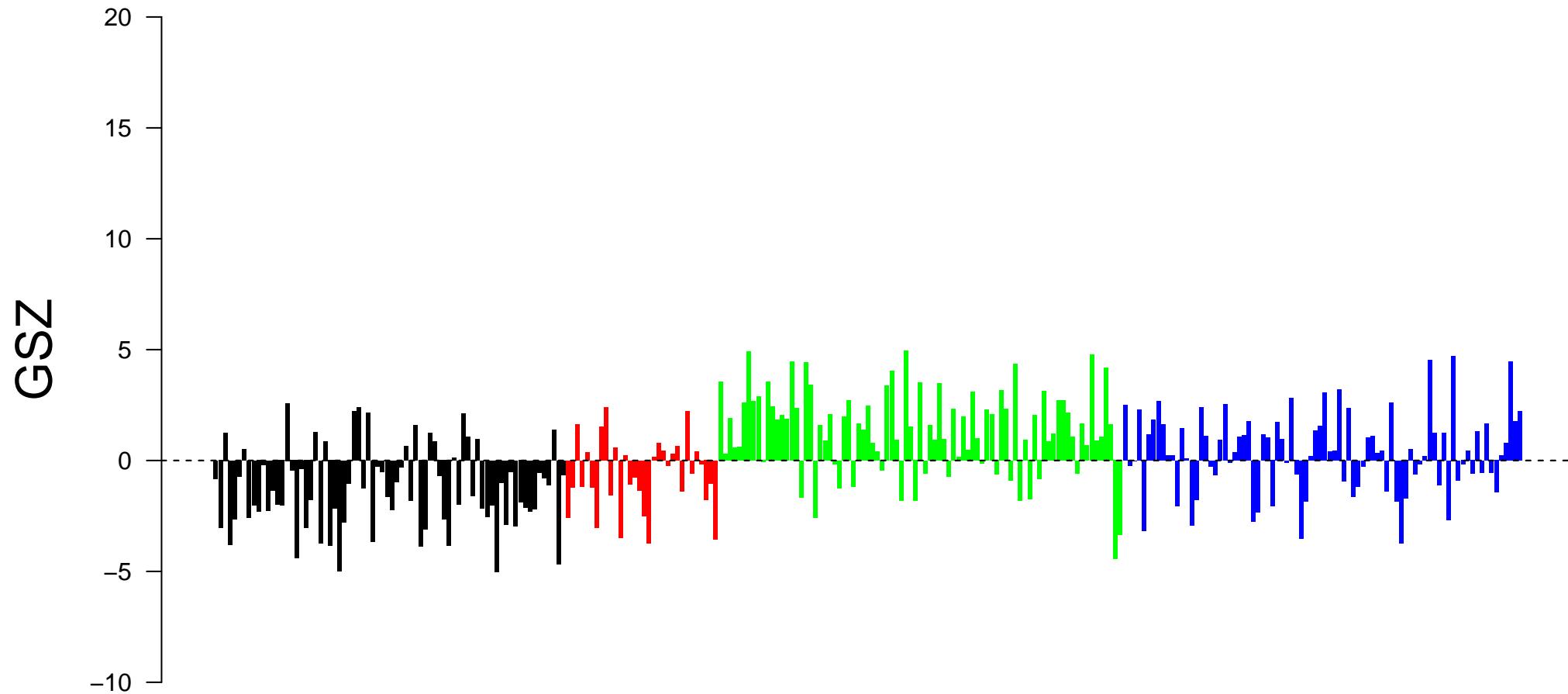
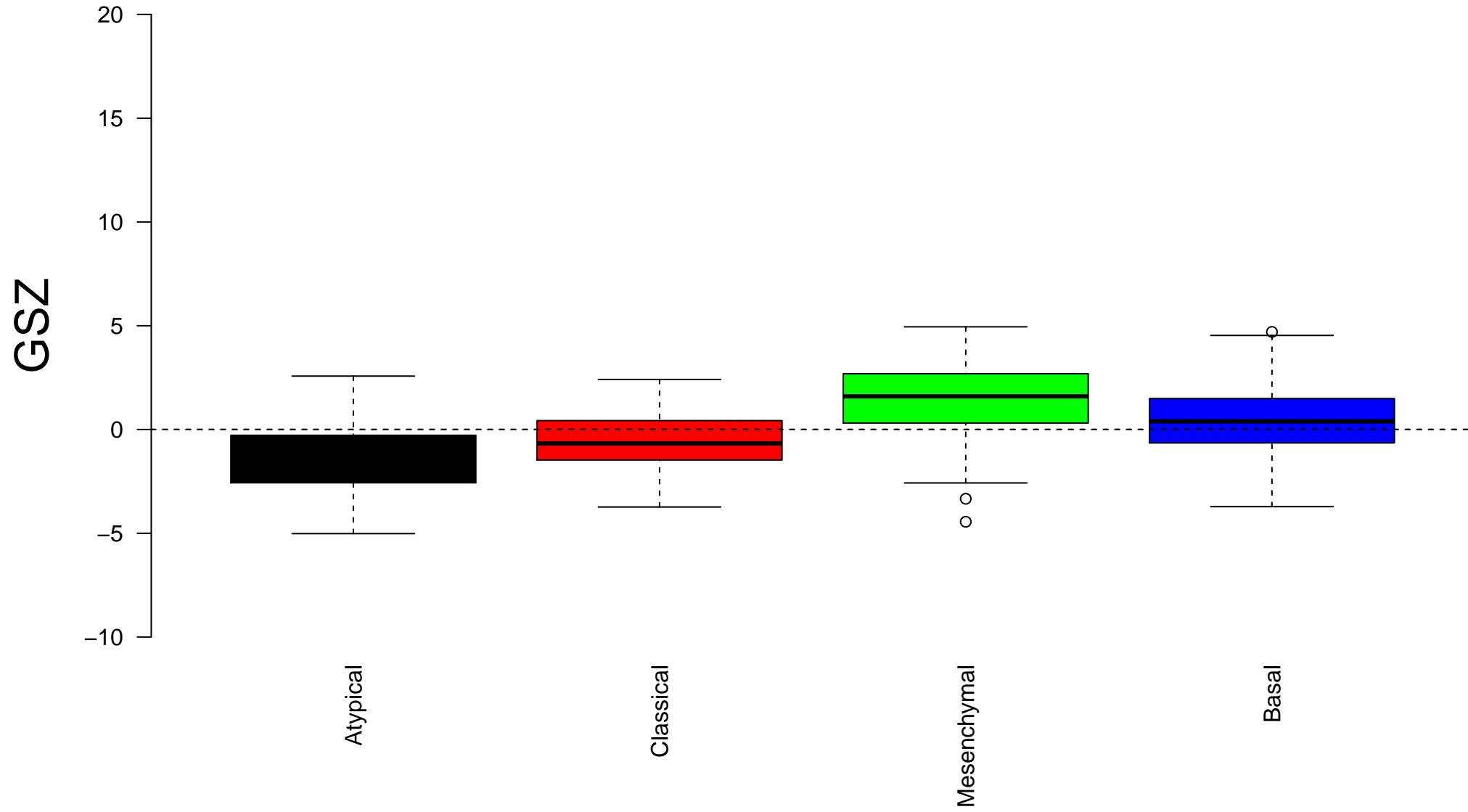


