

Chemical reprogramming of *Caenorhabditis elegans* germ cell fate

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Small molecules can control cell fate *in vivo* and may allow directed induction of desired cell types, providing an attractive alternative to transplant-based approaches in regenerative medicine. We have chemically induced functional oocytes in *Caenorhabditis elegans* adults that otherwise produced only sperm. These findings suggest that chemical approaches to therapeutic cell reprogramming may be feasible and provide a powerful platform for analyzing molecular mechanisms of *in vivo* cell reprogramming.

Cell fate reprogramming within organisms has the potential to revolutionize regenerative medicine¹. By reprogramming endogenous cells into a desired cell type, major difficulties associated with transplant-based approaches (for example, cell isolation and immune rejection) may be minimized. So far, most reprogramming has been limited to *in vitro* applications, although a recent breakthrough reported reprogramming of cells in mice with virally delivered transcription factors². Small molecules can modulate cell fates

*in vivo*³, and chemical induction of a desired cell type could provide an attractive alternative to transplant- or viral-based cell replacement therapies. Notably, chemical modulation of prostaglandin E2 synthesis can regulate hematopoietic stem cell proliferation⁴, and retinoid-based “differentiation therapy” is used clinically to treat M3 acute myeloid leukemia (AML)⁵. However, to our knowledge, the chemical induction of functional cells that were absent before treatment has not been demonstrated in any organism. The ability to induce missing cells could have profound implications for the treatment of diseases of cellular deficiency.

The germline of the nematode *C. elegans* provides a tractable system for studying cellular reprogramming⁶. Normally, *C. elegans* hermaphrodites make sperm as larvae and oocytes as adults, yielding an adult gonad that uses stored sperm to fertilize its developing oocytes (Fig. 1a). Genetic screens for mutant hermaphrodites that make only sperm (masculinized germlines) or only oocytes (feminized germlines) have identified key regulators

