

Olfactory chemosensation extends lifespan through TGF- β signaling and UPR activation

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Supplementary Table 1. Survival information for lifespan experiments.

Strain	Treatment	Median lifespan (days)	Control	Median lifespan (days)	P value (Mantel-Cox)
N2	+ 1-undecene	20	- 1-undecene	18	0.0227
N2	+ 1-undecene	20	- 1-undecene	18	0.0357
N2	+ 1-undecene	26	- 1-undecene	21	0.0185
N2	+ 1-undecene	25	- 1-undecene	21	0.0007
N2	+ 1-undecene	25	- 1-undecene	21	<0.0001
N2	+ 1-undecene	23	- 1-undecene	19	<0.0001
<i>xbp-1(zc12)</i>	+ 1-undecene	14	- 1-undecene	14	0.7015
<i>xbp-1(zc12)</i>	+ 1-undecene	18	- 1-undecene	20	<0.0001
<i>xbp-1(zc12)</i>	+ 1-undecene	22	- 1-undecene	22	0.0066
<i>daf-1(m40)</i>	+ 1-undecene	21	- 1-undecene	25	0.9948
<i>daf-1(m40)</i>	+ 1-undecene	21	- 1-undecene	23	0.2990
<i>daf-7(e1372)</i>	+ 1-undecene	21	- 1-undecene	21	0.4488
<i>daf-7(e1372)</i>	+ 1-undecene	21	- 1-undecene	23	0.7628
N2	+ pyrrole	24	- pyrrole	19	<0.0001
N2	+ pyrrole	22	- pyrrole	22	0.0046
<i>xbp-1(zc12)</i>	+ pyrrole	17	- pyrrole	17	0.0071
<i>xbp-1(zc12)</i>	+ pyrrole	19	- pyrrole	17	<0.0001
<i>xbp-1(zc12)</i>	+ pyrrole	22	- pyrrole	17	<0.0001

10 **Supplementary Table 2. List of *C. elegans* strains used.**

<i>Caenorhabditis elegans</i> strains used		
Wild type	CGC	N2
<i>zcls4 [hsp-4::GFP] V</i>	CGC	SJ4005
<i>ire-1(zc14) II; zcls4 [hsp-4::GFP] V</i>	CGC	SJ30
<i>zls356 [daf-16p::daf-16a/b::GFP + rol-6(su1006)] IV</i>	CGC	TJ356
<i>rmls132 [unc-54p::Q35::YFP]</i>	CGC	AM140
<i>dvlN70 [hsp-16-2p::GFP; rol-6]</i>	CGC	CL2070
<i>daf-11(m47) V</i>	CGC	DR47
<i>daf-1 (m40) IV</i>	CGC	DR40
<i>daf-7 (e1372) III</i>	CGC	CB1372
<i>rmls110 [F25B3.3p::Q40::YFP]</i>	Imanikia et al., 2019	AGD1397
<i>uthls393 [vha-6p::Q40::YFP+rol-6(su1006)]</i>	Imanikia et al., 2019	AGD1395
<i>rmls1s9 [daf-7p::xbp-1s::unc-54 3'UTR, myo-3p::mKate]; zcls4 [hsp-4p::GFP] V</i>	Özbey et al., 2020	RCT206
<i>rmls1s8 [xbp-1p::xbp-1::GFP]</i>	Özbey et al., 2020	RCT21
<i>rmls1s7 [tdc-1p::mKate2::let-858 3'UTR, cc::GFP]; rmls1s8 [xbp-1p::xbp-1::GFP]</i>	Özbey et al., 2020	RCT192
<i>xbp-1 (zc12) III</i>	Taylor & Dillin, 2013	AGD1049
<i>xbp-1 (zc12) III; zcls4 [hsp-4::GFP] V</i>	Taylor & Dillin, 2013	AGD972
<i>unc-13 (e450) I; zcls4 [hsp-4::GFP] V</i>	Taylor & Dillin, 2013	AGD1137

