

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: gRNA editing efficiency of ABE and CBE measured by SURRO-seq technology.

File Name: Supplementary Data 2

Description: Model architecture of CRISPRon-BE and detailed information on hyper-parameter screening.

File Name: Supplementary Data 3

Description: Detailed performance of CRISPRon-BE compared with other base editor predictors.