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* You should state whether an appropriate sample size was computed when the study was being designed
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* Indicate how samples were allocated into experimental groups (in the case of clinical studies, please specify allocation to treatment method); if randomization was used, please also state if restricted randomization was applied
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Additional data files include source data files for Figure 4, supplementary files for quantitative data corresponding to heat-map representations in Figure 1, Figure 1 – figure supplements 1-4 and Figure 2. TaDa sequencing data and experimental information have been deposited in the ArrayExpress database at EMBL-EBI ([www.ebi.ac.uk/arrayexpress](https://www.ebi.ac.uk/arrayexpress/)) and will be available under accession number E-MTAB-8935. Processed data and intermediate files, including the read coverage at GATC motifs, are available at GitLab (<https://gitlab.com/roman.arguello/ir_tada>), as described in the “Data availability” section.