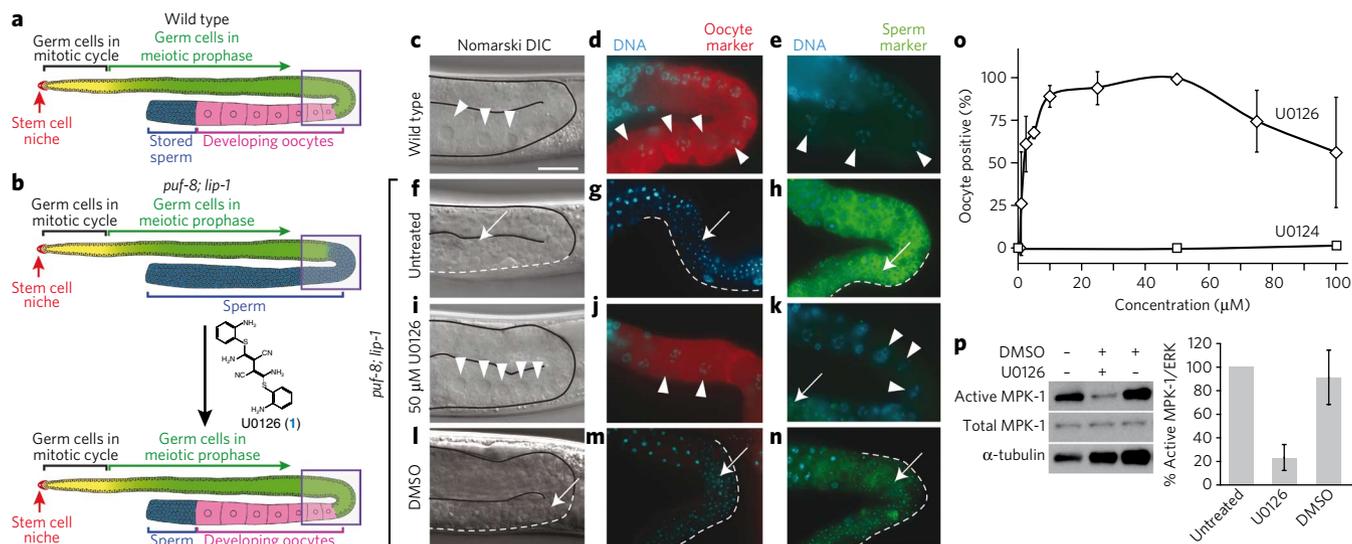


Figure 1 | Small molecule-mediated reprogramming of sperm/oocyte fate. (a) Organization of the wild-type hermaphrodite gonad. Boxed area indicates the region shown in panels. (b) The *puf-8; lip-1* gonad produces only sperm, but oocytes can be induced chemically. (c–n) Genotypes and treatments are labeled in the figure. Oocytes are indicated with arrowheads. The extent of sperm is indicated with white dashed lines and representative single sperm with arrows. Scale bar, 10 μm. (c, f, i, l) Differential interference contrast (DIC) images reveal morphology of large granular oocytes (c, i) or small condensed sperm (f, l). (d, g, j, m) Germlines stained for DNA using 4',6-diamidino-2-phenylindole (DAPI) (blue) and the RME-2 oocyte marker (red). (e, h, k, n) Germlines stained for DNA using DAPI (blue) and the SP56 sperm marker (green). (o) Dose-response analysis for oogenesis induction by U0126 (◇) and its inactive analog U0124 (□). Data are presented as mean ± s.d., n = 3–12 independent experiments with 3–149 (average 13) animals each. (p) Left, semiquantitative western blot analysis of activated MPK-1/ERK (top panel), total MPK-1/ERK (middle panel) and α-tubulin (bottom panel) in untreated, U0126-treated and DMSO-treated animals. Right, graph generated with active MPK-1/ERK signals normalized to α-tubulin (n = 3) and compared to untreated signal (set as 100% active MPK-1/ERK).

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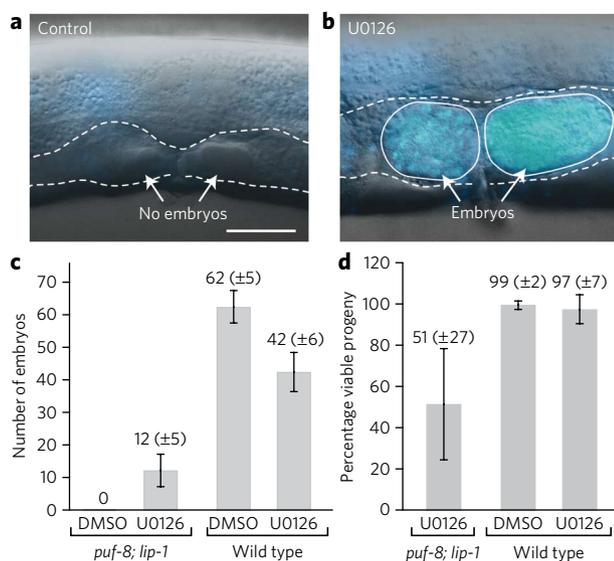


Figure 2 | Small molecule-induced oocytes are functional. (a,b) Whole animals were imaged by DIC and DAPI (blue) staining after treatment with control (DMSO) or U0126. (a) DMSO-treated mutant lacking embryos (dashed outline indicating empty uterus). (b) U0126-treated mutant with embryos containing numerous DAPI-stained nuclei. (c) Number of embryos produced by mutants and wild type treated with DMSO or U0126. (d) Percentage of embryos that developed to adulthood after parents were treated with DMSO or U0126. Approximately half of the embryos from chemically induced oocytes developed to adulthood. Data are presented as mean \pm s.d. Scale bar, 10 μ m.

