

Table 1. Content of 5me-Deoxycytidine (5-meC) in DNA Preparations of Ten Species and the Estimated Average Numbers of SmdC per (Haploid) Genome of *A. thaliana*, *M. musculus*, *D. melanogaster*, *E. coli*, *S. cerevisiae*, *S. boulardii*, *S. paradoxus*, *P. pastoris*, and *S. pombe*^a

species	genome size MB	% methylation	Est. no. SmdC per MB	Est. no. SmdC/genome
<i>A. thaliana</i>	119.7	14 ± 1.5	5.0 × 10 ⁴	6.0 × 10 ⁶
<i>M. musculus</i>	2800	7.6 ± 0.8	3.2 × 10 ⁴	8.9 × 10 ⁷
<i>D. melanogaster</i>	139.5	0.034 ± 0.013	1.4 × 10 ³	2 × 10 ⁴
<i>E. coli</i> DHSα	4.6	2.3 ± 0.1	1.2 × 10 ⁴	5.4 × 10 ⁴
<i>E. coli</i> GM2929 (<i>dcm-6</i>)	4.6	0.016 ± 0.002	81	3.8 × 10 ²
<i>S. cerevisiae</i> BY4741**	12.5	not detected (<0.002)	not detected (LOD = 8)	not detected (LOD = 96)
<i>S. cerevisiae</i> D27–310B**	12.5 (b.f.)	not detected	not detected	not detected
<i>S. cerevisiae</i> AWRI 796**	12.5 (b.f.)	not detected	not detected	not detected
<i>S. cerevisiae</i> SCHD0308	12.5 (b.f.)	not detected	not detected	not detected
<i>S. cerevisiae</i> SaflagerW-34/70	12.5 (b.f.)	not detected	not detected	not detected
<i>S. boulardii</i> CBS 5926	11.4	not detected	not detected	not detected
<i>S. paradoxus</i> KPN3829	11.8	not detected	not detected	not detected
<i>S. pombe</i> 972h-	13.8	not detected	not detected	not detected
<i>P. pastoris</i> SMD1168	9.4	not detected	not detected	not detected

^aTotal SmdC per genome and MB average estimations were calculated based on genome size and GC content.^{29–35} *n* = 3 from biological triplicates, (**) analysis of stationary, and exponential cultures yielded the same result. b.f. = bona fide. Est. = estimated.

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