

Figure S1. Correlation plot of sno/scaRNAs expression levels (red bar, right scale) and corresponding inferred CN (point, left scale). For each sno/scaRNA localization and Kendall corrected p-value are reported.

Figure S2. SCARNA22 and MMSET expression levels in the cohort of 55 MMs and 8 sPCLs (A) and in 18 Human MM cell lines (B).