



Figure S1 The distance distribution of intergenic unique poly(A) sites after 3'-UTRs. 1–1000 nt downstream of 3'-UTR is selected to investigate the distribution of poly(A) sites in intergenic region. X-axis shows the 200 sub-regions which are 5 nt in length (e.g., 1–5 and 6–10). Y-axis labels the percentage of poly(A) sites for each sub-region.