# Supplementary Information

# cGAL, a Temperature-Robust GAL4-UAS System for *C. elegans*

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#### **Supplementary Note 1** | Vector Map of the driver construct pHW393

(pHW393) *Prab-3::nls::GAL4-SK(DBD)::VP64::let-858 3'UTR*. New driver constructs can be generated by cloning new promoters into the MCS, between Sphl/Notl/Fsel and Ascl/BssHII/BamHI/Smal/Xmal/Mscl. The backbone of pHW393 (between Sphl and Kpnl) is derived from the plasmid pPD117.01 from the Fire lab *C. elegans* Vector Kit 1997. pPD117.01 has a 5' decoy and the *let-858* 3'UTR. The 5' decoy contains a splice acceptor (decoy intron) and a short terminated coding region (decoy ORF) upstream of the MCS, which was designed to catch any upstream transcription with a 3' splice site. Adding this 5' decoy sequence was shown to decrease ectopic expression in the posterior gut (A. Fire, personal communication). We highly recommend using the pPD117.01 vector for new neuronal driver constructs.



#### Supplementary Note 2 | Sequence of the driver construct pHW393

(pHW393) *Prab-3::nls::GAL4-SK(DBD)::VP64::let-858 3'UTR*, is available from Addgene (<u>www.addgene.org</u>; plasmid no. 85583). The color scheme here matches the vector map of pHW393 shown in Supplementary Note 1.



## Supplementary Note 3 | Vector map of the effector construct pHW394

(pHW394) *15xUAS::Δpes-10::gfp::let-858 3'UTR*.New effector constructs can be generated by cloning new effector genes into the MCS, between Sall/Hincll/Kpnl/Agel and EcoRl/AccIII. The backbone of pHW393 (between SphI and KpnI) is derived from the plasmid pPD117.01 from the Fire lab *C. elegans* Vector Kit 1997. pPD117.01 has a 5' decoy and the *let-858* 3'UTR. The 5' decoy contains a splice acceptor (decoy intron) and a short terminated coding region (decoy ORF) upstream of the MCS, which was designed to catch any upstream transcription with a 3' splice site. Adding this 5' decoy sequence was shown to decrease ectopic expression in the posterior gut (A. Fire, personal communication).