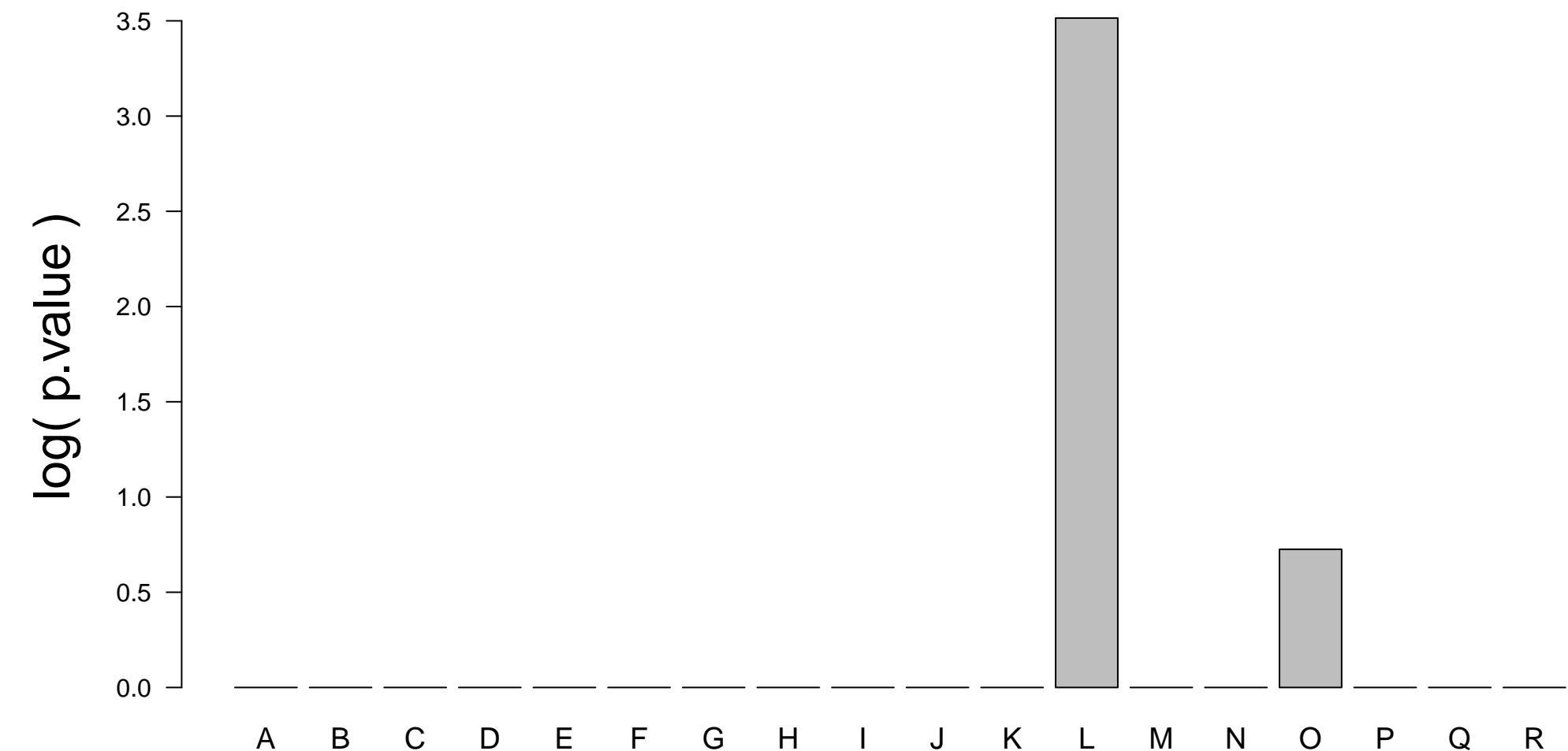
KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors



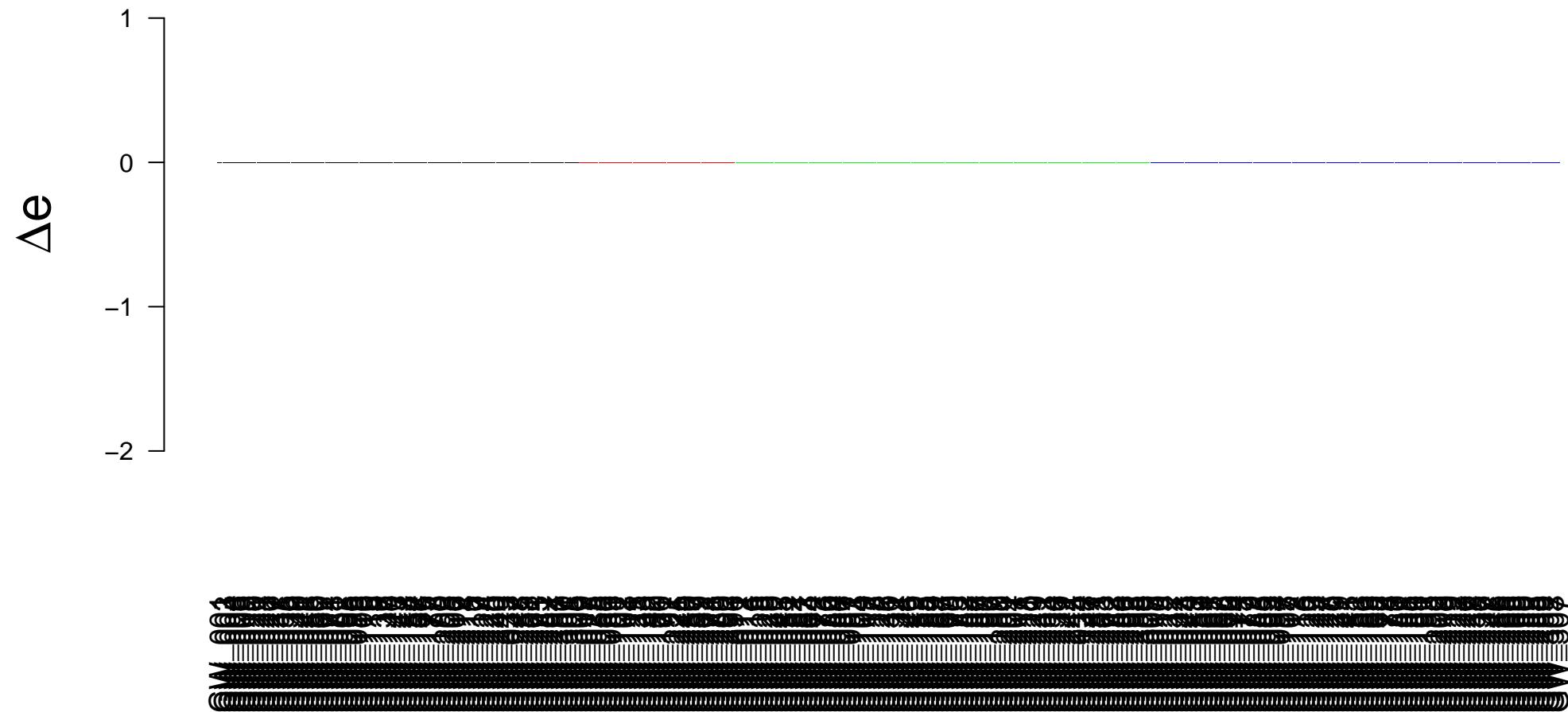
### KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors



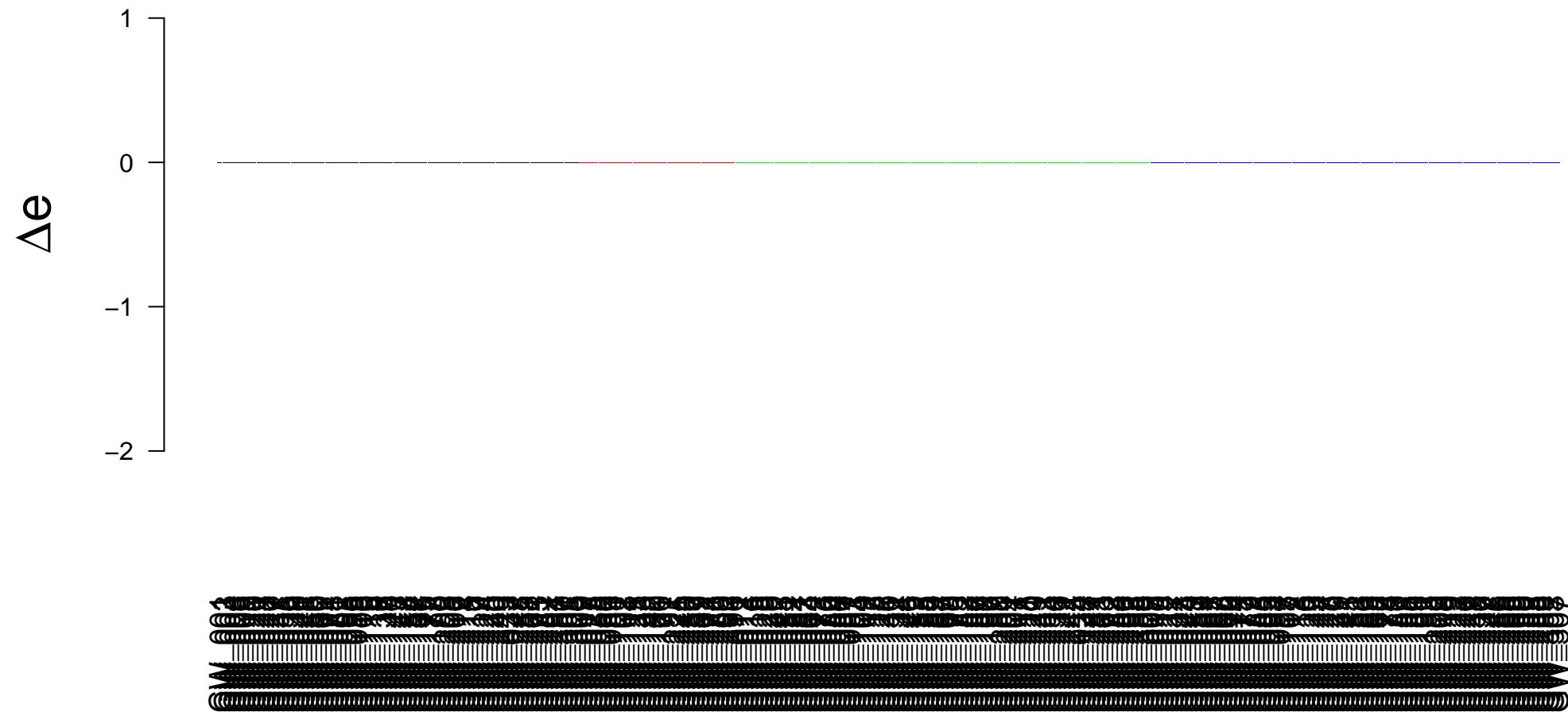
Enrichment in spots: KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivo



