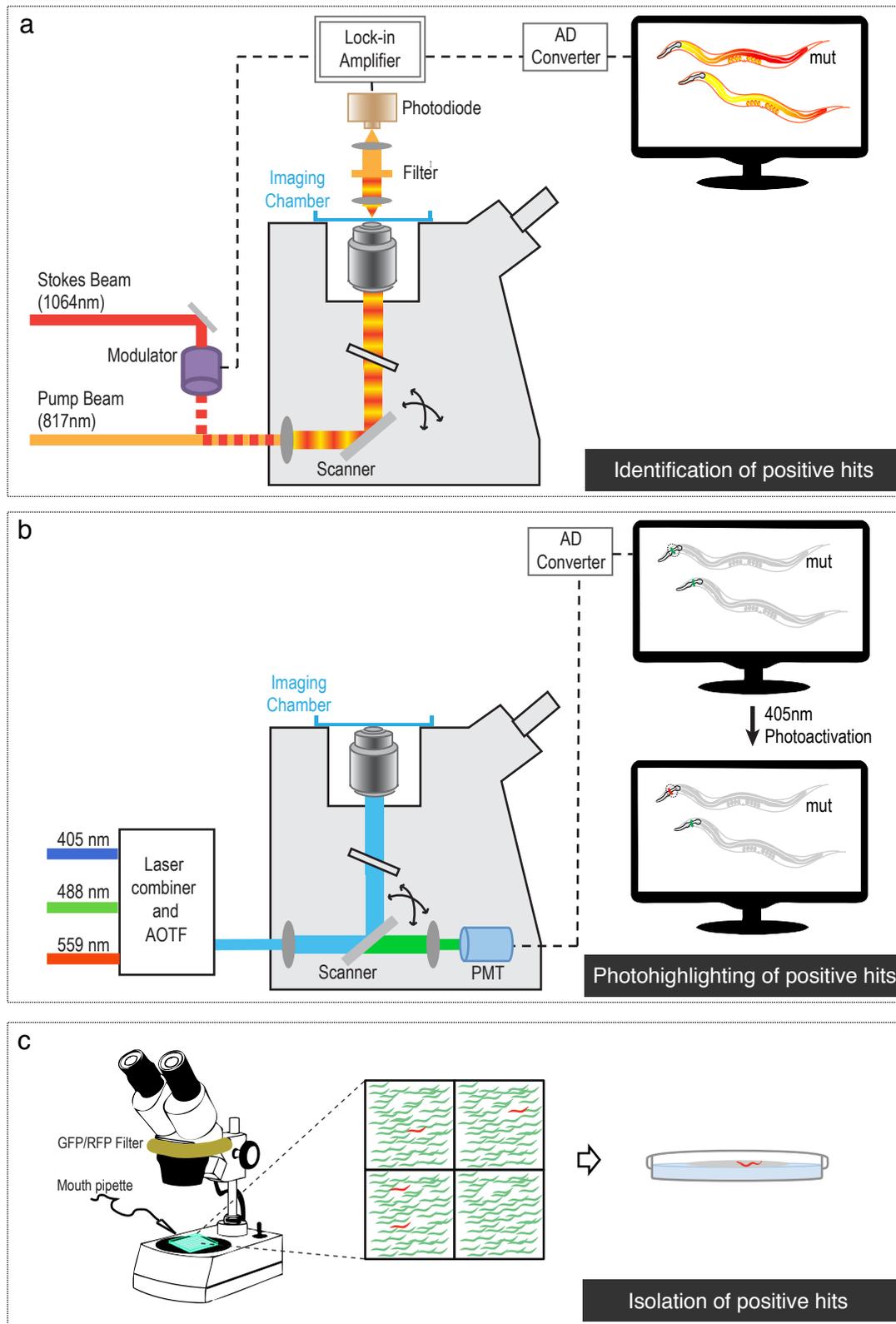
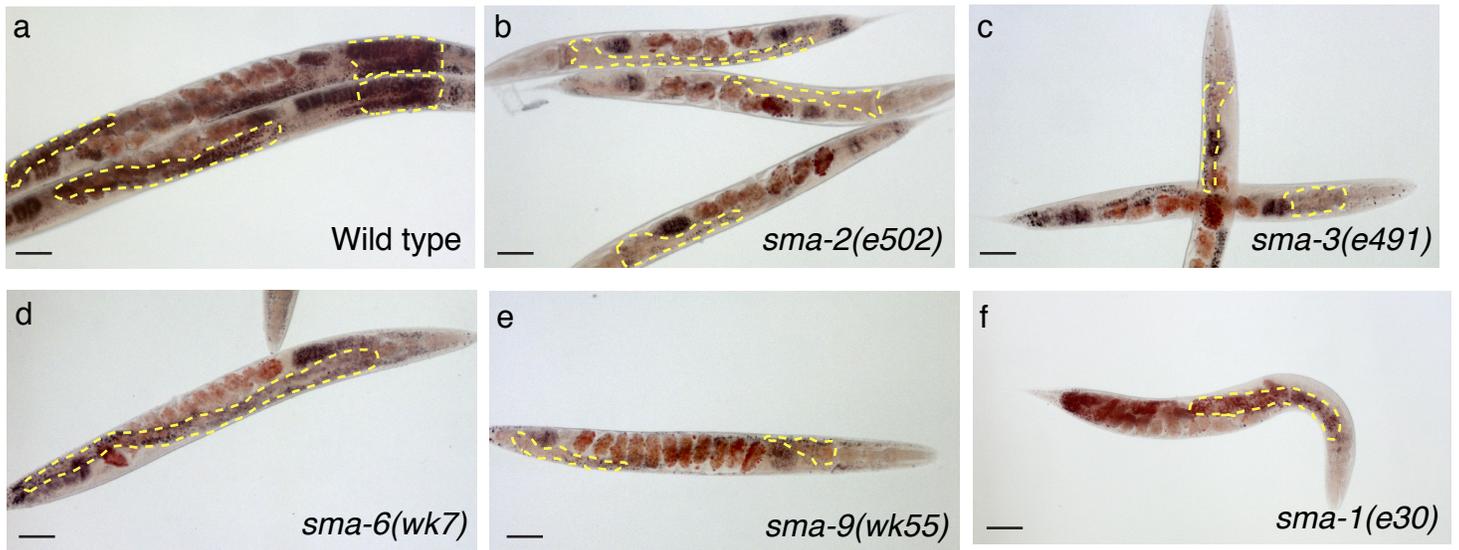


