

Table S2 Sequence statistics for all genotypes sequenced in this study. Percentage of mapped reads are to the New Mexico Valencia *A de novo* assembly

Genotype	Total Reads	Mapped Reads
		%
Florida 07	13,881,744	60.30
NC3033	15,736,991	55.98
SPT06-06	16,252,634	56.48
Tifrunner	35,457,450	61.97
C76-16	9,949,147	63.36