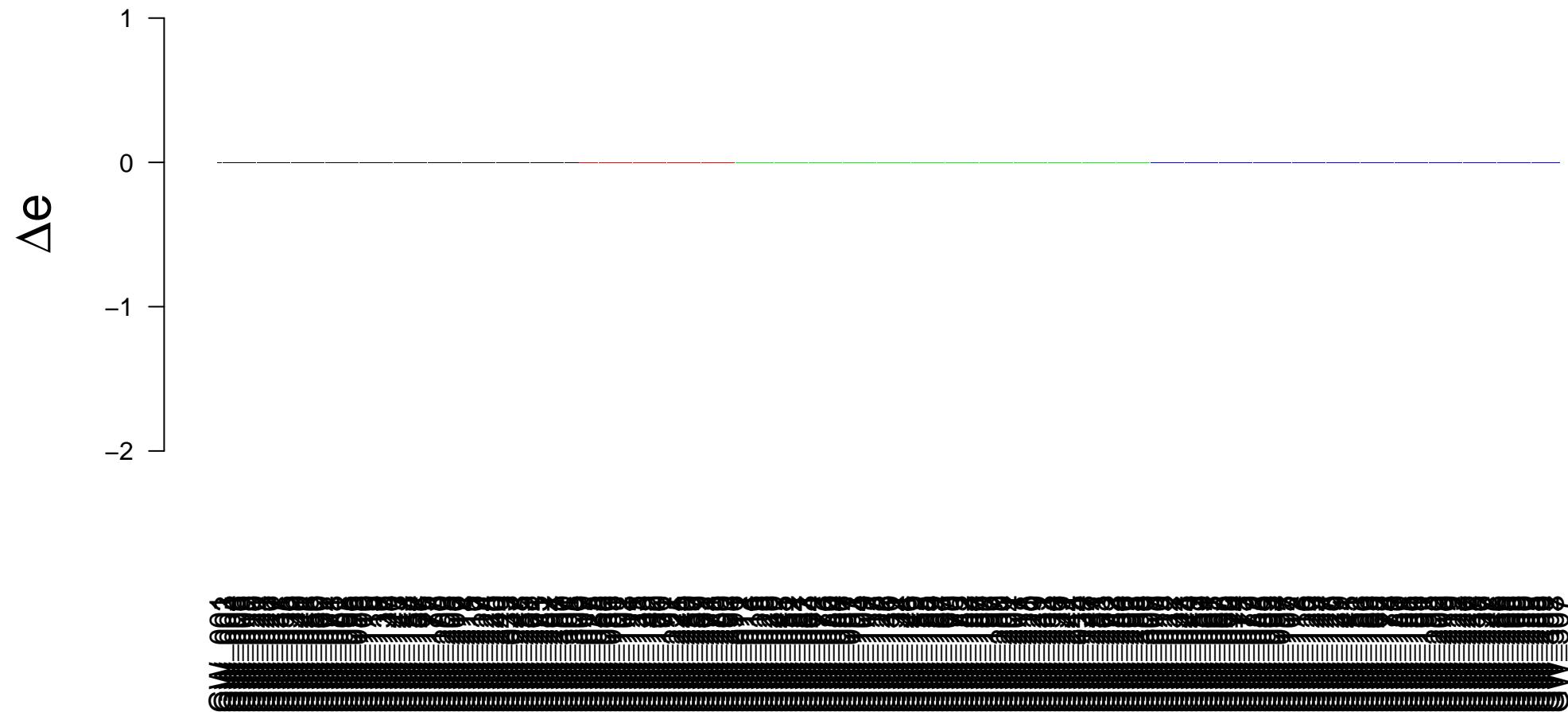
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot A



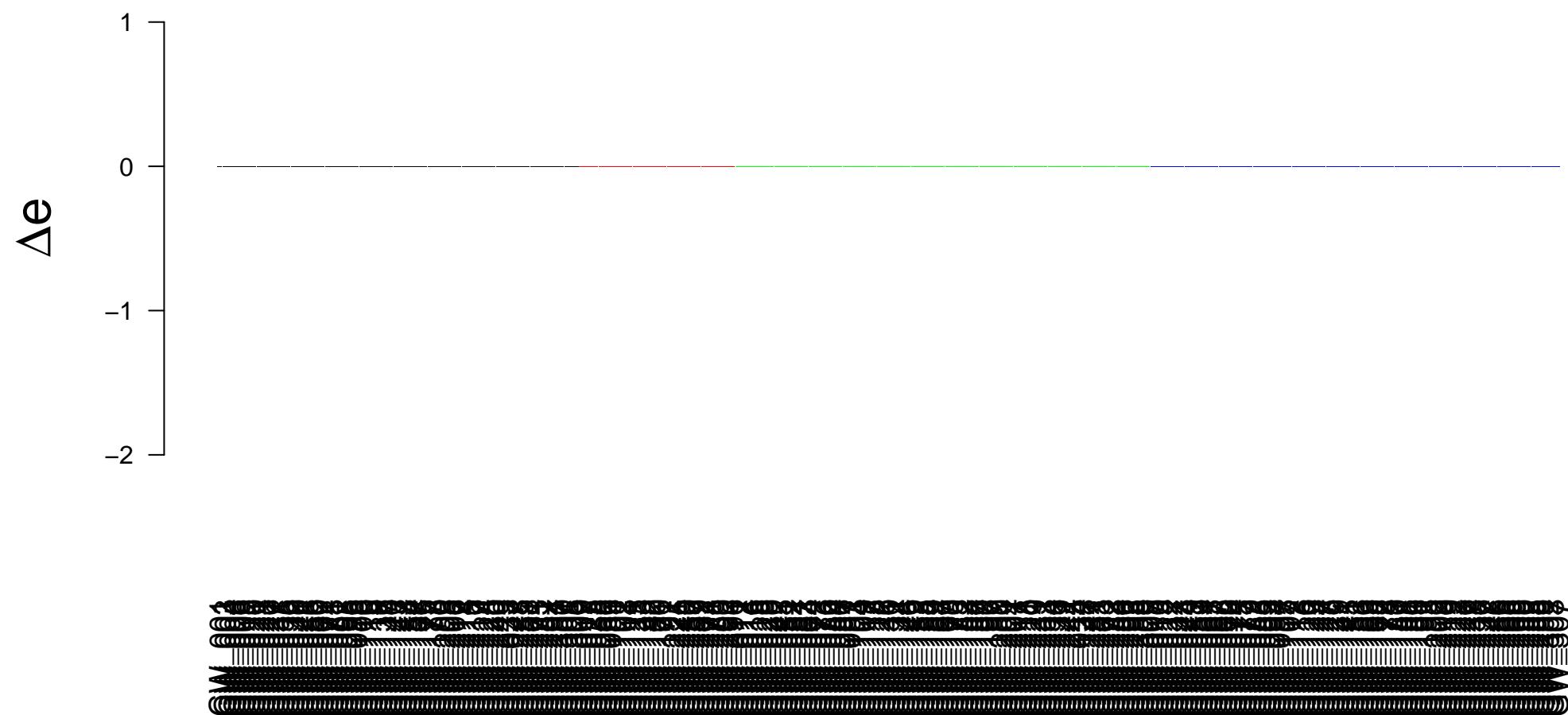
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot B



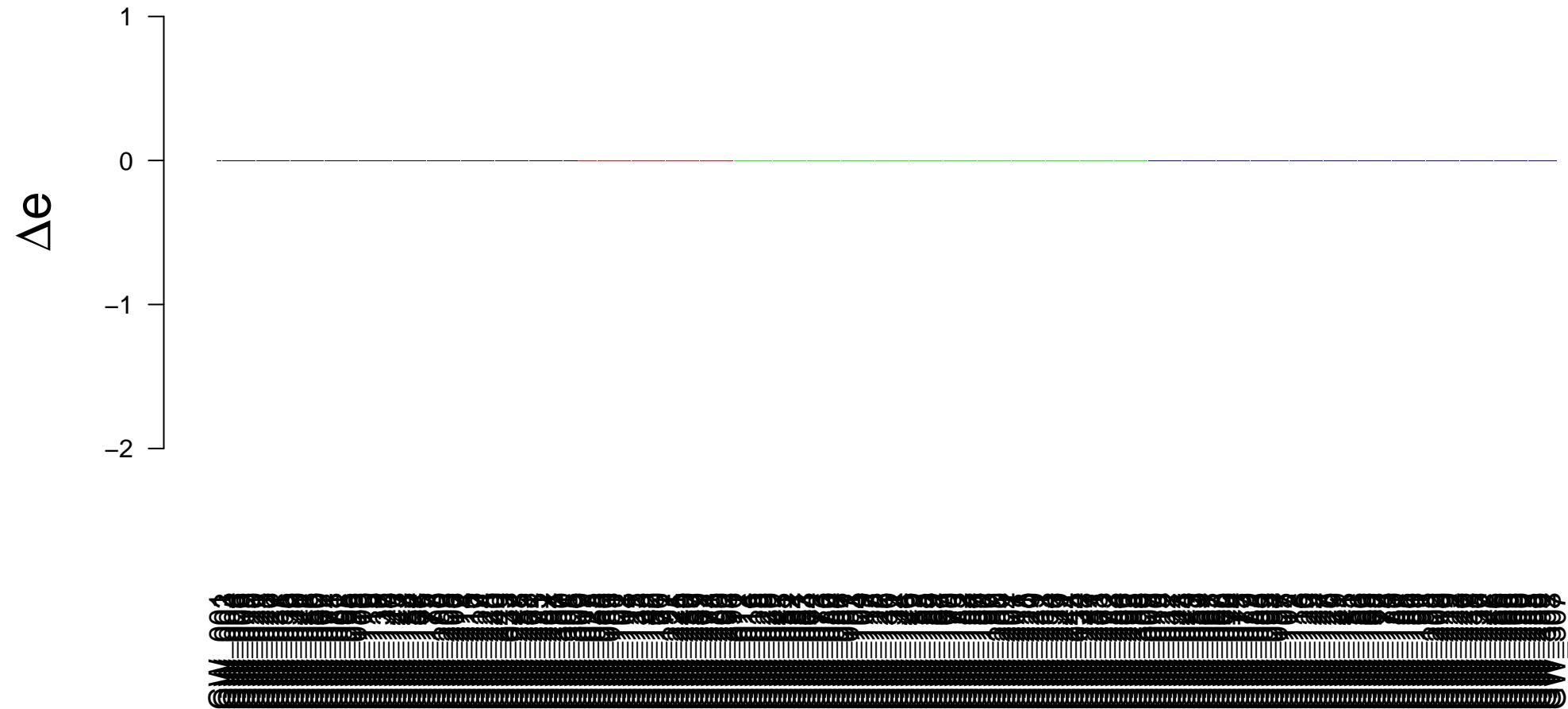
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot C



