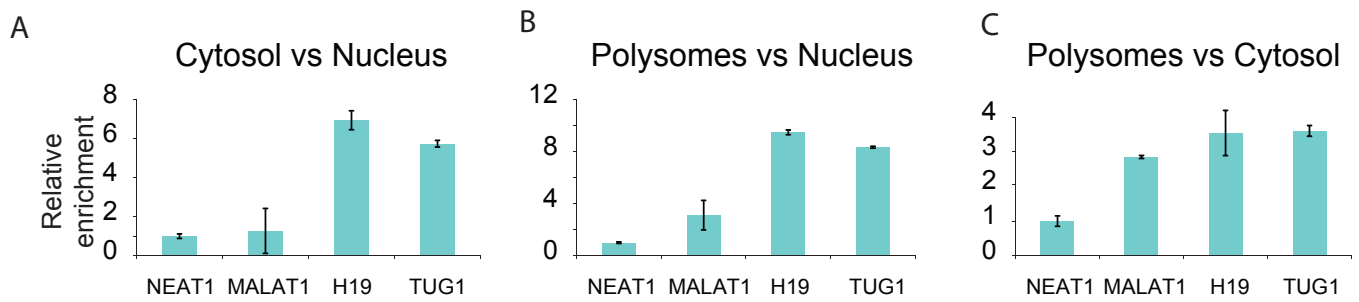




Additional file 9) Subcellular enrichment of specific lncRNAs is confirmed by qPCR.

Relative enrichment of the four tested lncRNAs is represented as $\Delta(\Delta CT)$. CT values were normalized first per sample to H19, resulting in a ΔCT value. Relative differences between lncRNAs, between subcellular samples are determined by calculating the difference between the ΔCT s between samples, resulting in $\Delta(\Delta CT)$ values. $\Delta(\Delta CT)$ values were determined between A) cytosol and nucleus, B) polysomes and nucleus and C) polysomes and cytosol. The lowest $\Delta(\Delta CT)$ is set to 1 (NEAT1 in all panels). Error bars depict the standard deviation of three different primersets used per lncRNA.