Supplementary Figure 4. Somatic copy-number alterations in the investigated samples for the most affected genes.
Whenever multiple tumor samples were available from the same patient, appropriate sample names are shown in blue. Genes listed in the NCCN guidelines for genetic testing of cancer are marked with red. (List of genes are presented in detail in Supplementary Table 2.) Tumor content of the samples estimated by Sequenza are presented as row annotations.