Supplementary material

Amel Oueslati, Amel Salhi-Hannachi, François Luro, Hélène Vignes, Pierre Mournet and Patrick Ollitrault. Genotyping By Sequencing reveal the interspecific C. maxima / C. reticulata admixture along the genomes of modern citrus varieties of mandarins, tangors, tangelos, orangelos and grapefruits. Plos One (submitted)

S1 fig: Distribution of missing data among markers (a) and individuals (b)

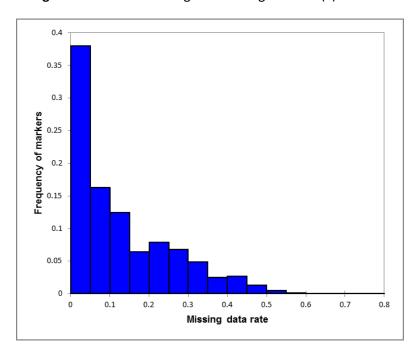


Fig S1A: distribution of missing data rate for markers

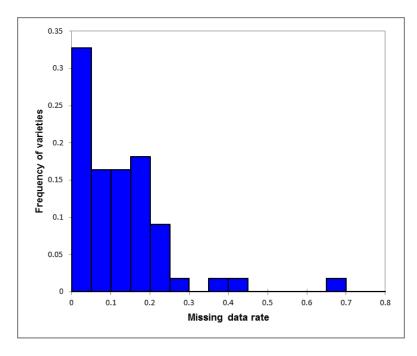


Fig S1B: distribution of missing data rate for varieties