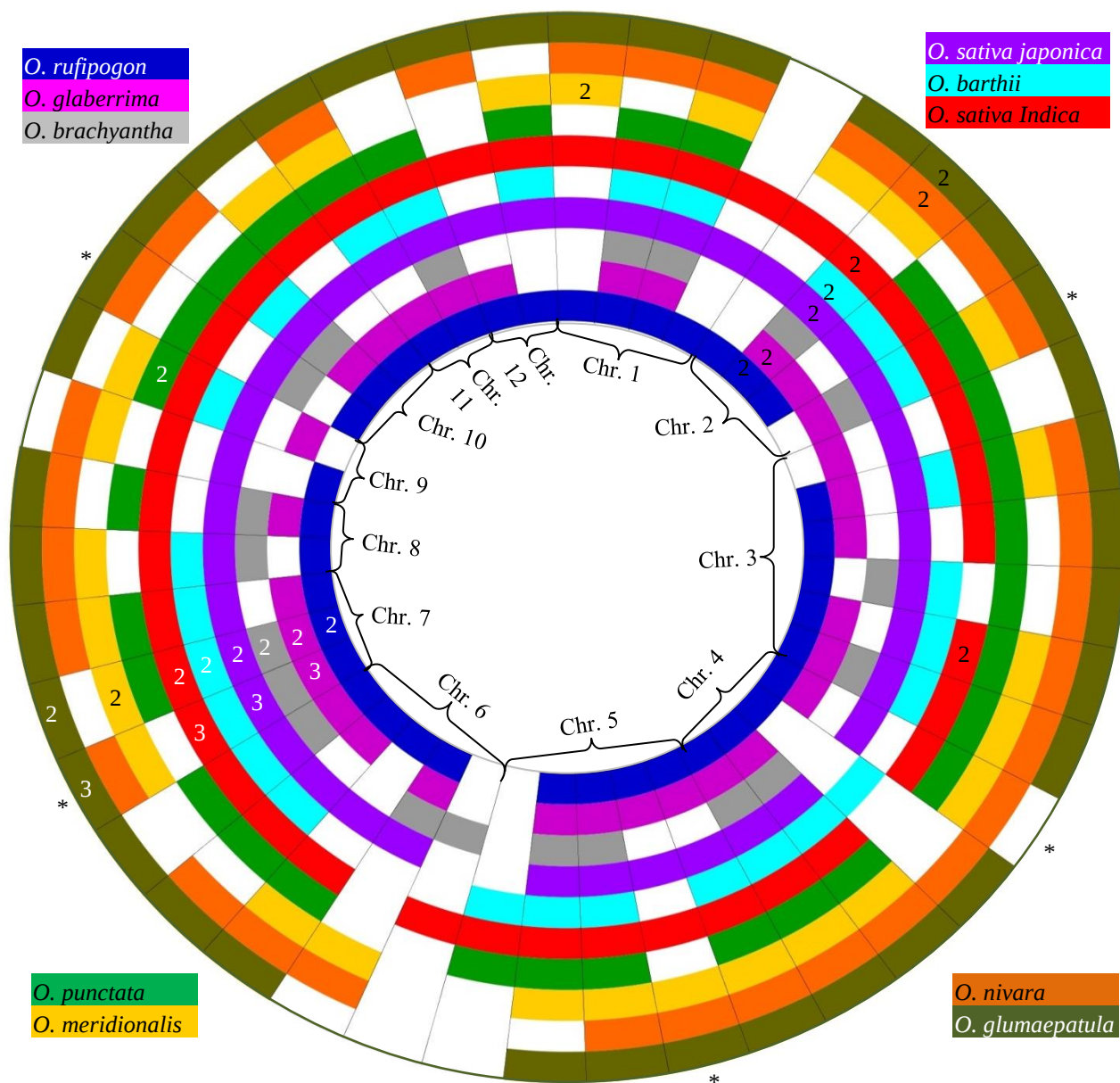




Additional file 6: Figure S2. Syntenic analysis of orthologous VQs among ten species of the *Oryza* genus. The coordinates were mapped using the GenomeRing program (Herbig et al. 2012. *Bioinformatics* **28**, i7-i15). The stars indicated that these loci were undergone positive selection with K_a/K_s value significantly larger than 1. The numbers indicated copies of tandemly duplicated genes in these loci among different species. Locus names and coordinates were listed in Table S3.