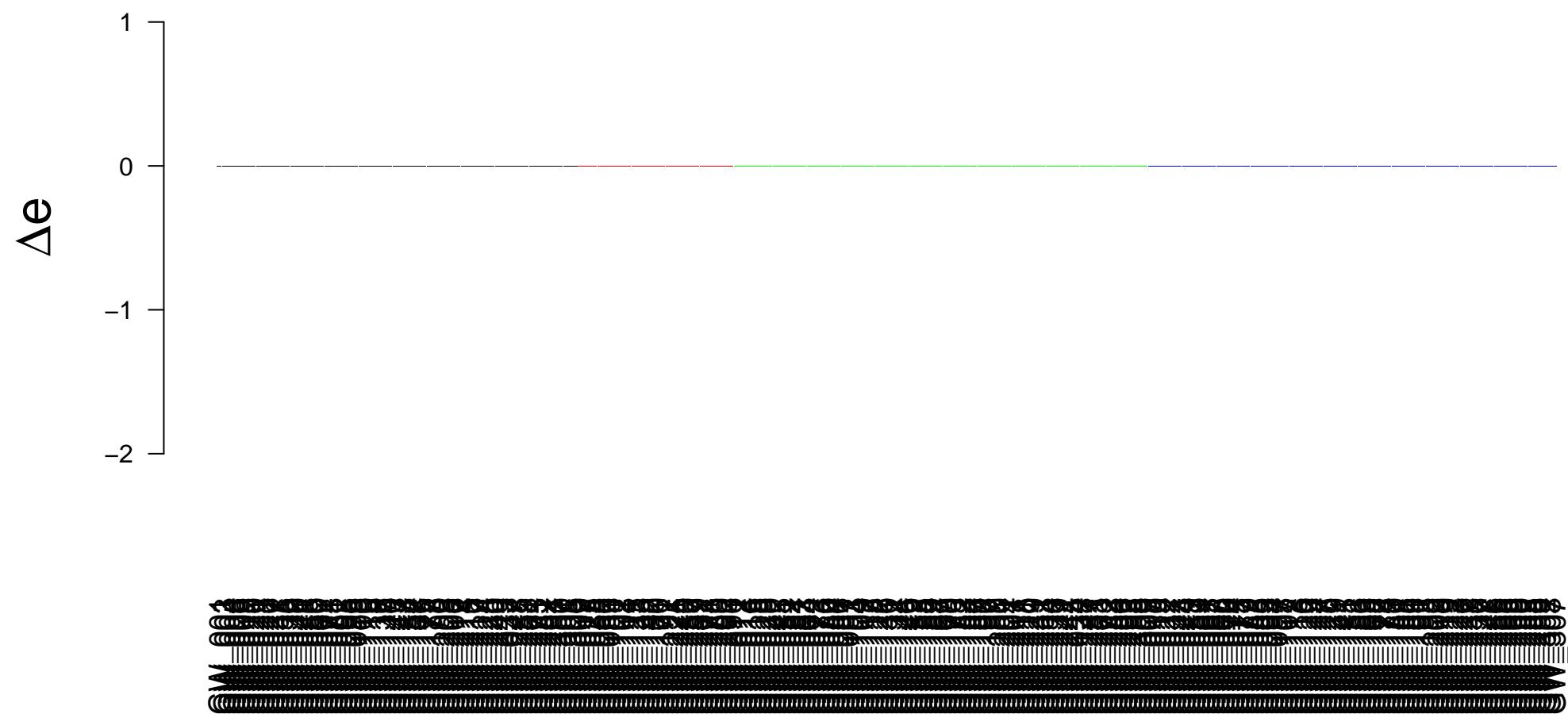
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot D



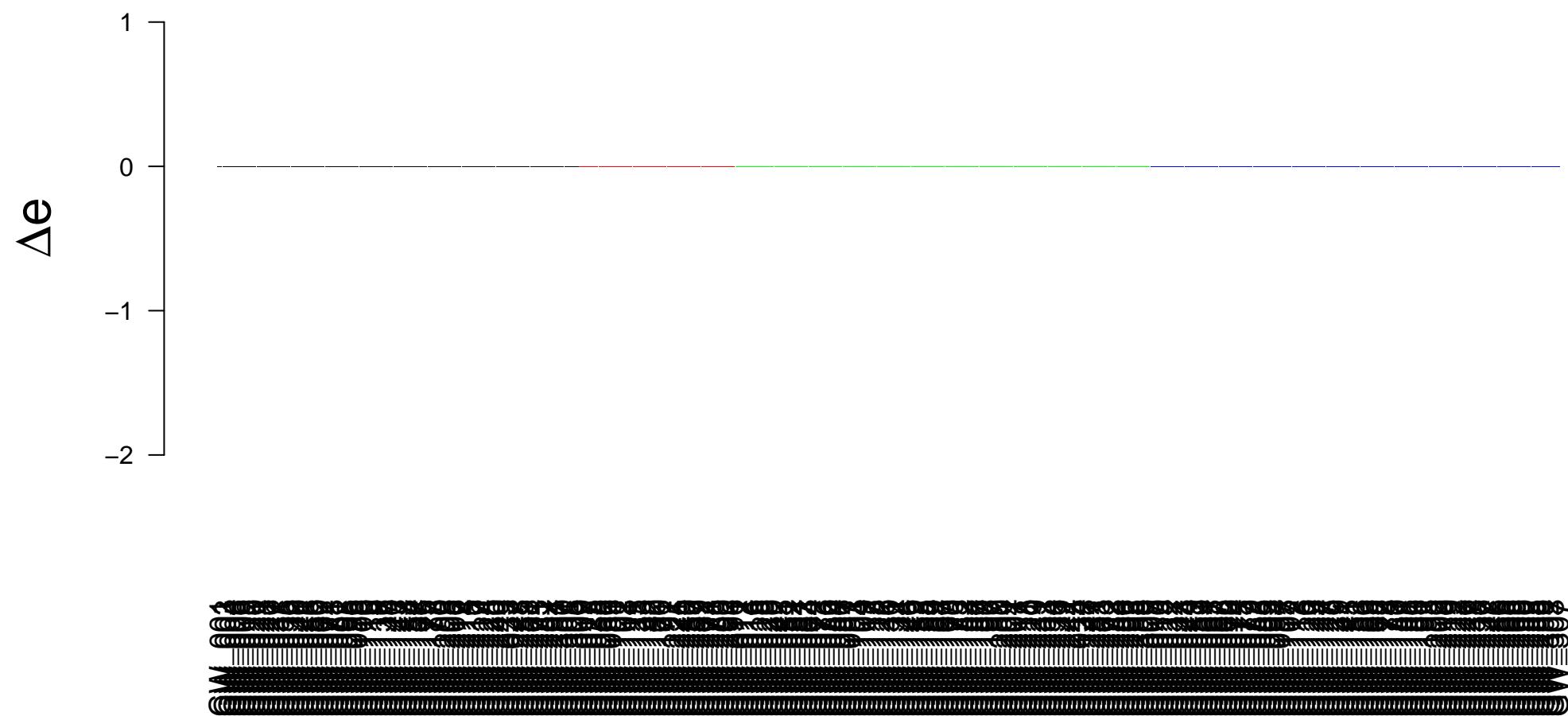
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot E



