

Table S6: Number of genes with each pattern of inheritance in hybrids							
	Additive	<i>sim</i> dominant	<i>sec</i> dominant	<i>mau</i> dominant	Underdominant	Overdominant	Conserved
<i>(sim x mau) x sim</i>	88	735		182	197	270	376
<i>(mau x sim) x mau</i>	97	733		210	168	260	412
<i>(sec x mau) x sec</i>	237		540	292	280	371	322
<i>(mau x sec) x mau</i>	220		489	302	329	382	313
<i>(sim x sec) x sim</i>	180	450	215		141	187	536
<i>sim x mau</i>	51	68		20	71	27	17
<i>mau x sim</i>	77	57		31	40	40	11
<i>sec x mau</i>	70		62	30	39	44	17
<i>mau x sec</i>	58		69	35	34	49	19
<i>sim x sec</i>	49	59	42		51	24	19

**Table S6:** Number of genes within each category of inheritance pattern in the hybrids. Stage 2 hybrids are shown in the first five rows and stage 5 hybrids are shown in the last five rows of the table.