



Figure S5. The full syntentic maps of chromosome regions containing Hsf genes across *L. japonicus*, *M. truncatula*, *G. max* and *C. cajan* genes. Figures in (A)–(Q) are 17 groups of syntentic segments. Arrows represent individual genes and their transcriptional orientations. Distances are roughly to scale. The Hsf genes are highlighted in red, and their flanking genes are shown in black. All genes are numbered from the left to the right for each interval. Duplicated genes are given the same number but ordered by a, b, c, Genes with syntentic matches are linked with lines.