Supplementary Figure 1. Design of imaging chambers for 3D printing.



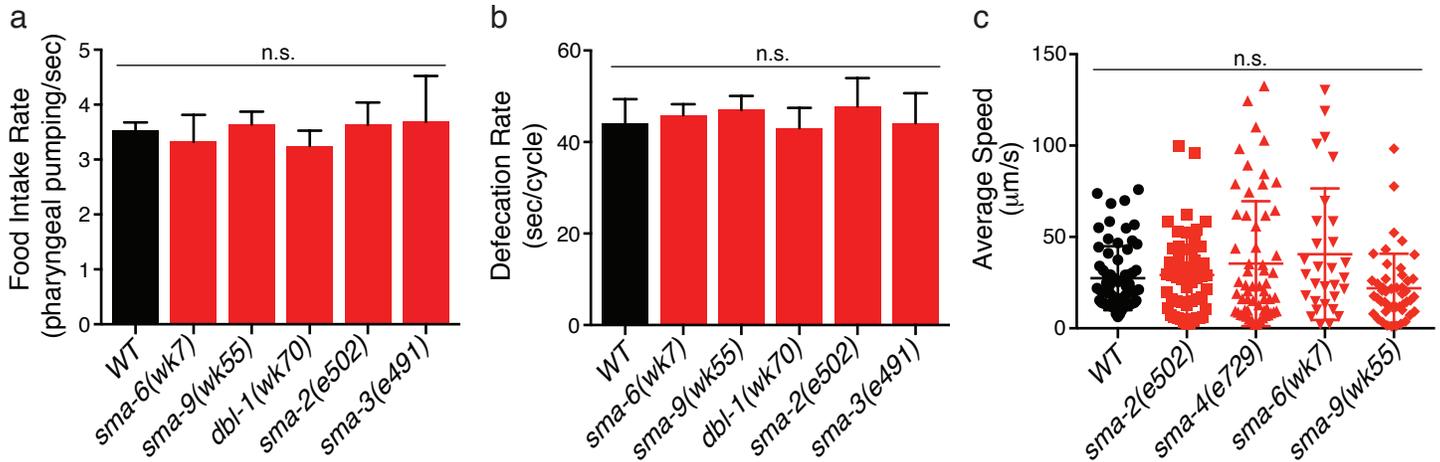
Supplementary Figure 2. Experimental scheme of high-throughput genetic screening.

