

Table S3. Pearson correlation between gene parameters estimated for 56 *Rosaceae* *PEBP* genes. Correlation coefficients with significant adjusted p-value <0.05 were shown in bold font. Positive correlation coefficients were colored in red and negative correlations colored in blue.

	Length	CAI	totalGC%	GC1%	GC2%	GC3%	ENC
Length		-0.10	0.19	0.32	-0.03	0.16	0.19
CAI	-0.10		0.56	0.09	0.16	0.72	-0.62
totalGC%	0.19	0.56		0.80	0.45	0.93	-0.34
GC1%	0.32	0.09	0.80		0.17	0.67	0.09
GC2%	-0.03	0.16	0.45	0.17		0.18	-0.26
GC3%	0.16	0.72	0.93	0.67	0.18		-0.43
ENC	0.19	-0.62	-0.34	0.09	-0.26	-0.43	