

SUPPLEMENTARY TABLE 7. — Identical sequences shared among taxa^a

Locus	Taxa Compared					
	MA-PA	MA-ME	MA-LU	PA-ME	PA-LU	ME-LU
<i>adh1</i>	2	1		1		
<i>asg11</i>	2	2		3	1	
<i>asg35</i>	1	2		1		
<i>asg64</i>	4	5		4		
<i>asg65</i>				2	1	
<i>bnl7-13</i>				3		
<i>bz2</i>	2	1		2		
<i>c1</i>	1					
<i>csu1132</i>	1	2		1		
<i>csu1138</i>	2	2	1	2	1	1
<i>csu1171</i>	2	1	1	1		2
<i>csu381</i>	1	1		1		
<i>csu636</i>	4	3		5		
<i>csu838</i>	3	3		2		1
<i>d8</i>	1			1		
<i>fus6</i>		1		1	2	1
<i>gl1510</i>	1	2		2	1	
<i>glb1</i>	1					
<i>mgs3020</i>	2	2		3		
<i>pepc1070</i>	2			1		
<i>pepc1150</i>	1	3		2		
<i>ts2</i>		3		1	1	2
<i>vp1010</i>					2	
<i>wip1</i>	1			1		
<i>wx1</i>	4	3		1		

^aShown is the number of distinct identical sequences shared between at least one individual of both taxa.

MA *Zea mays* ssp. *mays*; PA *Zea mays* ssp. *parviglumis*; ME *Zea mays* ssp. *mexicana*; LU *Zea luxurians*