(a) Diagram of the optical components for SRS microscopy built on a laser scanning confocal microscope. (b) Diagram of the conventional confocal imaging components on the same microscope used for photo-highlighting. PMT, photomultiplier tube. AOTF, acousto-optic tunable filters. (c) Recovery of photo-highlighted positive hits under a fluorescent dissection scope.



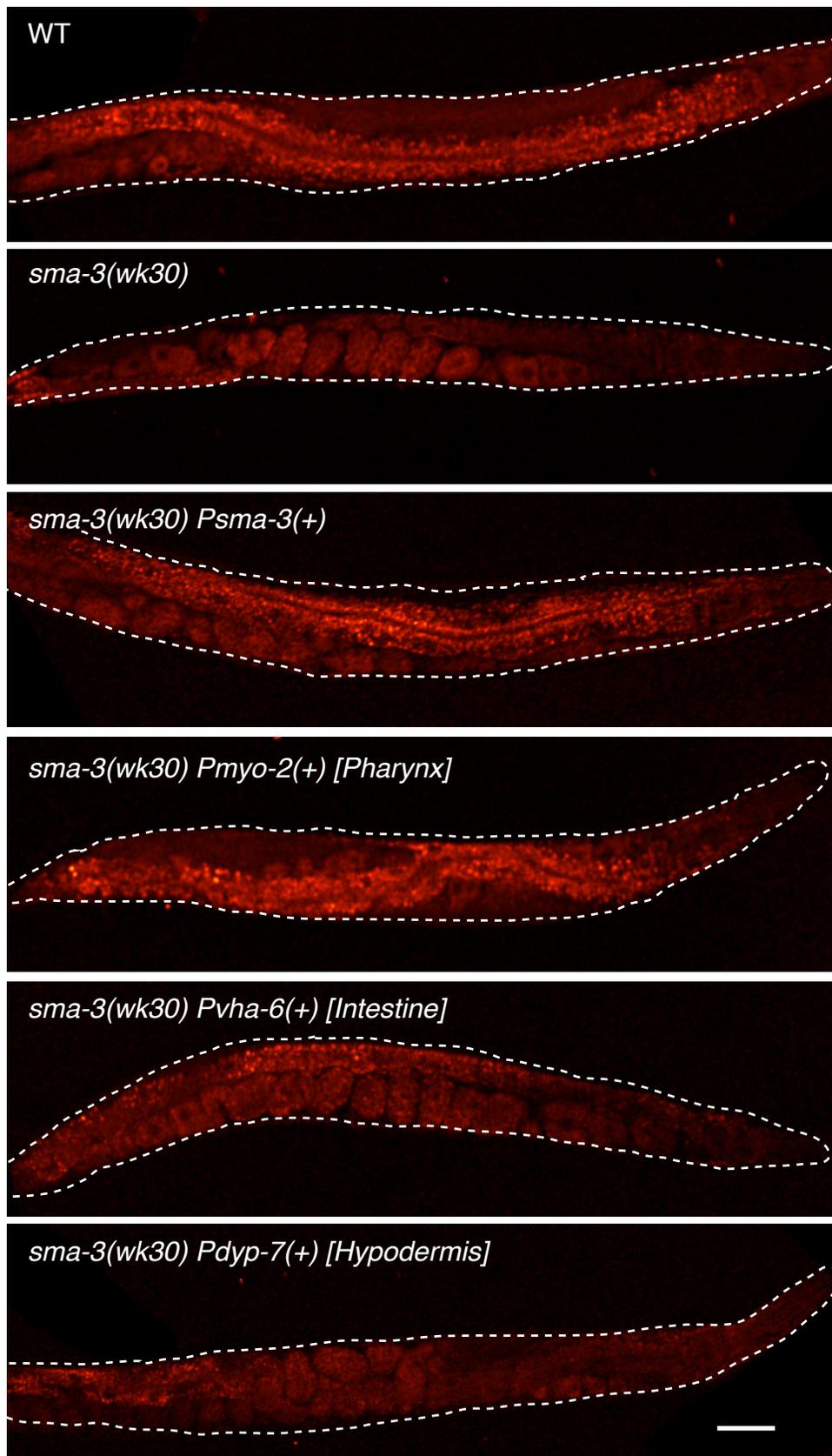
Supplementary Figure 3. Oil Red O staining images of *sma* mutants.