of sperm/oocyte cell fate^{6,7}. Indeed, genetic manipulation of these fate regulators revealed that germ cells continuously make the sperm/oocyte decision, even in adults⁸. Such continued cell fate specification in adults is typical of many vertebrate tissues, including blood and intestinal epithelium. In *C. elegans*, one regulator that promotes the sperm fate is MPK-1, the ERK-1/2 MAPK homolog^{9,10} and terminal kinase of canonical Ras-ERK signaling. Yet MPK-1/ERK hyperactivation, which occurs upon loss of the dual-specificity phosphatase LIP-1, fails to generate excess sperm¹¹. However, we have found that a double mutant, lacking *lip-1* and a second regulator (*puf-8*), produces only sperm (Fig. 1b). The *puf-8* gene encodes a PUF (for Pumilio and FBF) RNA-binding protein that affects the sperm/oocyte decision redundantly with another PUF protein, FBF-1 (ref. 12). Because PUF-8 targets are largely unknown, the mechanism underlying the interaction between PUF-8 and Ras-ERK signaling remains unclear. However, it is likely to involve parallel controls that impinge on the sperm/oocyte decision redundantly.

We scored sperm and oocyte production in wild-type and *puf-8(q725); lip-1(zh15)* (hereafter referred to as “*puf-8; lip-1*” or “mutant”) adult germlines by morphology and staining for molecular markers (Fig. 1c–n). Wild-type oocytes are large cells with granular cytoplasm, and their nuclei have progressed to diplotene or diakinesis of meiotic prophase¹³ (Fig. 1c). Moreover, oocytes stained with the oocyte marker RME-2 (Fig. 1d), but not with the sperm-specific marker SP56 (Fig. 1e). By contrast, mutant germlines lacked oocytes and instead made excess sperm. Sperm have compact nuclei (Fig. 1f–h) and fail to stain with oocyte marker (Fig. 1g), but stain strongly with the sperm marker (Fig. 1h). Indeed, all mutant germlines were masculinized (100%, $n = 199$).

We used masculinized *puf-8; lip-1* germlines to develop a small molecule-based method to switch germ cell fates. Specifically, we investigated whether inhibition of Ras-ERK signaling could reinstate oogenesis. To this end, we grew synchronized mutants on

normal growth media until adulthood (to avoid effects on somatic development) and treated them with U0126 (1), a small-molecule MEK1/MEK2 inhibitor that blocks MPK-1/ERK activation^{14,15}. After 24 h of treatment, virtually all mutants made oocytes (99%; $n = 158$). The induced oocytes had oocyte-specific cellular morphology (Fig. 1i), expressed the oocyte marker (Fig. 1j) and were negative for the sperm marker (Fig. 1k). These germlines with induced oocytes also contained sperm (Fig. 1k), so not all germ cells were reprogrammed. Induction of RME-2-positive oocytes was also observed when U0126 was solubilized in a different solvent (EtOH), while vehicle (DMSO or EtOH)-treated control animals remained masculinized as assessed by morphology and markers (Fig. 1l–n and Supplementary Fig. 1).

One interpretation of the induced oocytes in the U0126-treated mutant germlines is that germ cells normally destined for sperm differentiation had been reprogrammed into the oocyte fate. Alternatively, germ cells destined for oogenesis might not be detected in the mutant due to differential proliferation or apoptosis of cells specified as oocytes. To compare proliferation, we determined the percentage of S-phase germ cells (see Supplementary Methods) in spermatogenic and oogenic mutants and found $52 \pm 7.2\%$ ($n = 3$) and $57 \pm 1\%$ ($n = 3$) S-phase cells, respectively. Therefore, sex-specific differences in germline proliferation cannot explain the appearance of U0126-induced oocytes. We then investigated whether mutant germ cells undergo apoptosis instead of oocyte differentiation. To this end, we blocked apoptosis by RNAi of the Apaf-1 homolog *ced-4*, an essential regulator of programmed cell death¹⁶. However, no oocytes were observed in *ced-4* RNAi-treated mutants ($n = 70$). We conclude that U0126 induced oocytes by chemically reprogramming the germline, generating a cell type that was absent in untreated mutants.

