium and Host Calcium Signaling in *Toxoplasma gondii* egress

Stephen A. Vella et al

## Supporting Information Supplemental Figures:



Fig. S1. Impact of host histamine signaling on T. gondii Ca<sup>2+</sup>: A, stable HeLa cells expressing jRGECO1a were infected with tachyzoites expressing jGCaMP7f localized to the PV. Dashed white outlines indicates the area used for the green fluorescence (PVs) analysis and the red squares show the region of the host cell used for the analysis of the jRGECO1a fluorescence shown in the graphs. B, HeLa cells transiently expressing the red GECI RGECO infected with tachyzoites expressing cytosolic GCaMP6f. Dashed white outlines show the parasites used for the fluorescence analysis shown in the graphs and the dashed red square shows the region of the host cell used to analyze red fluorescence. C. HeLa cells infected with tachyzoites expressing cytosolic GCaMP6f and RGECO in the PV. Dashed white outlines indicate parasites and dashed red outlines indicate the PV region used to analyze the fluorescence shown in the graph. Under all conditions 100 µM Histamine was added at the 1 min mark and the response recorded. Numbers at the upper right of each panel indicate the time frame of the video. Tracings to the right of each panel shows green fluorescence (GCaMP6f or jGCaMP7f) and red fluorescence (iRGECO1a or RGECO) fluctuations with the scales for the green fluorescence shown on the left Y axis and for the red fluorescence on the right Y scale. Ca<sup>2+</sup> fluctuations of PVs are shown in A (green) and C (red). Single parasites fluctuations are shown in B and C (green). Host cytosolic changes are shown in A and B (*red*). Bar graphs represent quantification of the average  $\Delta F$  values of a minimum of three independent trials (red, jRGECO1a or RGECO) (green, GCaMP6f or jGCaMP7f) with Standard Error of the Mean (S.E.M.) represented as error bars.



% of responsive tachyzoites expressing cytosolic GCaMP6f to carbachol or histamine. The experiment are the ones presented in Fig. 1B and S1B. C, % of tachyzoites responding to Carbachol or Histamine addition. The experimental conditions are the ones from Figs 1C and S1C. Light green portions are used to represent the percentage of responsive cells and dark green portions are used to represent the percentage of non-responsive cells.



Figure S3. Extracellular tachyzoites expressing cytosolic GCaMP6f do not respond to carbachol or histamine: The fluorescence changes of  $2 \times 10^7$  cells/mL of GCaMP6f tachyzoites resuspended in Ringer Buffer supplemented with 2 mM Ca<sup>2+</sup> were analyzed in a fluorescence spectrophotometer, Hitachi 7000. Excitation was set at 485 nm and emission at 520 nm. Traces shown are an average of three independent experiments conducted on separate cell preparations. (A) Addition of 1 mM Carbachol (A) or 100  $\mu$ M Histamine (B) at 60 seconds lead to no increase in the GCaMP6f signal and at 200 second 1  $\mu$ M of Ionomycin was added as control for responsiveness.



**Fig S4. Calibration of Ca<sup>2+</sup> threshold:** (A) RH tachyzoites (5 x 10<sup>7</sup>) were loaded with FURA2-AM and resuspended in 2.5 mL of Ringer's Buffer supplemented with 100  $\mu$ M EGTA. The cytosolic Ca<sup>2+</sup> response to the addition of various concentrations of ionomycin is shown. The figure shows the average of at least 2 trials of each condition; (B) Quantification of ionomycin titration curve. The first 15 seconds of the slope after addition of ionomycin was determined and averaged from at least two independent trials



Fig. S5. Ionomycin Induced Egress in Ca<sup>2+</sup> Free Buffer: A) Hela cells were infected with GCaMP6f-expressing tachyzoites in buffer supplemented with 100  $\mu$ M EGTA to create a Ca<sup>2+</sup>-free extracellular buffer. 1  $\mu$ M Ionomycin was added to trigger egress. B) Representative fluorescence tracings of ionomycin addition in Ca<sup>2+</sup> free buffer. Inset: quantification of ionomycin induced egress between Ca<sup>2+</sup>-rich and Ca<sup>2+</sup>-free media. Error bars represent the Standard Error of the Mean (S.E.M.)



