



Correction to: 2D morphometric analysis of *Arabidopsis thaliana* nuclei reveals characteristic profiles of different cell types and accessions

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The original version of this article unfortunately contained a mistake.

The Tables 1 and 2 were published erroneously.

The corrected Tables 1 and 2 were shown in the next page.

The publisher sincerely apologizes for this error and any inconvenience caused to the authors and readers of the journal.

The original article has been corrected.

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The original article can be found online at <https://doi.org/10.1007/s10577-021-09673-2>.

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Table.1 Morphometric parameters used for chromatin profiling together with their Biological equivalent, Description, and Mathematical formula.

	Biological feature	Parameter (trait)	Description	Formula	Unit
Nucleus	Size	Area	Sum of pixels within area of <i>Nu</i>	Σ pixels	pixel
	DNA density	Intensity mean	Average fluorescent intensity of DAPI stain of <i>Nu</i>	Σ intensity / number pixels	
	Variation in DNA density	Heterogeneity	Fraction of pixels that deviate more than 10% from the averaged intensity mean of <i>Nu</i>		
	DNA content	Intensity sum	Sum of fluorescent intensity of DAPI stain of <i>Nu</i>		
chromocenter	Total size	Total area	Sum of areas of all <i>CCs</i> in <i>Nu</i>	Σ pixels x number <i>CC</i>	pixel
	Average size	Area average	Average area of individual <i>CCs</i>	Σ pixels / number <i>CC</i>	pixel
	DNA density	Intensity (mean)	Average fluorescent intensity of DAPI stain of individual <i>CCs</i> in <i>Nu</i>	Σ intensity / number pixels	
	DNA content	Intensity (sum)	Sum of fluorescent intensity of DAPI stain of all <i>CCs</i> in <i>Nu</i>	Σ intensity	
	Heterochromatin content in <i>Nu</i>	Relative Heterochromatin Fraction (RHF)	Fraction of DNA content of <i>CCs</i> per <i>Nu</i>	Σ (area <i>CC</i> x intensity mean <i>CC</i>) / (area <i>Nu</i> x intensity mean <i>Nu</i>)	
	Number <i>CCs</i>	# <i>CC</i>	Sum of all <i>CCs</i> in <i>Nu</i>	Σ <i>CCs</i>	
euchromatin	Size	Area	Area of <i>Eu</i>	Σ pixels <i>Nu</i> - Σ pixels <i>CC</i>	pixel
	DNA density	Intensity (mean)	Average fluorescent intensity of DAPI stain of <i>Eu</i>	$[\Sigma$ intensity (sum) <i>Nu</i> - Σ intensity (sum) <i>CC</i>] / area <i>Eu</i>	
	DNA content	Intensity (sum)	Sum of fluorescent intensity of DAPI stain of <i>Eu</i>	Σ intensity (sum) <i>Nu</i> - Σ intensity (sum) <i>CC</i>	

Note: Chromocenter (*CC*) is used in the meaning of a microscopically visual heterochromatin body in the interphase nucleus. DNA density as a biological feature is used as equivalent to DNA concentration. *Nu* = nucleus, *Eu* = euchromatin

Table.2 Fraction of transposable elements in heterochromatin and euchromatin

		cell types				accessions				
		GC	PC	VC	EC	Col	Cvi	C24	Ler	Ws
125 Mb	RHF	0.19	0.13	0.12	0.09	0.13	0.08	0.12	0.13	0.10
	DNA in CC (Mb)	23.75	16.25	15.00	11.25	16.25	10.00	15.00	16.25	12.50
	amount of TEs in CC	7.49	-0.01	-1.26	-5.01	-0.01	-6.26	-1.26	-0.01	-3.76
	fraction of TE in CC	0.60	0.00	-0.10	-0.40	0.00	-0.50	-0.10	0.00	-0.30
	fraction of TE in eu	0.40	1.00	1.10	1.40	1.00	1.50	1.10	1.00	1.30
157 Mb	RHF	0.19	0.13	0.12	0.09	0.13	0.08	0.12	0.13	0.10
	DNA in CC (Mb)	23.75	16.25	15.00	11.25	16.25	10.00	15.00	16.25	12.50
	amount of TEs in CC	7.49	-0.01	-1.26	-5.01	-0.01	-6.26	-1.26	-0.01	-3.76
	fraction of TE in CC	0.60	0.00	-0.10	-0.40	0.00	-0.50	-0.10	0.00	-0.30
	fraction of TE in eu	0.40	1.00	1.10	1.40	1.00	1.50	1.10	1.00	1.30
191 Mb	RHF	0.19	0.13	0.12	0.09	0.13	0.08	0.12	0.13	0.10
	DNA in CC (Mb)	36.29	24.83	22.92	17.19	24.83	15.28	22.92	24.83	19.10
	amount of TEs in CC	20.03	8.57	6.66	0.93	8.57	-0.98	6.66	8.57	2.84
	fraction of TE in CC	1.05	0.45	0.35	0.05	0.45	-0.05	0.35	0.45	0.15
	fraction of TE in eu	-0.05	0.55	0.65	0.95	0.55	1.05	0.65	0.55	0.85
211 Mb	RHF	0.19	0.13	0.12	0.09	0.13	0.08	0.12	0.13	0.10
	DNA in CC (Mb)	40.09	27.43	25.32	18.99	27.43	16.88	25.32	27.43	21.10
	amount of TEs in CC	23.83	11.17	9.06	2.73	11.17	0.62	9.06	11.17	4.84
	fraction of TE in CC	1.13	0.53	0.43	0.13	0.53	0.03	0.43	0.53	0.23
	fraction of TE in eu	-0.13	0.47	0.57	0.87	0.47	0.97	0.57	0.47	0.77

CC=chromocenter, TE=transposon element, eu=euchromatin, GC=gard cell, PC=pavement/parenchyma cell, VC=vascular cell. EC=endopolyploid cells, Mb=megabasepairs

The negative values for the *fraction of TEs in CC* (red) suggests that in addition to all TEs also a fraction of the major tandem repeats is in euchromatin. The negative values for the *fraction of TEs in euchromatin* (i.e., GC values in 191 Mb and 211 Mb) suggest more DNA in the CCs than the sum of tandem repeats and TEs. This implies that genes are also present in CCs.