

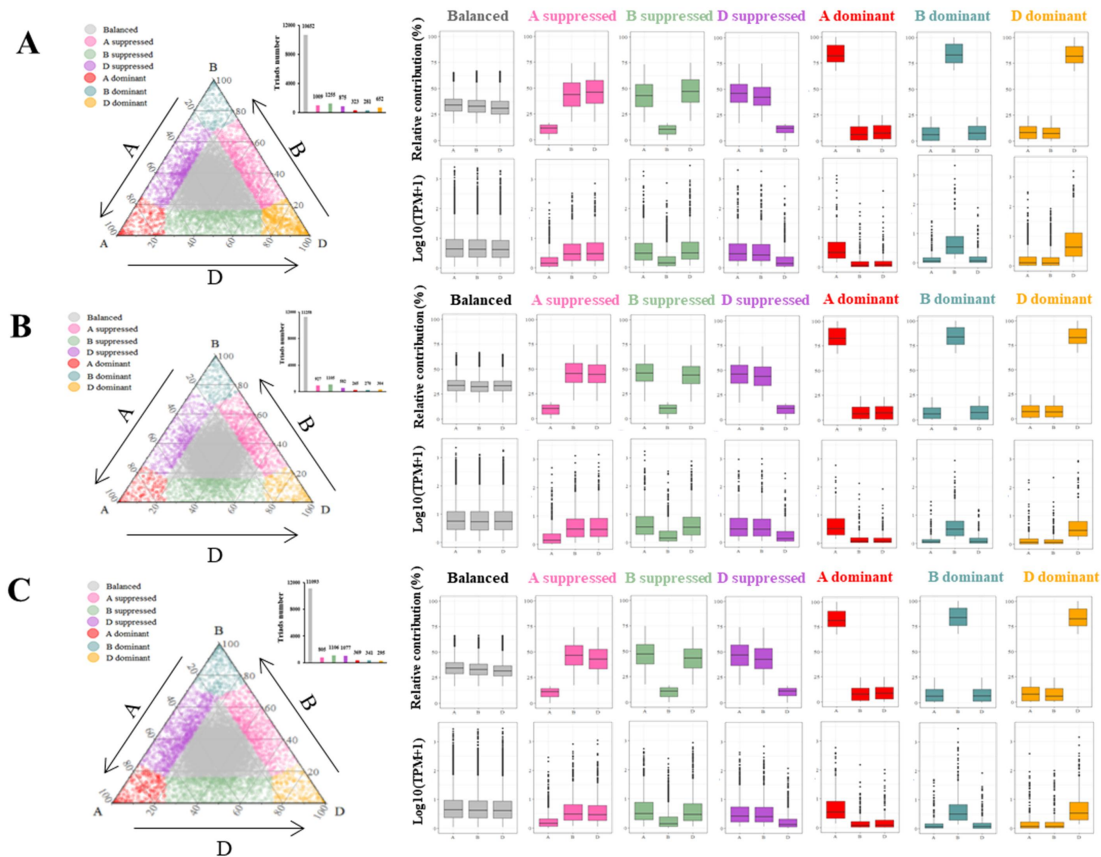


Figure. S2 – Homoeolog expression bias in homoeolog triads. (A) Ternary plot showing relative expression abundance of triads in the parents of 6x-1. (B) Ternary plot showing relative expression abundance of triads in 6x-2. (C) Ternary plot showing relative expression abundance of triads in the parents of 6x-2. Each circle represents a gene triad with an A, B, and D coordinate consisting of the relative contribution of each homoeolog to the overall triad expression. The distribution of the numbers of homeoallele transcription bias categories is shown at the top right. Box plots indicate the relative contribution of each subgenome based on triad assignment to the seven categories.