Figure S3. Scatter plots of τ values for highly expressed genes in respective japonica or indica tissues. Nipponbare τ, Minghui63 τ, and Zhenshan97 τ values were calculated using 44, 36, and 39 tissues, respectively. Red points represent changed-tissues jDE genes. Diagonal lines mean same japonica and indica τ values. The top and right panels of each scatter plot show the percentages of changed-tissues (solid line) and global (broken line) jDE genes on a window of 0.1 τ window for highly expressed genes in Nipponbare and indica strain tissues, respectively.