1	$\label{eq:constraint} \texttt{A} \texttt{GCTTGCTAGC} \texttt{constraint} \texttt$
101	$\verb+tgtcctccgagcggagtactgtcctccgagcggagactcCCTagCccatgattacgcccaagctacgggcggagtactgtcctccgagcggagtactgtcc$
201	$\verb+ccgagcggagtactgtcctccgagcggagtactgtcctccgagcggagtactgtcctccgagcggagactcCCTagCccatgattacgccaagctacgg$
301	g <mark>cggagtactgtcctccg</mark> agcggagtactgtcctccgagcggagtactgtcctccgagcggagtactgtcctccgagcggagtactgtcctccgagcggag
401	<mark>gacto</mark> CCTagggGCGGCCGCactgactgGGCCGGCCaaaaatgcataaggttttgctggcactgttctttatctttctggcaccagcatccgcactggca
501	$\tt gtctccgaaccggcctgcagg togattttttgcaaattacgagcgttgtaggggggggggg$
601	$\tt tcattggtacattcatttacccaccttcctctttctgagcttctctgtgcttccttttttcccttattttatactgtaatttttaactttc$
701	${\tt ag} {\tt gcattgattGGCGCCCtctagaggatccccgggattggccaaaggacccaaaggtatgtttcgaatgatactaacatagaacattttcaggatcgcaaaggacccaaaggtatgtttcgaatgatactaacatagaacattttcaggatcgcaaaggacccaaaggtatgtttcgaatgatactaacatagaacattttcaggatcgcaaaggacccaaaggtatgtttcgaatgatactaacatagaacattttcaggatcgcaaaggacccaaaggtatgtttcgaatgatactaacatagaacattttcaggatcgcaaaggacccaaaggtatgtttcgaatgatactaacatagatcaacatagaacattttcaggatcgcaaaggacccaaaggtatgtttcgaatgatactaacatagatcacatagaacattttcaggatcgcaaaggacccaaaggtatgtttcgaatgatactaacatagatcacatagaacattttcaggatcgcaaaggacccaaaggtatgtttcgaatgatactaacatagatacatagaacattttcaggatcgcaaaggaccaaaggaccaaaggaccaaaggtatgtttcgaatgatactaacatagatacatagaacattttcaggatcgcaaaggaccaaaggaccaaaggaccaaaggaccaaaggaccaaaggattggcaaaggatggttggcaaaggaccaaaggaccaaaggaccaaaggatggttggt$
801	ggacccttgGCTAGCgtcgacGGTACCggtagaaaaa <mark>ATGAGTAAAGGAGAAGAACTTTTCACTGGAGTTGTCCCAATTCTTGTTGAATTAGATGGTGAT</mark>
901	GTTAATGGGCACAAATTTTCTGTCAGTGGAGAGGGTGAAGGTGATGCAACATACGGAAAACTTACCCTTAAATTTATTT
1001	TTCCATGG gtaagtttaaacatatatatactaactaaccctgattatttaaattttcag CCAACACTTGTCACTACTTTCTGTTATGGTGTTCAATGCTT
1101	CTCGAGATACCCAGATCATATGAAACGGCATGACTTTTTCAAGAGTGCCATGCCCGAAGGTTATGTACAGGAAAGAACTATATTTTTCAAAGATGACGGG
1201	<mark>AACTACAAGACAC</mark> gtaagtttaaacagttcggtactaactaaccatactttaaattttaag <mark>GTGCTGAAGTCAAGTTTGAAGGTGATACCCTTGTTA</mark>
1301	atagaatcgagttaaaaggtattgattttaaagaagatggaaacattcttggacacaaattggaatacaactataactcacacaatgtatacatcatggc
1401	<mark>AGACAAACAAAAGAATGGAATCAAAGTT</mark> gtaagtttaaacatgattttactaactaatctgatttaaattttcag <mark>AACTTCAAAATTAGACACAAC</mark>
1501	attgaagatggaagcgttcaactagccgatcattatcaacaaaatactccaattggcgatggccctgtccttttaccagacaaccattacctgtccacac
1601	AATCTGCCCTTTCGAAAGATCCCAACGAAAAGAGAGACCACATGGTCCTTCTTGAGTTTGTAACAGCTGCTGGGATTACACATGGCATGGATGAACTATA
1701	CAAAATAGCATTCGTAGAATTCGCTAGCCGGCCATACAAGTAATCC <mark>GGATGATCGACGCCAACGTCGTTGAATTTTCAAATTTTAAATACTGAATATTTGT</mark>
1801	${\tt TTTTTTCCTATTATTTATTTATTCTCTTTGTGTTTTTTTT$
1901	<pre>teccaattegtatteegeteeteatetgaacacaatgtgeaagtttatttatettetegettteatttea</pre>
2001	gggaaacacacaaaaggatgatggaaatgaaataaggacacacaatatgcaacaacattcaattcagaaatatggaggaaggtttaaaagaaaacataaa
2101	aatatatagaggaggaaggaaaactagtaaaaataagcaaagaaattaggcgaacgatgAGAATTGTCCTCGCTTGGGCCCAAAAGGCCTTTTTTTTTT
2201	ggcgccgacgtcaggtggcacttttcgggggaaatgtgcgcggaacccctatttgtttatttttttt
2301	TAACCCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCCGTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTGCCTTC
2401	CTGTTTTTGCTCACCCAGAAACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGGGGTTACATCGAACTGGATCTCAACAGCGGTAA
2501	GATCCTTGAGAGTTTTCGCCCCGAAGAACGTTTTCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTGACGCCGGGCAA
2601	GAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGG
2701	
2801	Gegega trate tractice of the construction of t
2901	
3001	Getter G
3101	
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3701	
3801	
3901	
4001	TACTGCCTTTTGCTCCCCATGTTCTTTCCTGCGTTATCCCCTGATTCTGCGGTAACCGTATTACCGCCTTTGAGTGAG
4101	GAACGACCGAGCGAGCGAGCGAGCGAGAGCGGAAGAGCGGCAATACGCAACAACCGCCTCTCCCCGGCGTTGGCCGATTCATTAATGCAGC
4201	GCCACGACAGGTTTCCCGACTGGAAAGCGGGCAGTGAGCGCAACGCAATTAATGTGAGTTAGCTCACTCA
4301	
	TCCGGCTCGTATGTTGTGGGAATTGTGAGCGGATAACAATTTCACA <mark>CAGGAAACAGCTATGACCATG</mark> ATTACGCCAAGCT <mark>gtaagtttaaacatgatct</mark>

**Supplementary Note 4** | Sequence of the effector construct pHW394 (pHW394) *15xUAS::Δpes-10::gfp::let-858* 3'UTR is available from Addgene (<u>www.addgene.org</u>; plasmid no. 85584).The color scheme here matches the vector map of pHW394 shown in Supplementary Note 3.

Supplementary Note 5 | Information for extrachromosomal arrays and integrants

The co-injection markers used include *KP708 (Pttx-3::rfp), KP1369 (Pmyo-2::nls::mCherry), KP1106 (Pmyo-2::nls::gfp), unc-119*(+) rescue plasmid, *Pofm-1::rfp and Punc-122::gfp*.

All initial descriptions of extrachromosomal arrays (**syEx#####**) and integrants (**syIs#####**) are highlighted for convenience. All integrants were generated by X-ray irradiation.

