

SUPPLEMENTARY TABLE 2. — Loci and sample sizes. Numbers without superscript are data new to this paper.

Locus	Alignment length (bp)	<i>Zea mays</i>			<i>Z. luxurians</i>	<i>Tripsacum</i>
		<i>mays</i>	<i>parviglumis</i>	<i>mexicana</i>		
adh1	1620	31 ^{d,h}	8 ^a	8 ⁱ	8 ^h	1 ^d
asg11	512	24 ^d	6 ^a	7	5	
asg35 ^k	1010	22	12	12	10	1
asg64	658	20	16	11	11	
asg65	809	17	14	12	11	1
bnl7-13 ^k	934		12	11	11	1
bz2	640	25 ^a	12 ^a	7	11	1 ^d
c1	814	15 ^g	11 ^g		7 ^g	1 ^g
csu1132	589	25 ^a	8 ^a	10	10	1 ^d
csu1138	373	25 ^d	14	10	10	1 ^d
csu1171 ^k	471	25 ^a	9 ^a	10	8	1 ^d
csu381 ^k	1186	25 ^a	10 ^a	9	9	1 ^d
csu636 ^k	647	22	15	13	9	1
csu838	786	21	13	12	11	1
d8	827	25 ^d	13 ^a	10	8	1 ^d
fus6	273	23	10	12	12	1
gl1510	777	22	15	13	12	
glb1	1323	30 ^{c,d}	8 ^c		6 ^c	1 ^c
mgs3020	978	22	14	13	13	1
pepc1070	1000	18	13	8	10	
pepc1150	705	18	16	12	11	
tb1	3029	23	7	7	11	1 ^j
ts2	1104	24 ^a	10 ^a	9	8	1
vp1010	1299	22	12	9	6	1
wip1	761	12 ^b	15 ^b	10 ^b	9 ^b	1 ^b
wx1	1054	37 ^e	19 ^e	20 ^e	21 ^e	1 ^f

^a Tenaillon *et al.* (2004)

^b Tiffin and Gaut (2001)

^c Hilton and Gaut (1998)

^d Tenaillon *et al.* (2001)

^e Includes previously unpublished sequences from our lab (Genbank accessions AF29500-39).

^f Genbank accession DQ369871

^g Hanson *et al.* (1996) and unpublished sequences from our lab (Genbank accessions AF229540-53)

^h Eyre-Walker *et al.* (1998) and Gaut and Clegg (1993)

ⁱ Unpublished sequences from our lab (Genbank accessions AF293887-94)

^j Genbank accession AF322118

^k Noncoding loci