

Additional file 13: 3D distribution of gene sequence SNPs according to their haplotypic  $G_{ST}$  value ; a:  $G_{ST}$  value for three horticultural groups (mandarins, pummelos and citrons); b:  $G_{ST}$  values for three basic taxa (*C. reticulata, C. maxima, C. medica*) after introgression information corrections.