A full list of the available integrated drivers and effectors are listed in Supplementary Table 1.

**syEx1452** [15xUAS::Δpes-10::gfp::unc-54 3'UTR, 25ng/μL; Pttx-3::rfp, 40ng/μL; pBlueScript, 35 ng/μL], injected into N2, used to generate **syIs300** and **syIs302**.

**syEx1431** and **syEx1432** [*Pmyo-2::GAL4*<sub>SC</sub>::VP16::unc-54 3'UTR, 10ng/μL; unc-119(+), 50ng/μL; pBlueScript, 40 ng/μL], injected into the strain unc-119(ed3); syIs300.

**syEx1433** and **syEx1434** [*Pmyo-2::GAL4<sub>SC</sub>::VP64::unc-54 3'UTR, 10ng/μL; unc-119(+), 50ng/μL; pBlueScript, 40 ng/μL*], injected into the strain *unc-119(ed3); syIs300*.

**syEx1435** and **syEx1436** [*Pmyo-2::GAL4<sub>SK</sub>::VP64::unc-54 3'UTR; 10ng/μL, unc-119(+), 50ng/μL; pBlueScript, 40 ng/μL*], injected into the strain *unc-119(ed3); syIs300*.

**syEx1437** and **syEx1438** [*Pmyo-2::gfp::unc-54 3'UTR, 10ng/μL; unc-119(+), 50ng/μL; pBlueScript, 40 ng/μL*], injected into the strain *unc-119(ed3)*.

**syEx1448** and **syEx1449** [Pnlp-40::GAL4<sub>SK</sub>::VP64::unc-54 3'UTR, 10ng/μL; Pmyo-2::nls::mCherry, 10ng/μL; pBlueScript, 80ng/μL], injected into syIs302. syEx1449 was used to generate **syIs318**, **syIs319** and **syIs320** as intestine drivers.

**syEx1450** and **syEx1451** [*Pmyo-3::GAL4<sub>SK</sub>::VP64::unc-54 3'UTR, 10ng/μL; Pmyo-2::nls::mCherry, 10ng/μL; pBlueScript, 80ng/μL],* injected into *syIs302. syEx1451* was used to generate **syIs321** as the body wall muscle driver.

**syEx1471** [*Punc-47::GAL4<sub>SK</sub>::VP64::unc-54 3'UTR, 60ng/µL; Pofm-1::rfp, 40ng/µL*], syEx1471 was used to generate **syIs322**, **syIs323**, **syIs324** and **syIs325**, as GABAergic neuron drivers (These GABAergic drivers were weak, we suggest using drivers built in the pPD117.01 backbone with the *let-858 3'UTR*).

**syEx1475**, **syEx1476**, and **syEx1477** [5xUAS::Δpes-10::gfp::unc-54 3'UTR, 25ng/µL; unc-119(+), 50ng/µL; pBlueScript, 25 ng/µL], injected into the strain unc-119(ed3); syIs301.

**syEx1478** and **syEx1479** [10xUAS::Δpes-10::gfp::unc-54 3'UTR, 25ng/µL; unc-119(+), 50ng/µL; pBlueScript, 25 ng/µL], injected into the strain unc-119(ed3); syIs301.

**syEx1480** and **syEx1481** [15xUAS::Δpes-10::gfp::unc-54 3'UTR, 25ng/µL; unc-119(+), 50ng/µL; pBlueScript, 25 ng/µL ], injected into the strain unc-119(ed3); syIs301.

**syEx1482** and **syEx1483** [20xUAS::Δpes-10::gfp::unc-54 3'UTR, 25ng/µL; unc-119(+), 50ng/µL; pBlueScript, 25 ng/µL], injected into the strain unc-119(ed3); syIs301.

**syEx1443** and **syEx1444** [15xUAS::Δpes-10::aex-2(+) cDNA::unc-54 3'UTR, 25ng/µL; Pmyo-2::nls::gfp, 10ng/µL; pBlueScript, 65 ng/µL], injected into the strain aex-2(sa3).

**syEx1433** and **syEx1447** [Prab-3::GAL4<sub>SK</sub>::VP64::let-858 3'UTR, 10 ng/μL; Pofm-1::rfp, 40ng/μL; pBlueScript, 50 ng/μL], injected into N2. syEx1447 was used to generate **syIs334**, **syIs335** and **syIs336** as pan-neuronal driver lines.

**syEx1430**[*Pmyo-2::GAL4*<sub>SC</sub>::VP64::unc-54 3'UTR; 10ng/μL, Pofm-1::rfp 40ng/μL; 1kb DNA ladder(NEB), 150 ng/μL], also used to generate the **syIs301** as the pharyngeal muscle driver.

