Supplemental Figure S9. B. Litter Progression from Founder #25 (D-line)

L denotes Litter #
Supplemental Figure S9 (S9A, S9B). Expression of cell-type- and function- specific transcripts in captured and captured nuclei from BAT.

qRT-PCR analysis was performed on cDNA prepared from captured adipocyte nuclei from BAT (CAP) and uncaptured nuclei (UnCap-BAT). Nuclear mRNAs were assayed for transcripts encoding (A-F) Cell-type and function markers. SDHA was used as the endogenous control as it was nearly equivalently expressed in captured and uncaptured samples relative to the amount of input cDNA. Bar graphs show the Mean ± SEM with p<0.05 *, p<0.01 **, p<0.001 *** indicated for selected comparisons.
Supplemental Figure S9C, D (Continued). Expression of cell-type- and function-specific specific transcripts in captured and captured nuclei from BAT.
Supplemental Figure S9E, F (Continued). Expression of cell-type- and function- specific transcripts in captured and captured nuclei from BAT.