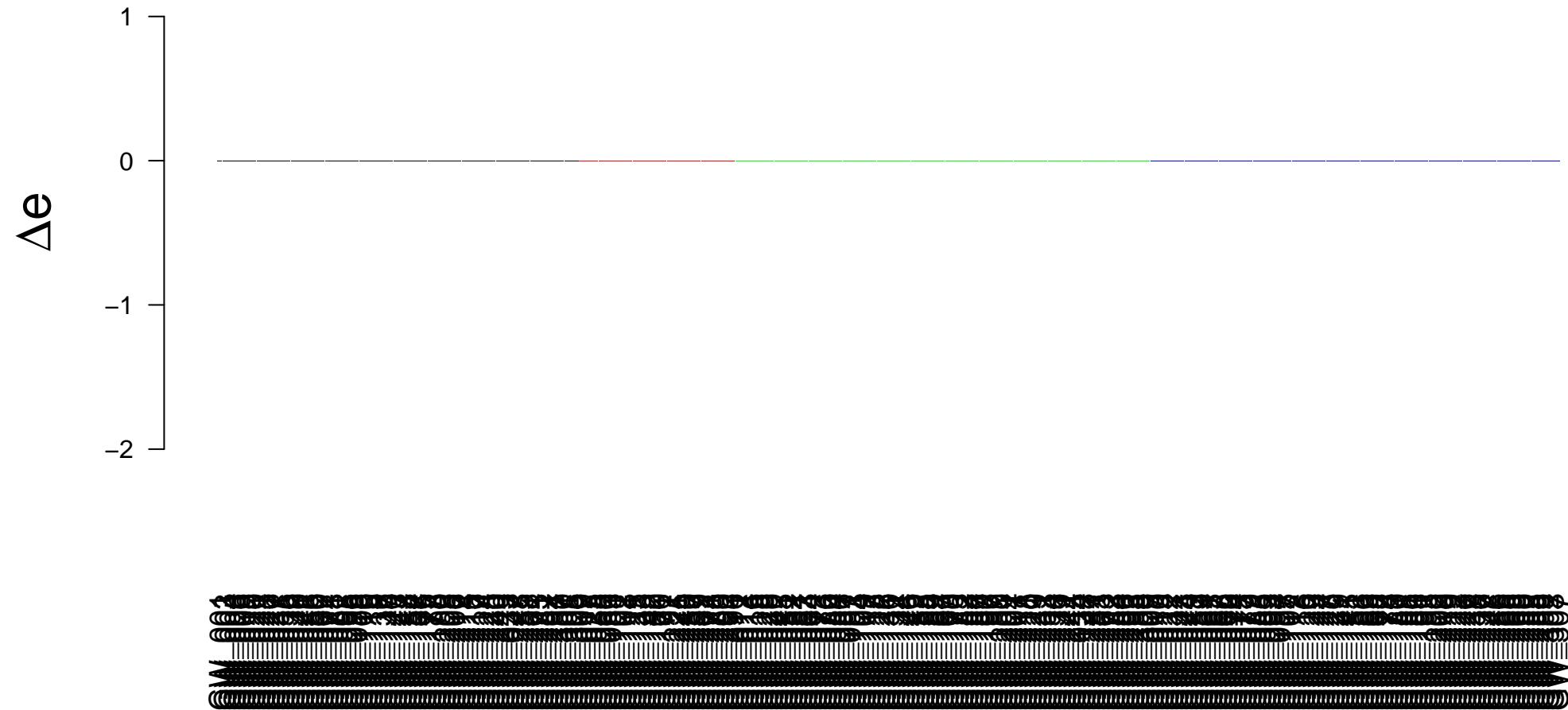
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot F



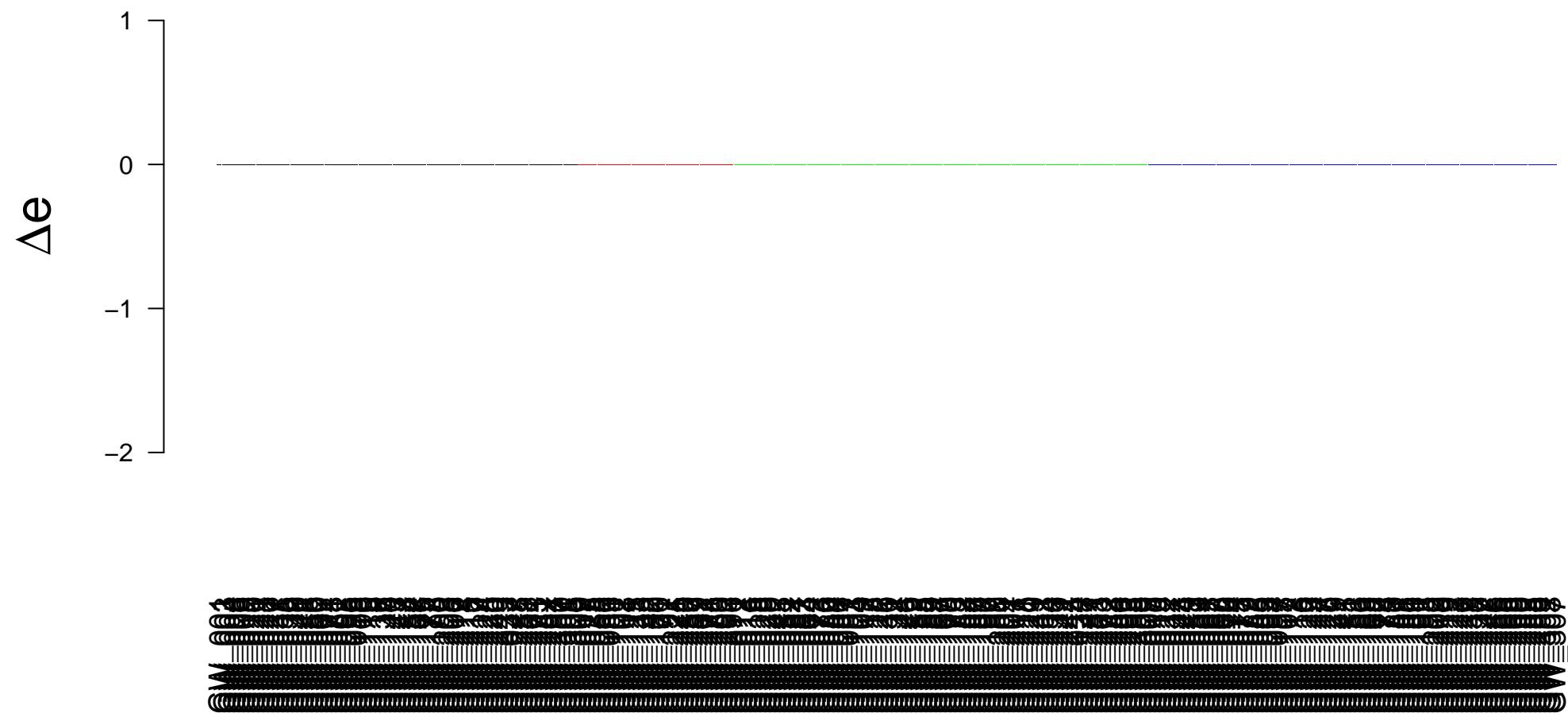
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot G



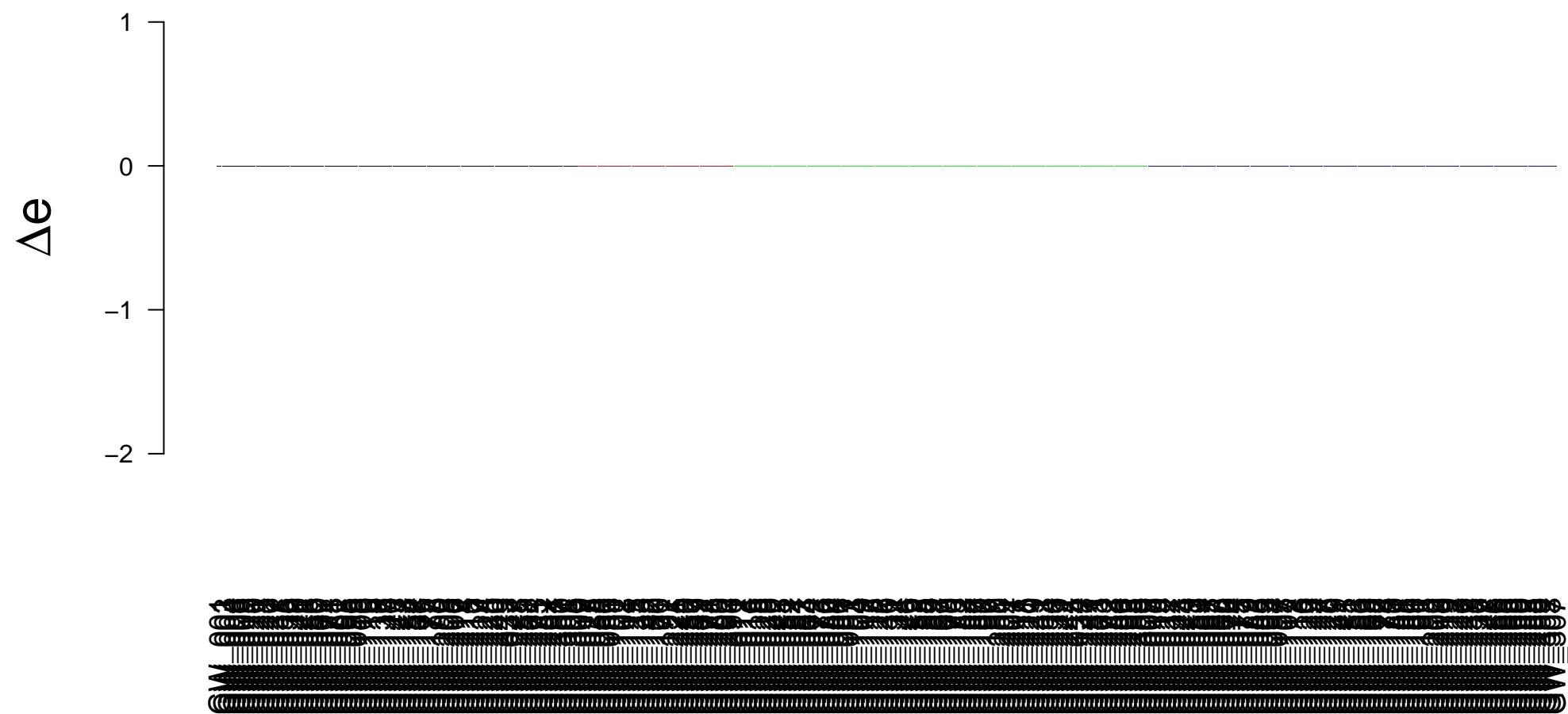
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot H