**syEx1488** [15xUAS::Δpes-10::gfp::let-858 3'UTR, 25 ng/µL; Pttx-3::rfp, 50 ng/µL; 1 kb ladder (NEB), 125 ng/µL], injected into N2, used to generate **syIs337** and **syIs390** for 15xUAS::gfp::let-858 3'UTR effector lines.

**syEx1484** [Punc-17:: GAL4<sub>SK</sub>::VP64::let-858 3'UTR, 25 ng/μL; Punc-17::mCherry, 25 ng/μL; unc-119(+), 50 ng/μL], injected into the strain syls390; unc-119(ed3).

**syEx1485** [Punc-47::GAL4<sub>SK</sub>::VP64::let-858 3'UTR, 25 ng/μL; Punc-47:: mCherry, 25 ng/μL; unc-119(+), 50 ng/μL], injected into the strain syls390; unc-119(ed3).

**syEx1486** [Peat-4::GAL4<sub>SK</sub>::VP64::let-858 3'UTR, 25 ng/µL; Peat-4:: mCherry, 25 ng/µL; unc-119(+), 50 ng/µL], injected into the strain syIs390; unc-119(ed3).

**syEx1460** [15xUAS::Δpes-10::hChR2(H134R)::eyfp::let-858 3'UTR, 25ng/µL; Pttx-3::rfp, 40ng/µL; pBlueScript, 35 ng/µL], injected into N2, used to generate **syIs340**, **syIs341** and **syIs342** for 15xUAS::hChR2(H134R)::eyfp::let-858 3'UTR effector lines.

**syEx1487** [*Punc-47::GAL4*<sub>SK</sub>::VP64::let-858 3'UTR, 25 ng/μL; Pofm-1::rfp, 40 ng/μL; 1 kb ladder (NEB), 35 ng/μL], injected into the strain *syls341*.

#### Supplementary Note 6 | Strains used in the study

For detailed information about arrays and integrants, see **Supplementary Note 5** and **Supplementary Table 1**.

Wild type N2

PS6041 unc-119(ed3) III

### Figure 1b, 1c and Supplementary Figure 1a:

PS6843 syls300 V PS6932 unc-119(ed3); syls300 PS6900 syEx1431; unc-119; syls300 PS6901 syEx1432; unc-119(ed3); syls300 PS6902 syEx1433; unc-119(ed3); syls300 PS6903 syEx1434; unc-119(ed3); syls300

### Figure 1d, 1e and Supplementary Figure 1b

PS6844 syls301 V PS6964 unc-119(ed3); syls301 PS7007 syEx1475; unc-119(ed3); syls301 PS7008 syEx1476; unc-119(ed3); syls301 PS7009 syEx1477; unc-119(ed3); syls301 PS7010 syEx1478; unc-119(ed3); syls301 PS7013 syEx1480; unc-119(ed3); syls301 PS7014 syEx1482; unc-119(ed3); syls301 PS7015 syEx1483; unc-119(ed3); syls301

#### Figure 2 and Supplementary Figure 2:

PS6902 syEx1433; unc-119(ed3); syIs300 PS6903 syEx1434; unc-119(ed3); syIs300 PS6904 syEx1435; unc-119(ed3); syIs300 PS6905 syEx1436; unc-119(ed3); syIs300 PS6906 syEx1437; unc-119(ed3) PS6907 syEx1438; unc-119(ed3)

## Figure 3a-3f:

PS6933 syls318 syls302 III PS7067 syls321; syls300 PS6987 syls337; syls334 PS7149 syls390 PS7184 syls390; unc-119(ed3) PS7018 syEx1484; syls390; unc-119(ed3) PS7019 syEx1485; syIs390; unc-119(ed3) PS7020 syEx1486; syIs390; unc-119(ed3)

Figure 3h:

JT3 aex-2(sa3) X PS6975 syEx1443; aex-2(sa3) PS6976 syEx1444; aex-2(sa3) PS6936 syIs321 PS6935 syIs320 PS6938 syIs323

The exact genotypes used for the quantification of the defecation assay in Figure 3h are (from left to right):

N2

aex-2(sa3) syEx1444; aex-2(sa3) syIs323/+; aex-3(sa3) syEx1444; syIs323/+; aex-2(sa3) syEx1444; syIs321/+; aex-2(sa3) syEx1444; syIs320/+; aex-2(sa3)

Figure 3i and Supplementary Video 1-2:

PS7021 syEx1487; syls341 PS7044 syls341

Supplementary Figure 3 and Supplementary Video 3-5:

PS6872 syls302 III PS6844 syls301 V PS6965 syls301; syls302 PS7186 syls407 PS7154 syls391 IV PS7136 syls378 V PS7190 syls409 X PS7167 syls396 syls337 III PS7192 syls413 IV PS6936 syls321 PS7205 syls424 III PS7199 syls371 PS7201 syls421