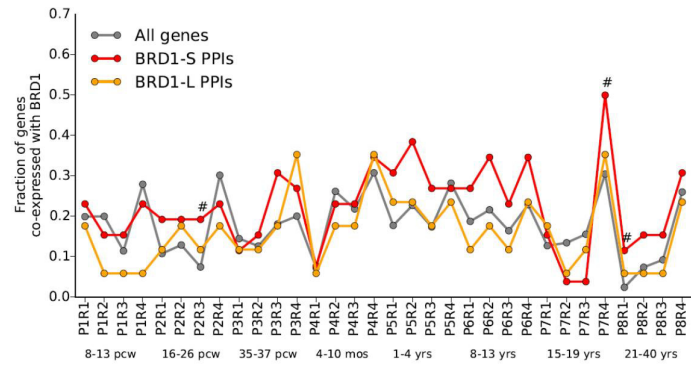
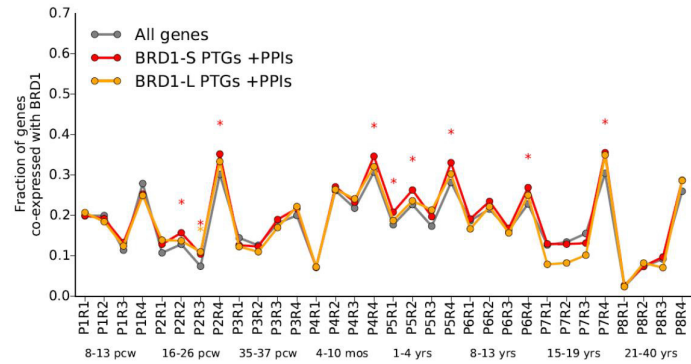


A



B



Spatiotemporal co-expression landscapes of BRD1-S and BRD1-L PPIs and PTGs in human brain. The fraction of co-expressed genes with BRD1 across 32 spatiotemporal intervals. (A) The fraction of genes co-expressed with BRD1 across all genes (grey) and genes in the BRD1-S and BRD1-L PPI networks (red and orange). None of these were significant at $P < 0.05$ after adjusting for multiple tests. (#) denote $P < 0.05$ before correcting for multiple testing. (B) The fraction of genes co-expressed with BRD1 across all genes (grey) and genes in the BRD1-S and BRD1-L PTGs + PPI networks (red and orange). A one-sided binomial test was performed for each BRD1 sub-network and spatiotemporal interval compared to the expected fraction in the background where the asterisk (*) denote $P < 0.05$ after adjusting the P-value for multiple tests.