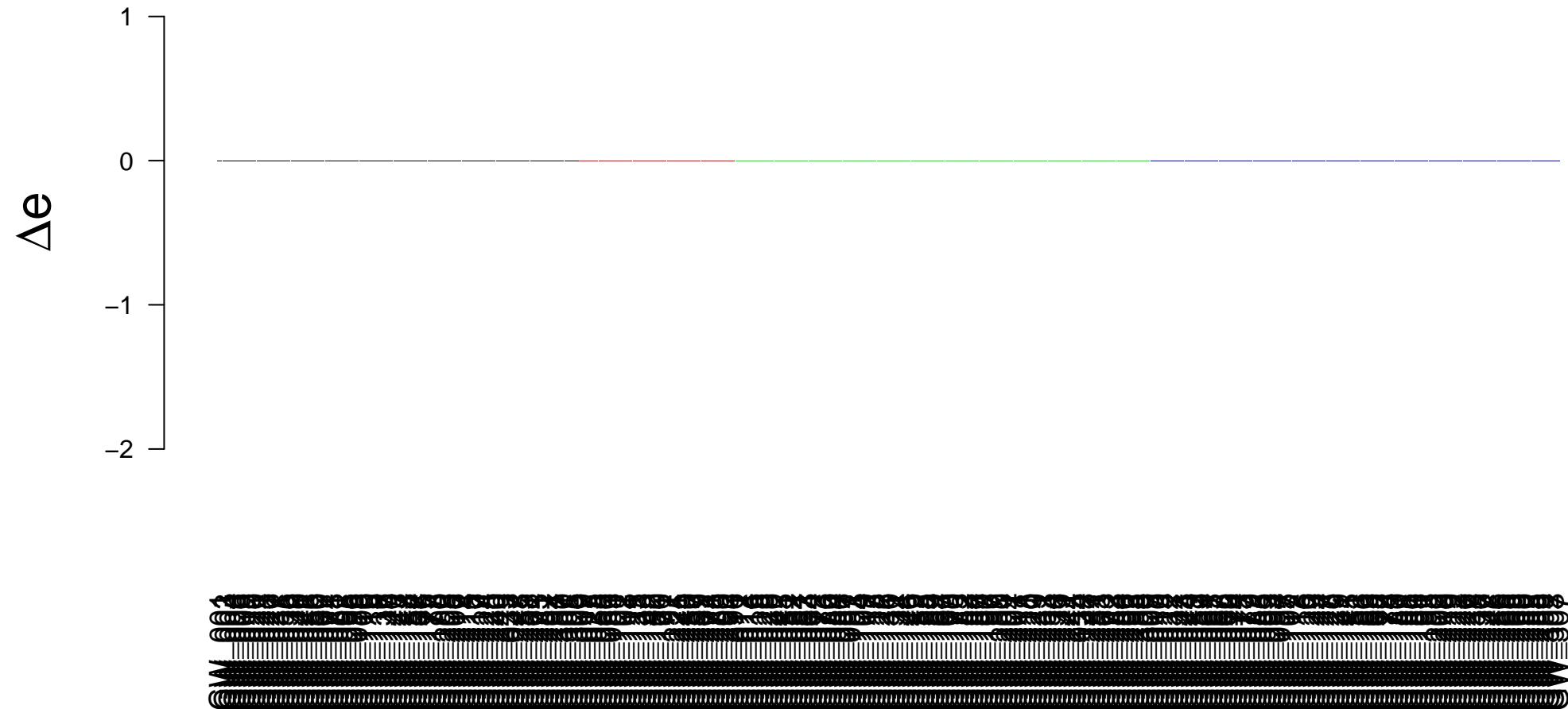
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot I



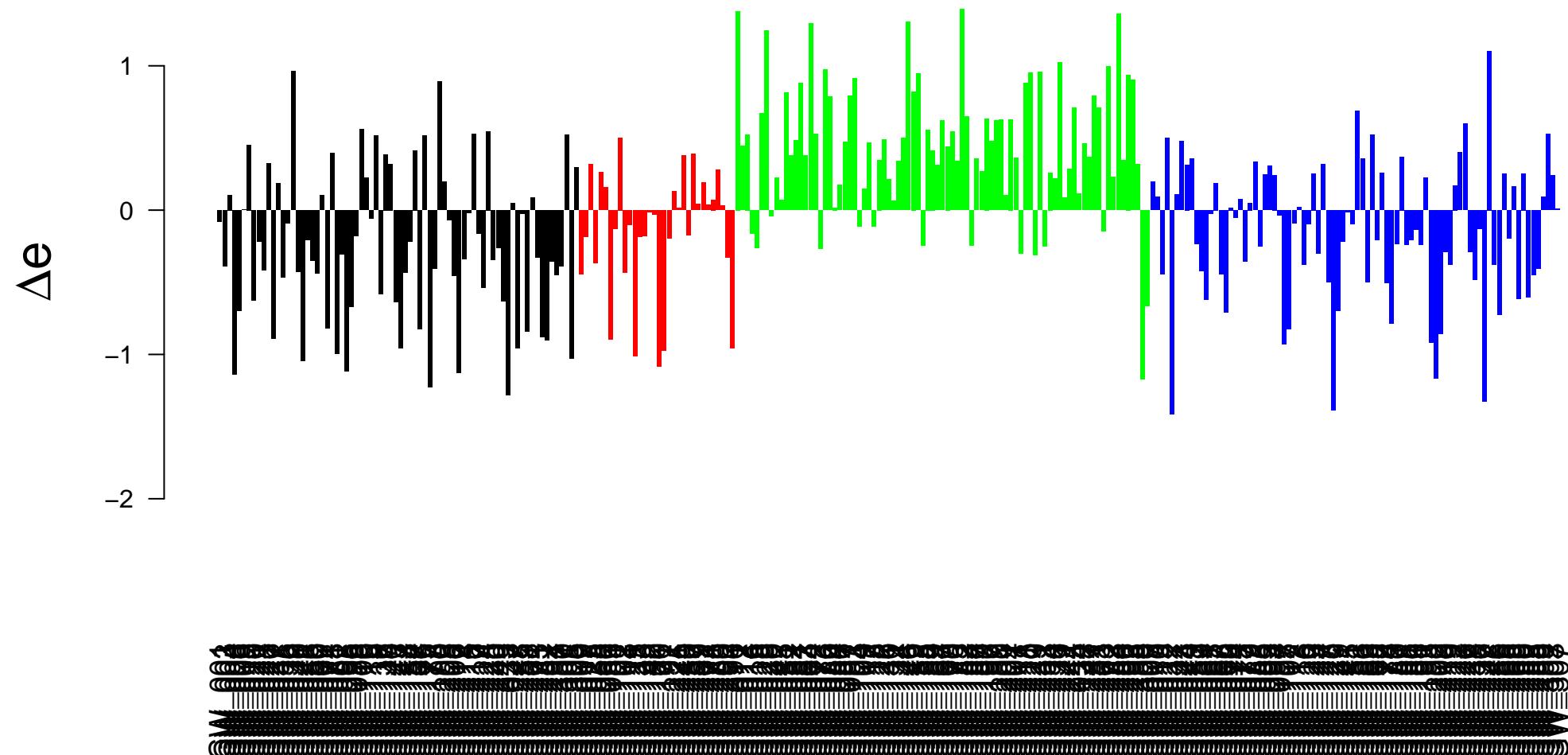
# Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot J