Fig S6. Off targeting effect of zaprinast on host cells: HeLa cells stably expressing jRGECO1a were used for video microscopy analysis. 100  $\mu$ M Zaprinast was added at approximately 60 seconds and 1  $\mu$ M ionomycin was added at 300 seconds.  $\Delta$ F represents the difference between the maximum fluorescence and minimum fluorescence 10 seconds before and 10 seconds after addition of either zaprinast or ionomycin



Fig S7. Parasites that do not show the second Ca<sup>2+</sup> peak are less likely to egress. A, Egress was induced in Hela cells infected with  $\Delta$ PLP1 GCaMP6f parasites using 100  $\mu$ M zaprinast; **B and C**, Representative fluorescence tracings of  $\Delta$ PLP1 GCaMP6f after addition of 100  $\mu$ M Zaprinast for egress and non-egressing parasites respectively. Notice, two peaks are evident for the parasites that were successfully able to egress and only one peak was visible for the parasites that were unable to egress. **D**, Quantification of egressing and non-egressing  $\Delta$ PLP1 GCaMP6f parasites. The average  $\Delta$ Fluorescence (F<sub>max</sub> - F<sub>min</sub>) of at least 10 cells per either egressing and non-egressing were averaged for 3 independent trials.

### Legends for Supplemental Videos

**Video S1. Carbachol addition in GCaMP6f Infected Host Cells:** Hela cells transiently expressing cytosolic localized RGECO were infected with cytosolic GCaMPf6 expressing parasites. 1 mM Carbachol was added at approximately the 1 min mark. (Fig. 1B shows still images from this video).

**Video S2. Thapsigargin stimulation in LAR-GECO expressing host cells:** Hela cells transiently expressing the low affinity mitochondrial Ca<sup>2+</sup> indicator LAR-GECO1.2 were infected with cytosolic GCaMP6f expressing parasite. 2  $\mu$ M Thapsigargin was added at approximately the 1 min mark. (Fig. 2B shows still images from this video).

**Video S3. Two-peaks of Ca<sup>2+</sup> contribute to parasite egress:** Hela cells infected with cytosolic GCaMP6f parasites were treated with 100  $\mu$ M of Zaprinast in media supplemented with 2 mM Ca<sup>2+</sup>. (Fig. 3E shows still images from this video).

**Video S4. Natural egress in cpd 1 synchronized parasites:** Hela cells stably expressing cytosolic jRGECO1a infected with cytosolic GCaMP6f expressing parasites. After 24 h post infection 1  $\mu$ M cpd1 was added. 24 hr post cpd1 addition parasite egress was imaged after cpd1 wash off. (Fig. 4A shows still images from this video).

Video S5. Whole-cell patch under high  $K^+$  and high  $Ca^{2+}$  conditions: Hela cells infected with GCaMP6f parasites were whole-cell patched with a patch pipette solution containing 140 mM  $K^+$  and 10  $\mu$ M  $Ca^{2+}$ . Note parasite egress is slow, methodical, and occurring one-by-one in comparison to detergents and ionophores. Video presented in real time. (Fig. 6B shows still images from this video).

**Video S6. Whole-cell patch under low K<sup>+</sup> and high Ca<sup>2+</sup> conditions:** Hela cells infected with GCaMP6f parasites were whole-cell patched with a patch pipette solution comprising 10 mM K<sup>+</sup>, 130 mM choline chloride, and 10  $\mu$ M Ca<sup>2+</sup>. Parasite egress was slower in comparison to detergents and ionophores, but more rapidly in comparison to high K<sup>+</sup> conditions. Parasite egress occurred almost explosively (all parasites egressed at once) under these conditions. Video presented in real time. (Fig. 6C shows still images from this video).