Representative images show lipid content levels in 1-day-old adult worms stained with Oil Red O. Yellow dashed lines indicate intestine areas. Scale bar=50 μ m. Compared to wild type controls (a), lipid content levels are reduced in the *sma* mutants of the BMP signaling pathway (b-e), but not in the non-BMP *sma* mutant (f).



Supplementary Figure 4. Physiological measurement of *sma*/BMP pathway mutants.

Compared to wild type (WT) worms, the *sma* mutants show similar food intake rate (**a**, $n=12$), defecation interval (**b**, $n=10$), and locomotion activity (**c**, $n=66$ for WT, $n=53$ for *sma-2*, $n=59$ for *sma-4*, $n=30$ for *sma-6*, $n=47$ for *sma-9*). Error bars represent means \pm s.d., n.s. $p>0.05$ by one-way ANOVA.



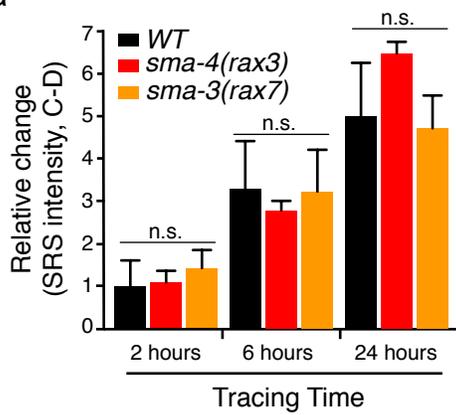
High  Low

SRS intensity at 2845 cm^{-1}

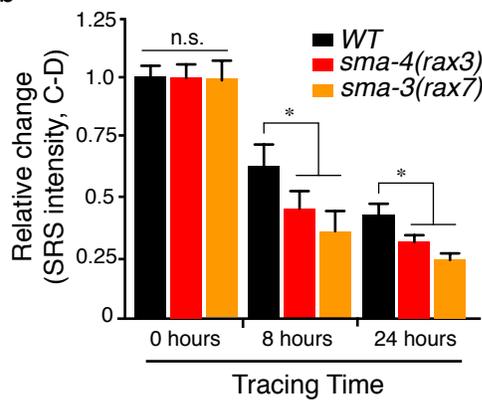
Supplementary Figure 5. Representative SRS images show lipid contents in wild-type, *sma-3* mutant, and *sma-3* tissue specific rescue strains.

Quantification results are shown in Fig. 3e. Dashed lines indicate the boundary of worms. Scale bar=50 μm .

