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Supplemental information

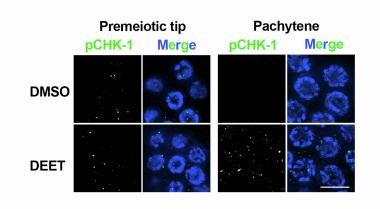
Altered gene expression linked

to germline dysfunction following exposure

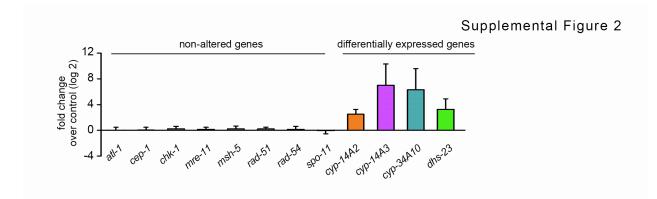
of Caenorhabditis elegans to DEET

Nara Shin, Laura I. Lascarez-Lagunas, Ayana L. Henderson, Marina Martínez-García, Rajendiran Karthikraj, Victor Barrera, Shannan Ho Sui, Kurunthachalam Kannan, and Mónica P. Colaiácovo

SUPPLEMENTAL FIGURES



Supplemental Figure 1. DEET exposure leads to CHK-1 activation. Related to Figure 2E. High-resolution images of premeiotic tip and mid pachytene nuclei from wholemounted gonads immunostained for phosphorylated CHK-1 (pCHK-1; green) and costained with DAPI (blue). Scale bar, 5 µm.



Supplemental Figure 2. RNA-sequencing validation by quantitative RT-PCR. Related to Figure 4. 8 genes involved in DSB formation (*spo-11*), DSB repair (*mre-11*, *msh-5*, *rad-51*, and *rad-54*), and DNA damage response (*atl-1*, *cep-1*, and *chk-1*) for which no change in expression was observed, and 4 DE genes, were examined by qRT-PCR see data S1J. Each analysis was performed with technical triplicates and normalized to *gpd-1* (GAPDH). Gene expression level change relative to vehicle alone is depicted in log2 scale with SEM. Only DE genes show gene expression changes.

#	Primer pair	F sequence	R sequence
1	atl-1	cgtcgaaccttcgtcttctc	gcattctcctgcgttttctc
2	cep-1	ttccgacgcaagtagtctcc	ccgtttgcattgaacaacac
3	chk-1	gtctggtcgtctgggattgt	ttgctgatccatcccatgta
4	mre-11	ctgtttggaaagcacagcaa	ttgaatgctcgaacaagacg
5	msh-5	ccccaaaacagctttccata	ggcgtcttgaatggatcact
6	rad-51	ccaggctgatgctaaaaagc	ttcggcttctggtaaattgg
7	rad-54	cgtcttcgaatgtggatcg	gtcgttttcttcggcttcag
8	spo-11	tggacctacgaaagaatttgc	tgatcgatggtgaaacgatg
9	cyp-14A2	ccactcccattcattggaaa	tgtccataatgtgaagcatgg
10	cyp-14A3	ccactacccgcagttgttct	ggacgattagcacggttgat
11	сур-34А10	ggctcccaaaaggacctact	acggcgcttcctaacttgat
12	dhs-23	gcatcgggacaagatgaagt	cagtcaaaagagcaccagca

 Table S1. Primers utilized for quantitative RT-PCR analysis. Related to STAR Methods.