

CORRECTION

Correction: Diatom genomes come of age

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Correction

After the publication of this minireview [1], we noted errors in the figures given in Table 1 and the text for the percentage of genome that is non-coding. The percentages should be 42% for *Thalassiosira pseudonana* and 45% for *Phaeodactylum tricorutum* (see Table 1 a corrected version of Table 1).

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Table 1: Comparison of the genome properties of *Thalassiosira pseudonana* and *Phaeodactylum tricorutum* genomes*

	<i>Thalassiosira pseudonana</i>	<i>Phaeodactylum tricorutum</i>
Genome size (Mb)	32.4	27.4
Predicted genes	11,776	10,402
Introns	17,880	8,169
Number of chromosomes	24	33
G+C content	About 48%	About 47%
Percentage of genome that is non-coding	About 42%	About 45%
ESTs in GenBank	61,913	133,871

*Data from [2,3].

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