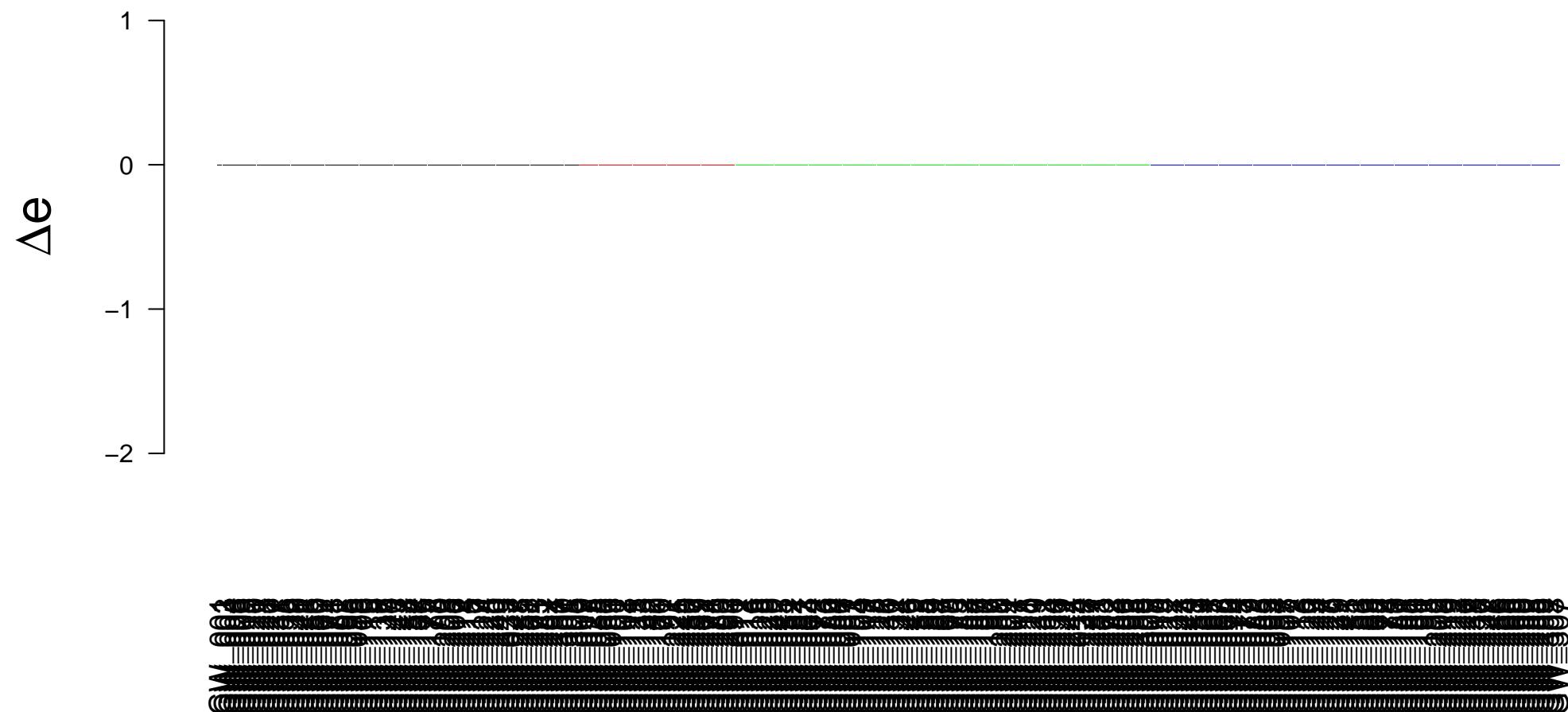
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot K



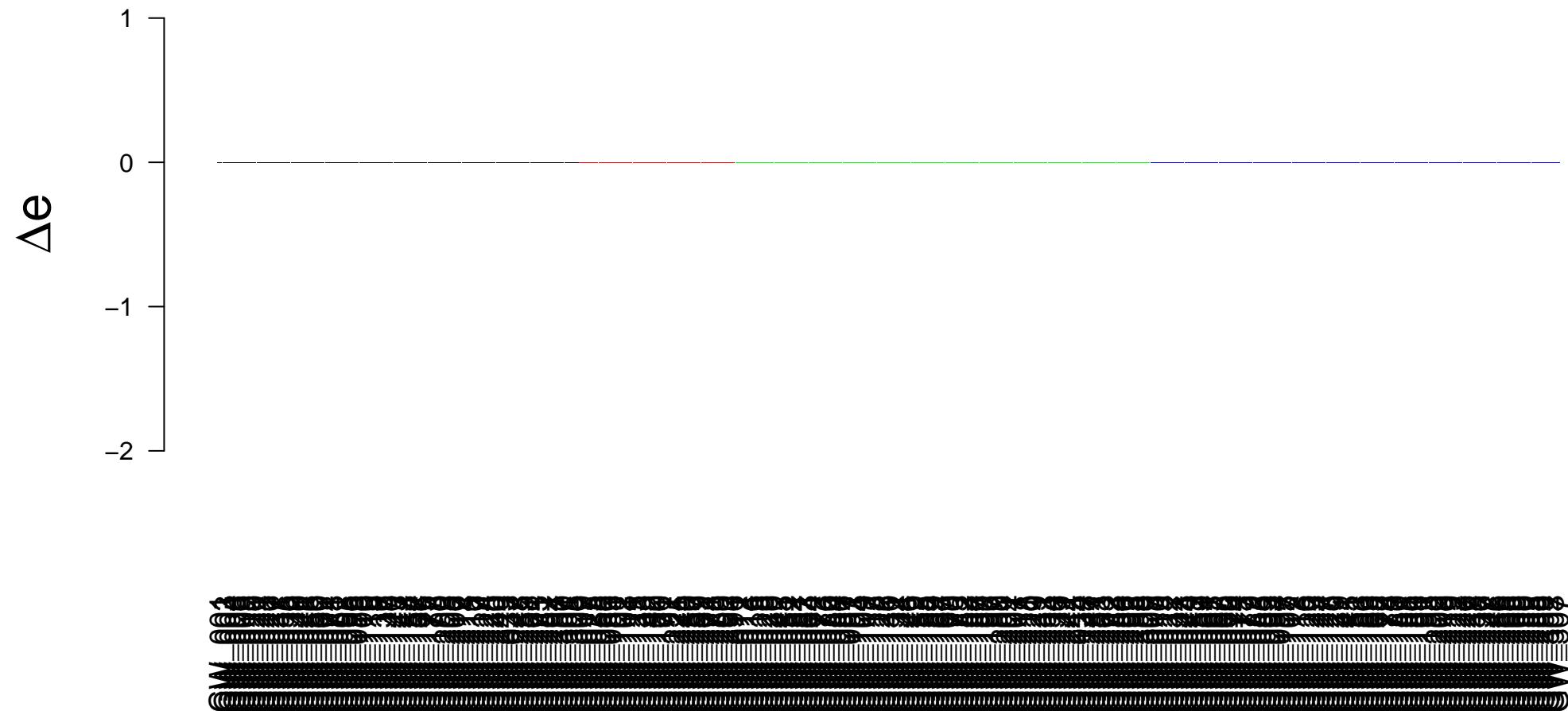
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot L



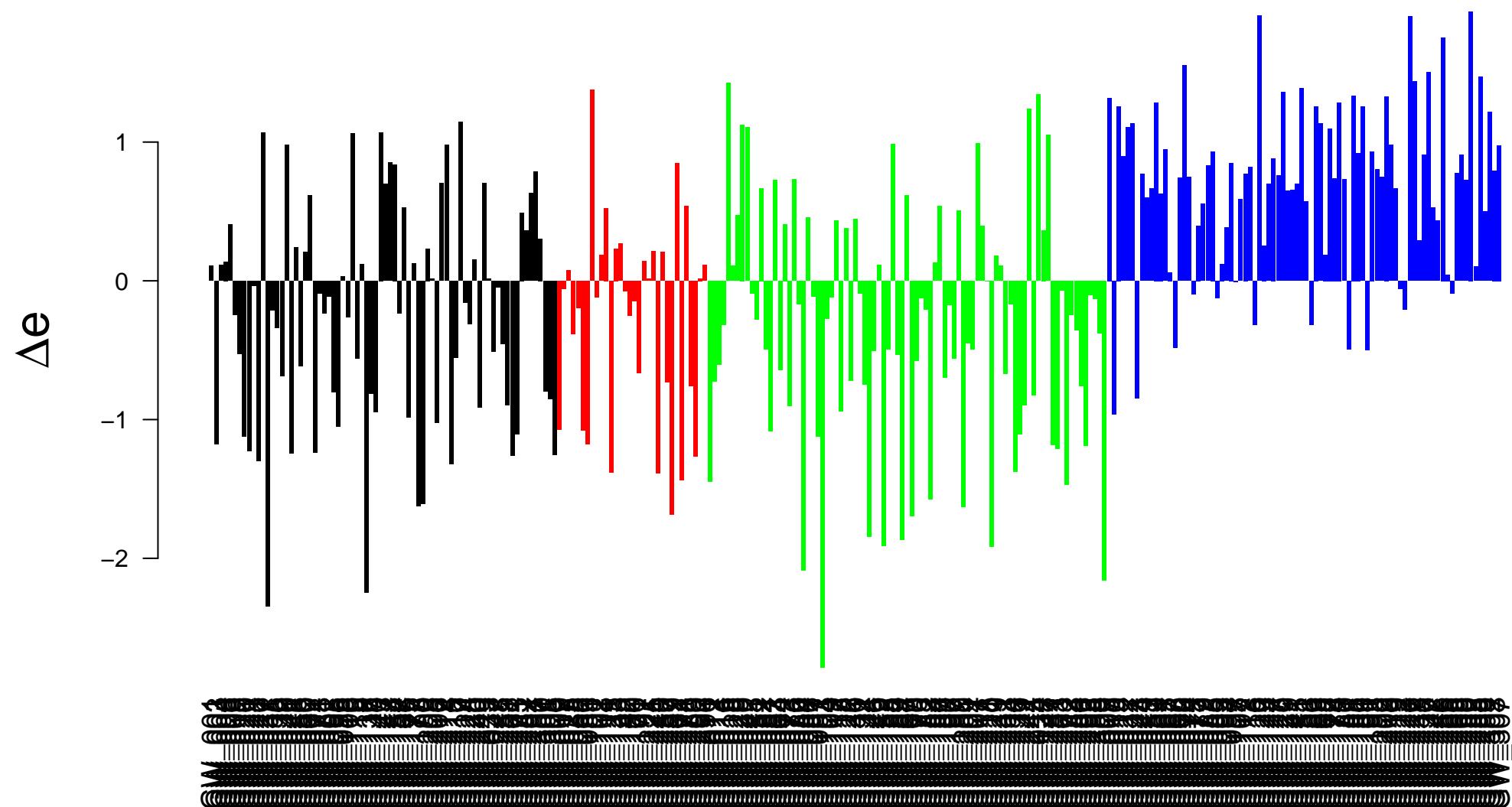
# Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot M



