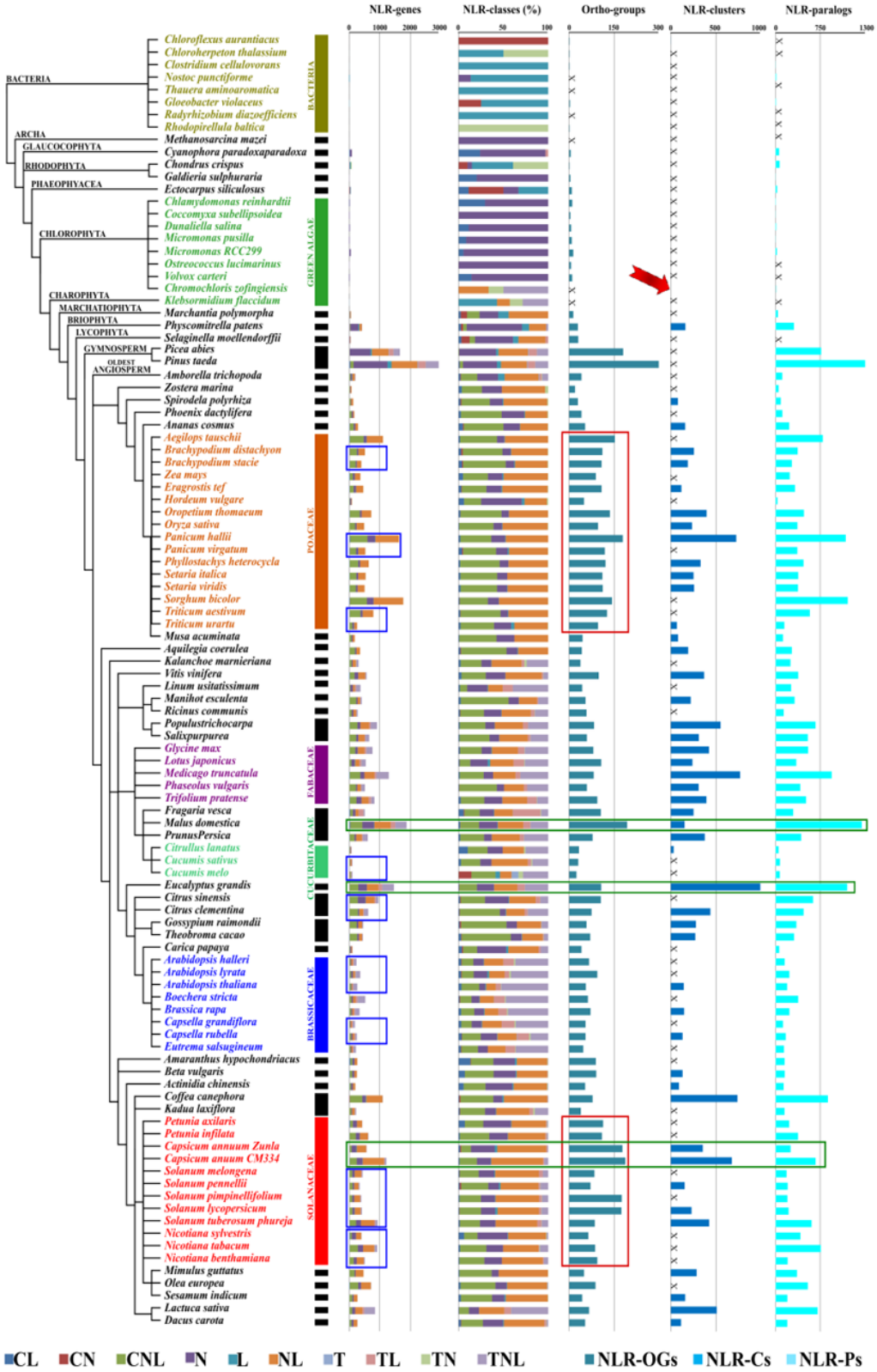

Figure S1. Arrangement and composition of NB-LRR genes in 104 genomes. The five bar charts, from left to right, represent for each genome the total number of NB-LRR genes (NLR-genes), the proportion of different NB-LRR protein classes (NLR-classes), the number of NB-LRR gene ortho-groups (NLR-OGs), NB-LRR clusters (NLR-Cs) and NB-LRR paralog genes (NLR-Ps). The colour boxes underlined the substantial differences of NB-LRR families. In the charts missing data are denoted with an "X" mark.