## Supplemental Tables

GECI	Kd (nM)	Dynamic Range	Use	Source or Reference
GCaMP6f	375	51.8	Standard, well established GECI, with fast on- off kinetics. We previously used this GECI to study <i>T. gondii</i> Ca <sup>2+</sup> dynamics	[1]
R-GECO	482	16	First generation red GECI, enables multi-color imaging when co-expressed with GCaMP6f or jGCaMP7f	[2]
jRGECO1a	148	12	Second generation red GECI with improved kinetics and enhanced sensitivity	[3]
jGCaMP7f	174	30.2	Successor to GCaMP6f with improved sensitivity and faster on/off kinetics	[4]
LAR- GECO1.2	12,000	8.7	Low affinity red GECI with optimized kinetics/sensitivity for the mitochondria of mammalian cells. Host mitochondria surrounds the PV and this indicator permits to see Ca <sup>2+</sup> fluxes through the host mitochondria	[5]

<b>Table S1: Genetically</b>	Encoded Calcium Indicators	(GECI) used in this study
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## Table S2: Parasite strains generated and used in this work

Strains	Characteristics	Source/Ref
RH with cytosolic GCaMP6f	Non-selectable drug RH strain expressing GCaMP6f in the cytosol	[6]
RH PV-jGCaMP7f	Expression of jGCaMP7f in the lumen of the PV	This study
RH PV-RGECO Cyto- GCaMP6f	Expression of PV-RGECO in the lumen of the PV and cytosolic expression of GCaMP6f	[6]

Table S3: Plasmids used for this work

Plasmid	Characteristics	Source or Reference
pGP-CMV-GCaMP6f	Transient expression of the GECI GCaMP6f in mammalian cells	[1]
pCMV-NLS-R-GECO	Transient expression of the GECI R-GECO in mammalian cells	[2]
pGP-CMV-NES-jRGECO1a	Transient expression of the GECI jRGECO1a in mammalian cells	[3]
pGP-CMV-jGCaMP7f	Transient expression of the GECI jGCaMP7f in mammalian cells	[4]
pCMV-mito-LAR-GECO1.2	Transient expression of the GECI LAR- GECO1.2 in the mitochondria of mammalian cells	[5]
ptub_IEα- DsRed_DHFR_sag1CATsa g1	Plasmid vector for expressing proteins in the lumen of the PV	[7]
рСТН3	Random Integration of a chloramphenicol selection plasmid expressing 3xHA epitopes	[8]
ptub_IEα-R- GECO_DHFR_sag1CATsag 1	Expressing the GECI R-GECO in the lumen of the PV	[6]
pUltra	Lentivirus expression of a gene of interest	[9]
jRGECO1a-pUltra	Lentivirus expression of the Red GECI jRGECO1a	This study

Table S4:	Oligonucleotide	nrimers	used in	this work
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To construct PV-jGCaMP7f			
SV229-jGCaMP7f-GIB- FWD	GGGCTGCAatgggttctcatcatcatc	This study	
SV72-jRCaMP1b-Gib-R	ccgggacgtcgtacgggtacctaggCTTCGCTGTCA TCATTTGTAC	This study	
SV77-P30-Gib-Fwd	TTTCTTGAATTCCCTTTTAGATCTATGTTT CCGAAGGCAGTGAGACG	This study	
SV-228-P30-Gib-Rev	gaacccatTGCAGCCCCGGCAAACTC	This study	
SV225-pCTH3-Gib-FWD	cagcgaagCctaggtacccgtacgac	This study	
SV226-Tub-Gib-Rev	CGGAAACATAGATCtaaaagggaattcaagaaaa aatg	This study	
To construct jRGECO1a-pUltra			
SV155V2-jRGECO1a- Agel-Fwd	aCCGGTATGCTGCAGAACGAGCTTGCTC TTA	This study	
SV156-jRGECO1a- ECORI-Rev	GAATTCGCCTACTTCGCTGTCATCATTTG TACA	This study	

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