

Table S14 tDNA numbers per chromosome in *P. sorbitophila* genome

chr.	Nbr per chr.			in eq. haploid
	Heterozygous	Homozygous	Total	
A	11	10	21	21
B	11	10	21	
C	6	4	10	10
D	6	4	10	
I (I/E)	0	0	0	0
E (I/E)	0	0	0	
J (J/F)	4	0	4	4
F (J/F)	4	0	4	
E (E/F)	17	0	17	17
F (E/F)	17	0	17	
I (I/J)	25	0	25	25
J (I/J)	25	0	25	
G	0	17	17	17
H	0	17	17	
K	0	25	25	25
L	0	25	25	
M	25	0	25	25
N	25	0	25	
Total	176	112	288	144

Genes encoding tRNAs were searched with tRNAscan-SE (Lowe and Eddy, 1997) and obtained sequences were analyzed for possible exceptions to the eukaryotic cloverleaf model, as described in Marck and Grosjean (2002). Initiator Met tRNA genes were distinguished from the elongator sequences thanks to the "GGG" sequence in positions 29-31. For chr. E/F/I/J, the syntenic heterozygous pairs considered are mentioned in brackets.