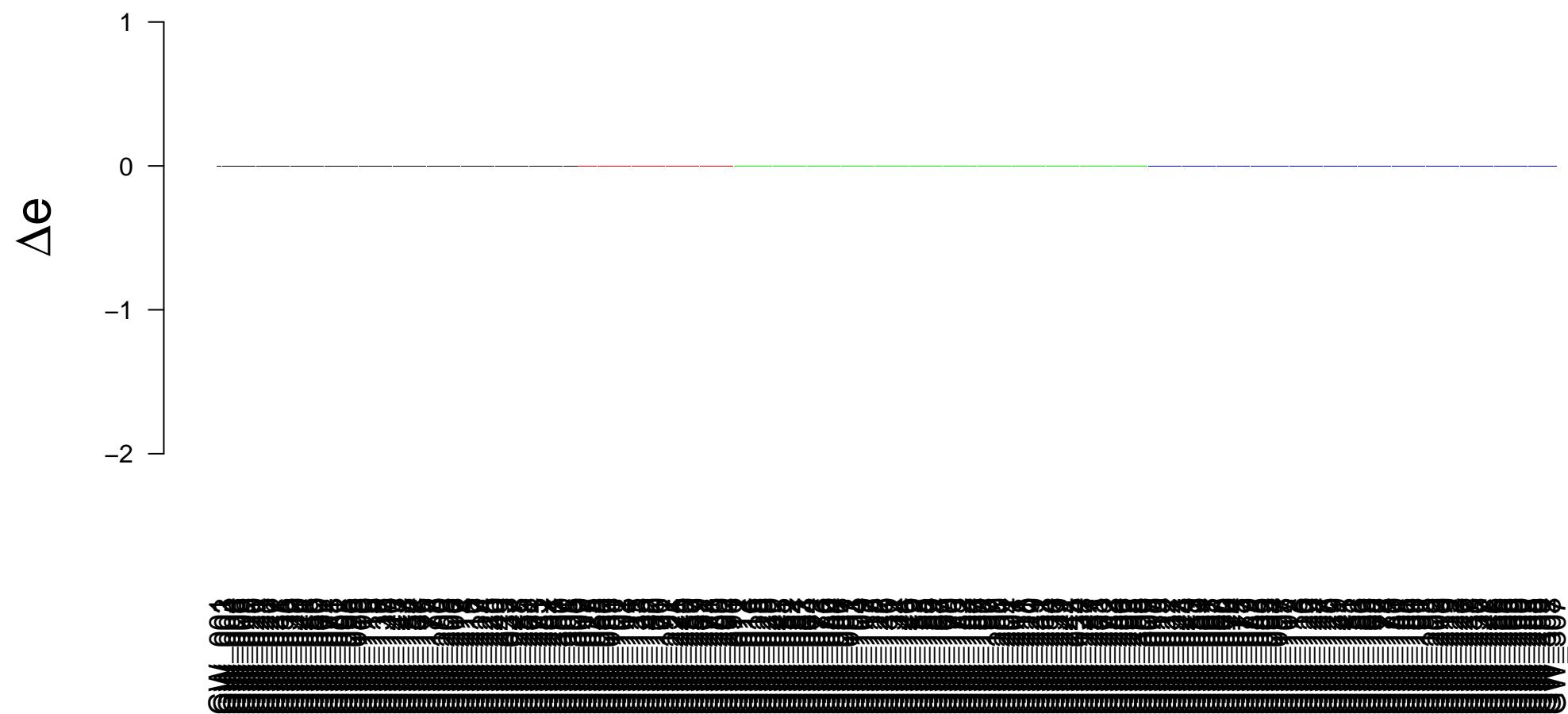
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot N



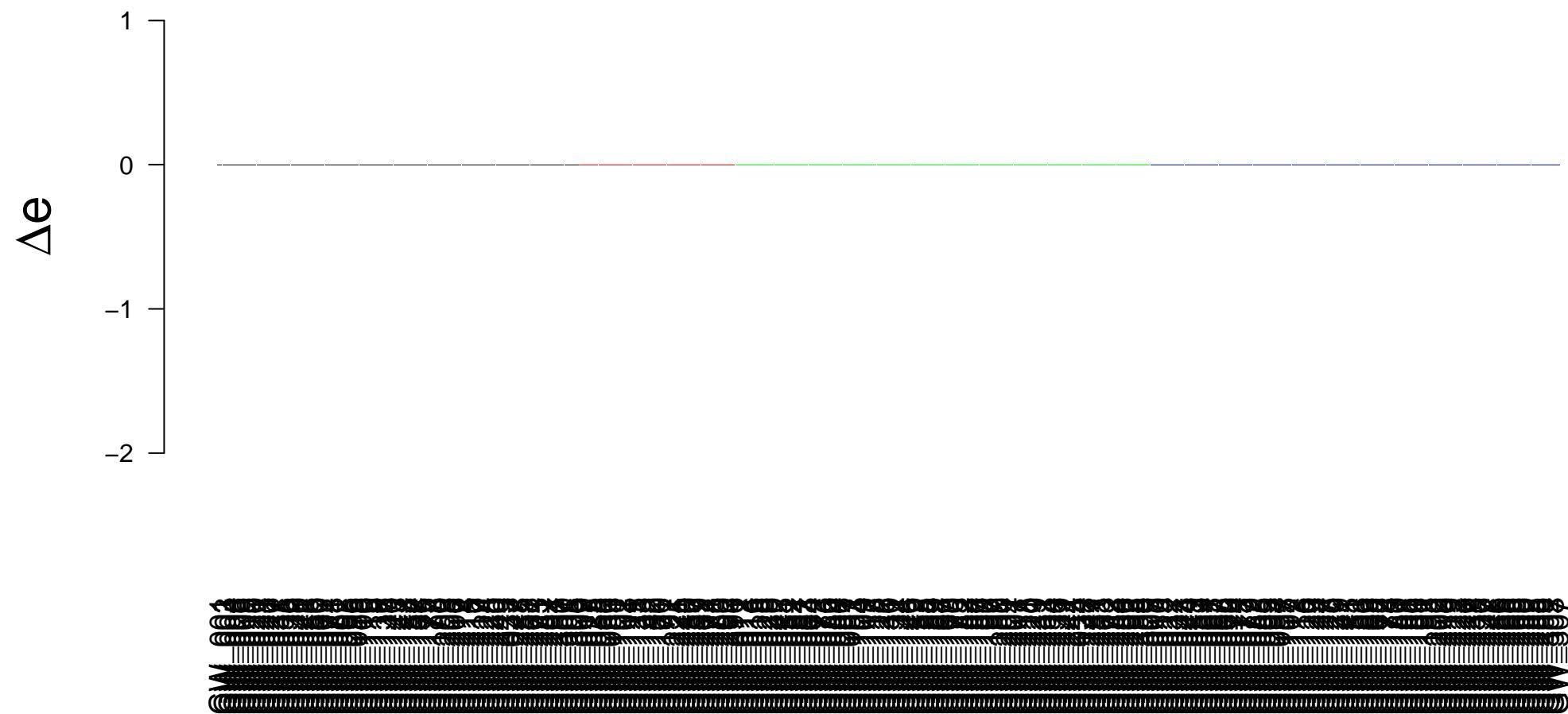
Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot O



Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot P



Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot Q



Expression of KIM\_epithelial–mesenchymal–transition related genes\_decreased expression in (GBM) LongTermSurvivors in Spot R

