



Figure S4 Pairwise comparison of developmental expression changes among diploid and polyploid cotton. Log₂ expression ratios were calculated for 20 dpa versus 10 dpa within each genome using 2-DE (A) and iTRAQ datasets (B). Lower-left panels of the comparison matrix show scatterplots while upper-right panels give Pearson's correlation coefficients, with the font size proportional to the absolute value of the correlation. Asterisks indicate significance of the adjacent correlation coefficient: * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$.