Table 2. Descriptive statistics of selected features

FEATURE NAME	UNITS	STATISTICS OF COMPUTED MORPHOMETRICS ($N = 100$)					
		EPC2		CP-A		CP-D	
		μ^{a}	σ^{b}	μ	σ	μ	σ
Cell volume	μm^{3}	1,010	321	1,173	249	1,803	787
Nuclear volume	μm^{3}	163	51	209.41	67.84	432	188
N/C volume ratio	–	0.20	0.03	0.22	0.07	0.33	0.10
Nuclear content	a.u.	113	35	146	47	299	130
Nuclear sphericity	–	1.27	0.06	1.33	0.09	1.31	0.14
Total nucleolar volume	μm^{3}	4.22	3.21	12.36	7.12	4.79	2.87
Average nucleolar margination	μm	1.12	0.45	1.55	0.80	2.24	0.70
Number of dense DNA clumps	–	1.97	1.28	1.93	1.46	5.58	3.92
Moderate-density DNA volume fraction	–	0.62	0.02	0.61	0.02	0.68	0.04
High-density DNA volume fraction	–	0.19	0.02	0.20	0.02	0.15	0.02
Chromatin density variability (Markovian contrast)	a.u.	3400	493	3512	660	3798	556
Chromatin texture smoothness (Markovian energy)	a.u.	0.73	0.04	0.73	0.05	0.64	0.05

[a] Mean.
[b] Standard deviation.