a



b

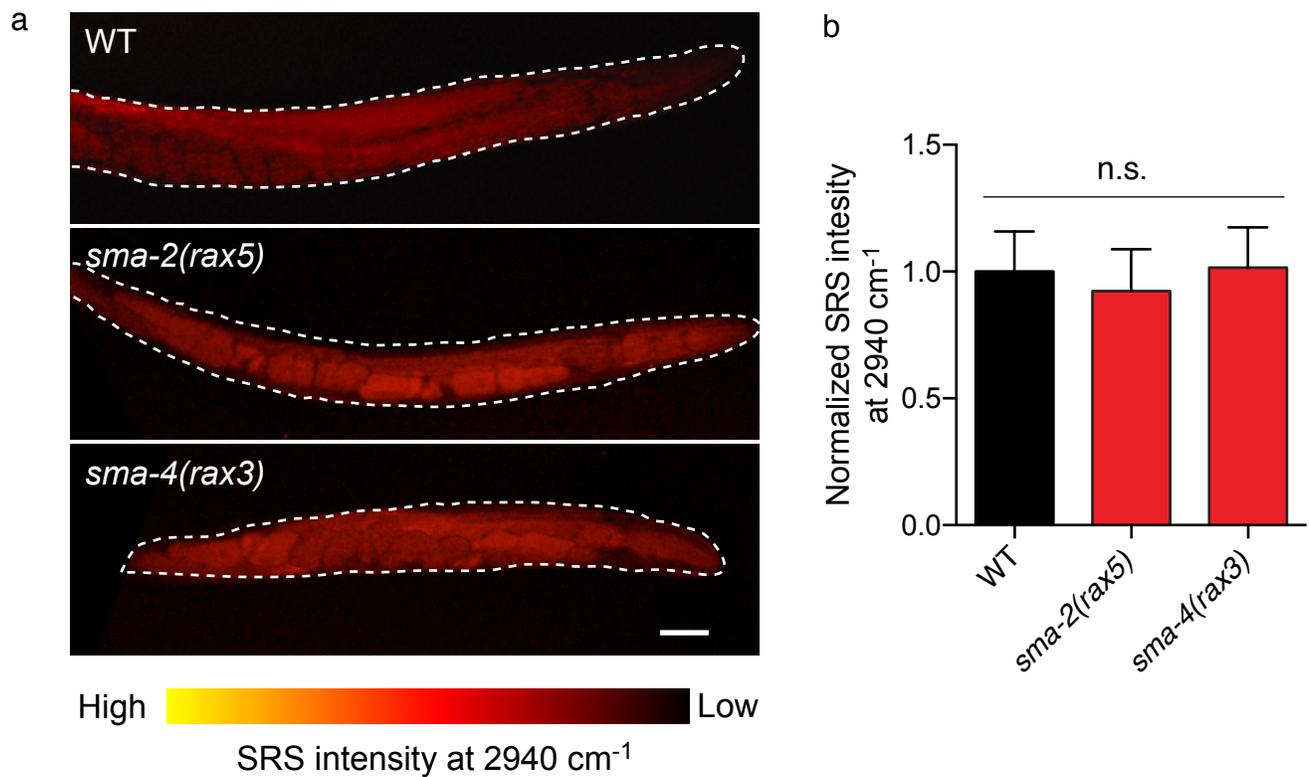


Supplementary Figure 6. Another iSRS replicate results reveal the regulation of lipid mobilization by BMP signaling.

(a) The BMP/*sma* mutants do not affect lipid synthesis, n.s. $p > 0.05$, $n = 5$.

(b) The BMP/*sma* mutants accelerate lipid mobilization, n.s. $p > 0.05$, * $p < 0.05$, $n = 5$.

In all bar charts, data represent mean \pm s.d., p value is determined by a Student's t-test (unpaired, two-tailed).



Supplementary Figure 7. SRS imaging at 2940 cm⁻¹ reveals unchanged protein levels in *sma* mutants.

(a) Example SRS images at 2940 cm⁻¹. Dashed lines indicate the boundary of animals. Scale bar=50 μ m.

(b) Quantification of SRS signal intensity at 2940 cm⁻¹. n.s. $p > 0.05$ Student's *t*-test, n=43 for WT, n=25 for *sma-2*, n=44 for *sma-4*.

Supplementary Table 1. qPCR primers

<i>acs-2</i> FWD	TTCGACCGGATGAGCCAGTAAACA
<i>acs-2</i> REV	GTTGTTGTTTCAGCCCGAAATGGGT
<i>cpt-4</i> FWD	CAGAAACATCGCCAGTCTCA
<i>cpt-4</i> REV	TCCATCCACTGACATCACAAA
<i>acs-18</i> FWD	TGTGGCTCCAGATCTCACAG
<i>acs-18</i> REV	GATCTGGGACAACGATTGCT
<i>act-1</i> FWD	GCTGGACGTGATCTTACTGATTACC
<i>act-1</i> REV	GTAGCAGAGCTTCTCCTTGATGTC
<i>rpl-32</i> FWD	AGGGAATTGATAACCGTGTCCGCA
<i>rpl-32</i> REV	TGTAGGACTGCATGAGGAGCATGT