drcSI7 [<i>daf-7p::Venus</i>]	Tullet Lab, U. of Kent	JMT50
<i>daf-1 (m40)</i> IV; <i>zcls4 [hsp-4::GFP]</i> V; <i>ftEx205[tdc-1p::daf-1:gfp + odr-1:dsRED]</i>	This study	RCT378
<i>pmk-1 (k25)</i> IV; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT379
<i>kgb-1 (um3)</i> IV; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT380
<i>lim-4 (ky403)</i> ; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT381
<i>oyls84 [gpa-4p::TU#813 + gcy-27p::TU#814 + gcy-27p::GFP + unc-122p::DsRed]; zcls4 [hsp-4::GFP]</i> V	This study	RCT237
<i>daf-7 (e1372)</i> III; <i>rmls132 [unc-54p::Q35::YFP]</i>	This study	RCT382
<i>zip-2 (ok3730)</i> III; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT369
<i>unc-31 (e928)</i> IV; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT370
<i>tdc-1 (n3419)</i> II; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT66
<i>tph-1 (mg280)</i> II; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT371
<i>cat-2 (e1112)</i> II; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT372
<i>unc-25 (e156)</i> III; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT373
<i>eat-4 (ky5)</i> III; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT374
<i>nsIs180 [hlh-17p::recCaspase-3, unc-122p::GFP]; zcls4 [hsp-4::GFP]</i> V	This study	RCT375
<i>cho-1 (tm373)</i> IV; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT189
<i>alh-11 (lj118)</i> III; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT376
<i>daf-1 (m40)</i> IV; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT67
<i>daf-7 (e1372)</i> III; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT68
<i>odr-3 (rms31)</i> V; <i>zcls4 [hsp-4::GFP]</i> V	This study	RCT377

Supplementary Table 3. List of qPCR primers used in this work.

	SOURCE	IDENTIFIER
qPCR Primers		
GTTCCCGTGTTCATCACTCAT	Sigma	<i>pmp-3</i> F
ACACCGTCGAGAAGCTGTAGA	Sigma	<i>pmp-3</i> R
CTGCTGGACAGGAAGATTACG	Sigma	<i>cdc-42</i> F
CTCGGACATTCTCGAATGAAG	Sigma	<i>cdc-42</i> R
GTCGCTTCAAATCAGTTCAGC	Sigma	Y45F10D.4 F
GTTCTTGTCAAGTGATCCGACA	Sigma	Y45F10D.4 R
CGTGCCTTGAATCAGCAGTG	Sigma	<i>xbp-1_s</i> F
CGAGGTGTCCATCTTCTTGTT	Sigma	<i>xbp-1_s</i> R
CAGATGAAAACTCAAATGCC	Sigma	<i>hsp-4</i> F
GGTTGCTTCCGAGCCACTCAA	Sigma	<i>hsp-4</i> R
CTTTATTGCTCCCTAACCGT	Sigma	Y41C4A.11 F
CTGCTCCATTCTCCACTCGTT	Sigma	Y41C4A.11 R
TTCTACACCGCCAAAGATCC	Sigma	<i>daf-7</i> F
CTGTGAGTGTGGCCTGAAGA	Sigma	<i>daf-7</i> R

Supplementary Table 4. List of CRISPR primers and oligonucleotides used in this work.

	SOURCE	IDENTIFIER
CRISPR Oligos		
GCTACCATAGGCACCACGAG	IDT	<i>dpy-10</i> cRNA
AAAATT CGGAAGGT AAC GCG; C TGC ATT T ACC GTT GG AAAA	IDT	<i>odr-3</i> cRNAs
CACTTGA ACTT CAATACGGCAAGATGAG AATGACTGGAAACCGTACCGCATGCGG TGCCTATGGTAGCGGAGCTTCACATGGC TTCAGACCAACAGCCTAT	IDT	<i>dpy-10</i> Repair Template
caattactcatagatcattgttttttagATATGGGCTCA TGCCAGAGCAATGAAAATT CGGAAGGT A ACAAATGGACAAAAAGAAGCAGAAAAGG CAATAGTTATGAAAGTACAGGAAAATGGAGAAG AAGGAGAAGCACTGACAGAAGAAGTTCGAAA GCAATTCAATCGTTATGGCAGATCCTGGCGTGAAGAA	IDT	<i>odr-3</i> Repair Template