We verified that U0126 inhibits Ras-ERK signaling to affect sperm/oocyte fate. U0126 induced oocytes in a dose-dependent manner with an *in vivo* effector concentration for half-maximum response (EC_{50}) of $\sim 2 \mu$ M (Fig. 1o), consistent with the previously reported half-maximal inhibitory concentration (IC_{50}) of $\sim 1 \mu$ M for the effect of U0126 on AP-1 activity in tissue culture¹⁵. Moreover, high U0126 doses (75–100 μ M) caused pachytene arrest, a defect diagnostic for loss of Ras-ERK signaling in *C. elegans* germlines¹⁷ (Fig. 1o). In western blots, U0126 substantially reduced the abundance of activated MPK-1/ERK compared to untreated and DMSO-treated controls (Fig. 1p, top panel), but levels of total MPK-1 were unchanged (Fig. 1p, middle panel). Therefore, U0126's effects on germline sex are likely to be the result of “tuning” ERK activity rather than complete inhibition. Taken together, our results show that U0126 inhibits MPK-1/ERK activity in the *C. elegans* mutant germline and confirm the role of MPK-1/ERK in sperm fate.

To rule out off-target effects of U0126 as the mechanism of reprogramming, we tested an inactive but structurally similar analog, U0124 (2) (Fig. 2a)¹⁴. U0124 failed to induce oogenesis (Fig. 1p and Supplementary Fig. 2b–d) or lower activated MPK-1/ERK (Supplementary Fig. 3 and Supplementary Table 1). Two structurally distinct small-molecule MEK-1/2 inhibitors, PD0325901 (3)¹⁸ and PD098059 (4)^{19,20}, also induced oocytes in the mutants. 50 μ M PD0325901 induced oocytes in 91% ($n = 35$) of the mutants, assayed by cellular morphology and gamete-specific markers (Supplementary Fig. 2e–g and Supplementary Table 1). Consistent with this result, PD0325901 substantially decreased activated MPK-1/ERK (Supplementary Fig. 3). PD098059 induced oocytes in only 5.3% ($n = 75$) at 100 μ M (Supplementary Table 1), consistent with PD098059's poor bioavailability and ~ 100 -fold lower affinity for MEK-1/2 (ref. 14) relative to U0126 (ref. 18). Thus, three structurally distinct inhibitors of Ras-ERK signaling elicited the same response: oocyte induction. These findings strongly suggest that the inhibitors elicit their effects via their common target, Ras-ERK signaling. We conclude that Ras-ERK inhibitors act as a chemical switch to reprogram germ cell fates in *puf-8; lip-1* mutants.

To confirm that MPK-1/ERK inhibition is sufficient for germ cell fate switching, we performed RNAi-mediated knockdown of the germline isoform of *mpk-1*, *mpk-1b*²¹. Mutants treated with *mpk-1b* RNAi made oocytes (99%; $n = 98$), as assessed by morphology and gamete-specific markers (Supplementary Fig. 2h–j). By contrast, all animals treated with RNAi vector made sperm only (100%, $n = 63$), as assessed by morphology and markers (Supplementary Fig. 2k–m). Therefore, inhibitors of Ras-ERK signaling can reprogram mutant germ cells. However, oogenesis was not induced by inhibition of several other major signaling pathways (Supplementary Table 1 and Supplementary Fig. 4). We conclude that germ cell fate switching does not occur by generalized perturbation, but specifically by inhibition of Ras-ERK signaling.

