

SupTable 1: Differentially expressed miRNAs between NE and BNE from Selection sample set A and B

SupTable 2: Differentially expressed miRNAs from microarray profiling of Selection sample set A

SupTable 3: Differentially expressed miRNAs from Nanostring nCounter profiling of Selection sample set B

SupTable 4: Putative miRNA targets of MIR192 and MIR194 identified by cross-referencing predicted targets with miRNAs known to be downregulated in BE.

SupTable 5: Primer sequences used for qRT-PCR and PCR