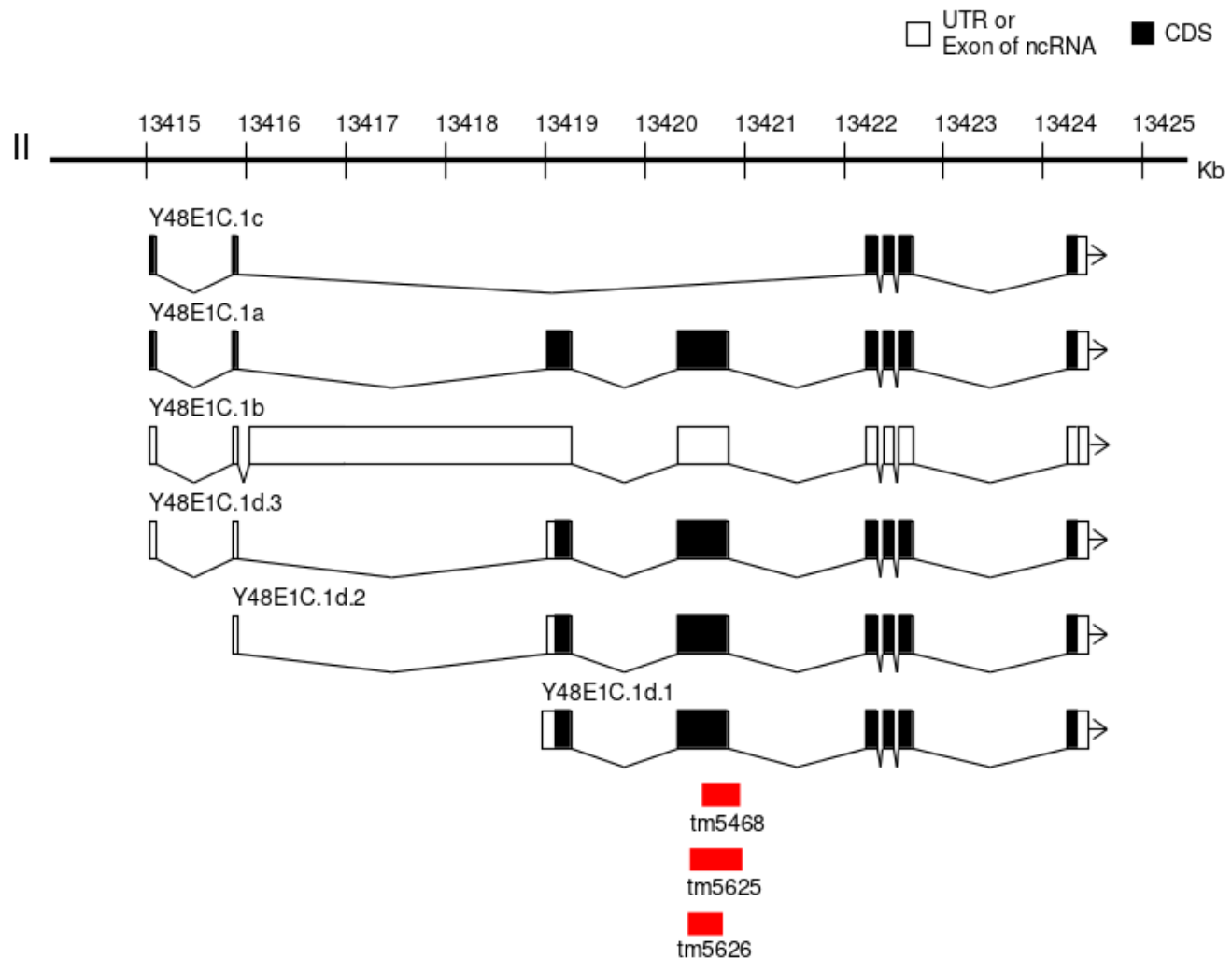


## Novel deletion alleles of a *C. elegans* gene Y48E1C.1, named as *tm5468*, *tm5625* and *tm5626*

Sayaka Hori<sup>1</sup>, Yuji Suehiro<sup>1</sup>, Sawako Yoshina<sup>1</sup> and Shohei Mitani<sup>1§</sup>

<sup>1</sup>Department of Physiology, Tokyo Women's Medical University School of Medicine, Shinjuku-ku, Tokyo, 162-8666, Japan