Finally, we investigated whether U0126-induced oocytes were functional gametes. To assess oocyte function, we examined their capacity to be fertilized, to make embryos and to support embryo viability. Control DMSO-treated mutants were sterile and made no embryos (Fig. 2a) ($n = 23$). By contrast, almost all U0126-treated mutants produced embryos (Fig. 2b) (96%, $n = 23$), indicating that the oocytes were functional and that reprogramming was highly efficient. To examine this effect in more detail, we first scored embryo number. U0126-treated mutants made 11.5 ± 4.8 embryos per animal ($n = 8$), whereas DMSO-treated mutants made none ($n = 8$) (Fig. 2c). For comparison, wild-type animals treated with DMSO or U0126 produced 62 ± 5.3 or 41.5 ± 5.7 embryos per animal, respectively ($n = 8$ for each) (Fig. 2c). Therefore, embryo production by U0126-treated mutants reached ~30% of wild-type levels. We next assessed embryo viability. About half the embryos made from chemically induced oocytes developed to adulthood (51%; $n = 92$) (Fig. 2d). Much of their embryonic lethality could be secondary to mutant defects, since untreated *puf-8* and *lip-1* single mutants exhibit partial embryonic lethality^{22,23} and U0126 did not affect wild-type embryo viability. To assay embryonic lethality of the double mutants in the absence of U0126, we genetically induced oocytes by RNAi knockdown of *fog-1*, a terminal sperm fate regulator⁶. The *fog-1* RNAi induced oocytes in all treated animals (100%, $n = 98$), but those oocytes were small and did not support embryonic development. Therefore, U0126 treatment had additional positive effects beyond inducing oogenesis. Among embryos from DMSO- and U0126-treated wild-type animals, 99.6 ± 2.1 ($n = 497$) and 96.6 ± 6.6 ($n = 332$) developed to adulthood, respectively (Fig. 2d). The *puf-8*; *lip-1* progeny arising from chemically induced oocytes were sterile as adults, because they made only sperm; yet oogenesis and fertility could be re-instated in these adults by U0126 treatment. Indeed, we expanded an initial population of homozygous mutants from 17 to ~15,000 animals in six generations by “pulsing” sterile adults with 50 μ M U0126 and allowing their progeny to grow to adulthood before the next pulse. Therefore, many of the U0126-induced oocytes are fully functional gametes.

In summary, we have shown that germ cell fate can be chemically reprogrammed within adult *C. elegans* and that reprogramming can induce a cell type that was absent without treatment. We speculate that the *in vivo* environment contributed significantly to the high efficiency and functionality of the reprogramming. We do not know whether the reprogramming occurs by lineage switching of progenitors or by direct conversion of spermatocytes to oocytes. *C. elegans* germlines have a stem cell reservoir that may provide a pool of cells with labile fate that can be efficiently reprogrammed. Indeed, the time required for reprogramming corresponds roughly to the time required for immature germ cells to enter meiosis and differentiate as sperm or oocyte. Some vertebrate tissues are similarly maintained in adults by a stem cell reservoir (for example, blood), and some human diseases similarly lack differentiated cells while maintaining

a progenitor pool (for example, acute myelogenous leukemias²⁴ and azoospermia²⁵). Such disease cells may represent good targets for therapeutic chemical reprogramming efforts—a notion supported by the success of differentiation therapy in M3 AML⁵. Our results in nematodes strongly support the idea that pharmacological approaches to cell reprogramming will be therapeutically feasible more broadly. In addition, the reprogramming of germ cell fates in *C. elegans* provides a powerful model for analyzing molecular mechanisms of *in vivo* cell reprogramming. Therefore, our findings provide a paradigm that may facilitate pharmacological approaches to therapeutic cellular reprogramming in other organisms.

Received 28 August 2009; accepted 19 October 2009;
published online 20 December 2009

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Acknowledgments

We thank S. Ward (University of Arizona) and B. Grant (Rutgers University) for providing reagents. We are grateful to L. Vanderploeg, A. Steinberg and A. Hellsley-Marchbanks for help preparing the figures and manuscript. We thank the Kimble lab members for helpful discussions. This work was supported by US National Institutes of Health grant RO1 GM069454. J.K. is an investigator with the Howard Hughes Medical Institute.

Author contributions

C.T.M. and M.-H.L. performed the experiments. C.T.M., M.-H.L. and J.K. analyzed the results. C.T.M. and J.K. wrote the manuscript.

Additional information

Supplementary information and chemical compound information is available online at <http://www.nature.com/naturechemicalbiology/>. Reprints and permissions information is available online at <http://npg.nature.com/reprintsandpermissions/>. Correspondence and requests for materials should be addressed to J.K.