Unsupervised clustering of tumor samples. Hierarchical clustering of probes passing unspecific filtering, i.e. IQR between tumor samples and expression above the background in at least four arrays.

(A) Hierarchical cluster tree of probes located in exons of protein-coding genes (N = 12061), and (B) of non-coding probes (N = 7263). Variance within clusters was minimized by applying Ward’s method on scaled log2-transformed data. Red numbers indicate cluster reliability in percent, here.

Detailed description of clinical, pathological and immunohistochemical data of presented tumor samples is provided in caption of Supplemental Table S.