

Supplemental Figure 14. Representative Sanger sequencing chromatographs of edited hPSCs enriched using XMAS-TREE and RoT-based approaches. Sanger sequencing chromatographs of Site-3 and PSEN1 of unsorted hPSCs as well as mCherry-positive and mCherry-positive/GFP-positive cells isolated using RoT-based and XMAS-TREE strategies, respectively.