CORRIGENDUM

In the paper by Andrew G. Clark and Jørgen Bundgaard (GENETICS **108:** 181–200) entitled "Selection components in background replacement lines of Drosophila," Table 1 was misprinted. The correct Table 1 appears below.

TABLE 1

Estimates of viability of eg/eg relative to eg/+ and standard errors

Genera-tion	Line	eg/eg × eg/+	eg/+ × eg/eg	eg/+ × eg/+
0		0.551 ± 0.021 (5778)[a]	0.770 ± 0.023 (7636)	0.776 ± 0.026 (14691)
10	GB-MIX	0.998 ± 0.076 (1120)	0.617 ± 0.031 (2077)	0.636 ± 0.041 (3014)
	GB-REP	0.903 ± 0.090 (576)	0.876 ± 0.093 (1162)	0.774 ± 0.068 (1668)
	Wd-MIX	0.932 ± 0.062 (1380)	0.756 ± 0.042 (2648)	0.693 ± 0.040 (2994)
	Wd-REP	0.489 ± 0.076 (817)	0.218 ± 0.032 (2463)	0.433 ± 0.031 (1758)
20	GB-MIX	0.938 ± 0.030 (2242)	0.926 ± 0.037 (1802)	1.121 ± 0.039 (2902)
	GB-REP	0.938 ± 0.030 (3801)	0.908 ± 0.031 (3612)	1.094 ± 0.043 (5960)
	Wd-MIX	0.882 ± 0.033 (3099)	0.997 ± 0.043 (3382)	1.212 ± 0.073 (4419)
	Wd-REP	0.835 ± 0.030 (2846)	0.920 ± 0.029 (2580)	0.927 ± 0.037 (4526)

[a] *A posteriori* comparisons are indicated by lines: estimates not connected are different at $P < 0.05$. Sample sizes are given in parentheses.