<sup>§</sup>To whom correspondence should be addressed: mitani.shohei@twmu.ac.jp



**Figure 1.** Location of the novel alleles

### Description

We report [tm5468](#), [tm5625](#) and [tm5626](#) as novel deletion alleles of the gene [Y48E1C.1](#) that is the only ortholog of human calmodulin-lysine N-methyltransferase (CAMKMT)1. CAMKMT encodes an evolutionarily conserved enzyme class I protein methyltransferase that acts in the formation of trimethyllysine in calmodulin for calcium-dependent signaling<sup>2</sup>. CAMKMT mutation is associated with Hypotonia-cystinuria syndrome in human<sup>2,3</sup>. The alleles were isolated from the comprehensive screening of gene deletions generated by TMP/UV4. In the screening, all the alleles were detected by nested PCR using the following primer sets, 5'-TCAAGCCACGCCACACTTA-3' and 5'-GAAGGCATACAGTGGGGGTA-3' for the first round PCR and 5'-CGCCCACTTAATGGTTAT-3' and 5'-GGCAGTGTAGGGATACTGT-3' for the second round PCR. By Sanger sequencing, the 30 bp flanking sequences of the alleles [tm5468](#), [tm5625](#) and [tm5626](#) were identified as

AATCCTTCACACACCACAACAGAAATCCTA – [384 bp deletion] -CGAGGTCACGCCCACACATTGGGCGGAGTT, CCGATGCTCCGTGCTGCTCCAAGTGCTCCG – [627 bp deletion + 9 bp insertion (TAATCTTGT)] – AGTACTCCTACAGTATCCCTACACTGCCCC, and AAAAAAGGATGACGTCACAGTTGCTCCGAT – [256 bp deletion] – ACGCCGATTCGGCAGCCGAATGATCTACAG, respectively. Based on the information about the splicing isoforms of [Y48E1C.1](#) (WormBase, <http://www.wormbase.org>, WS259), the forth exon of [Y48E1C.1a](#), [Y48E1C.1b](#) (annotated as non coding RNA) and the second exon of [Y48E1C.1d](#) transcripts are deleted in [tm5468](#), [tm5625](#) and [tm5626](#)(Fig. 1). Presumably, all of the alleles do not affect [Y48E1C.1c](#). According to information of protein in Wormbase, this exon contains a predicted some motif, suggesting hypothetical functional deficiency of [Y48E1C.1aY48E1C.1b](#), and [Y48E1C.1d](#) in the deletion mutants. In addition, these alleles are expected to be usable for comparing functions among the isoform c and the other isoforms. However, no visually obvious phenotypes (Let, Unc, and Dpy) were observed in [tm5468](#), [tm5625](#) and [tm5626](#).

## Reagents

FX05468 [Y48E1C.1 \(tm5468\)](#) II (Not outcrossed)

FX05625 [Y48E1C.1 \(tm5625\)](#) II (Not outcrossed)

FX05626 [Y48E1C.1 \(tm5626\)](#) II (Not outcrossed)

## References

Magnani R, Dirk LM, Trievel RC, Houtz RL. Calmodulin methyltransferase is an evolutionarily conserved enzyme that trimethylates Lys-115 in calmodulin. *Nat Commun.* 2010;1:43. DOI: 10.1038/ncomms1044 | PMID: 20975703.

Magen S, Magnani R, Haziza S, Hershkovitz E, Houtz R, Cambi F, Parvari R. Human calmodulin methyltransferase: expression, activity on calmodulin, and Hsp90 dependence. *PLoS One.* 2012;7(12):e52425. DOI: 10.1371/journal.pone.0052425 | PMID: 23285036.

Bartholdi D, Asadollahi R, Oneda B, Schmitt-Mechelke T, Tonella P, Baumer A, Rauch A. Further delineation of genotype-phenotype correlation in homozygous 2p21 deletion syndromes: first description of patients without cystinuria. *Am J Med Genet A.* 2013; Aug 161A(8):1853-1859. DOI: 10.1002/ajmg.a.35994 | PMID: 23794250.

Gengyo-Ando K, Mitani S. Characterization of mutations induced by ethyl methanesulfonate, UV, and trimethylpsoralen in the nematode *Caenorhabditis elegans*. *Biochem Biophys Res Commun.* 2000; Mar 5:269(1):64-69. DOI: 10.1006/bbrc.2000.2260 | PMID: 10694478.

**Funding:** The National BioResource Project.

**Reviewed By:** [James Lee](#)

**History:** Received August 30, 2017 Accepted October 3, 2017 Published October 3, 2017

**Copyright:** © 2017 by the authors. This is an open-access article distributed under the terms of the Creative Commons Attribution 4.0 International (CC BY 4.0) License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

**Citation:** Hori, S; Suehiro, Y; Yoshina, S; Mitani, S (2017). Novel deletion alleles of a *C. elegans* gene Y48E1C.1, named as *tm5468*, *tm5625* and *tm5626*. *microPublication Biology.* <https://doi.org/10.17912/W2CCQ14>