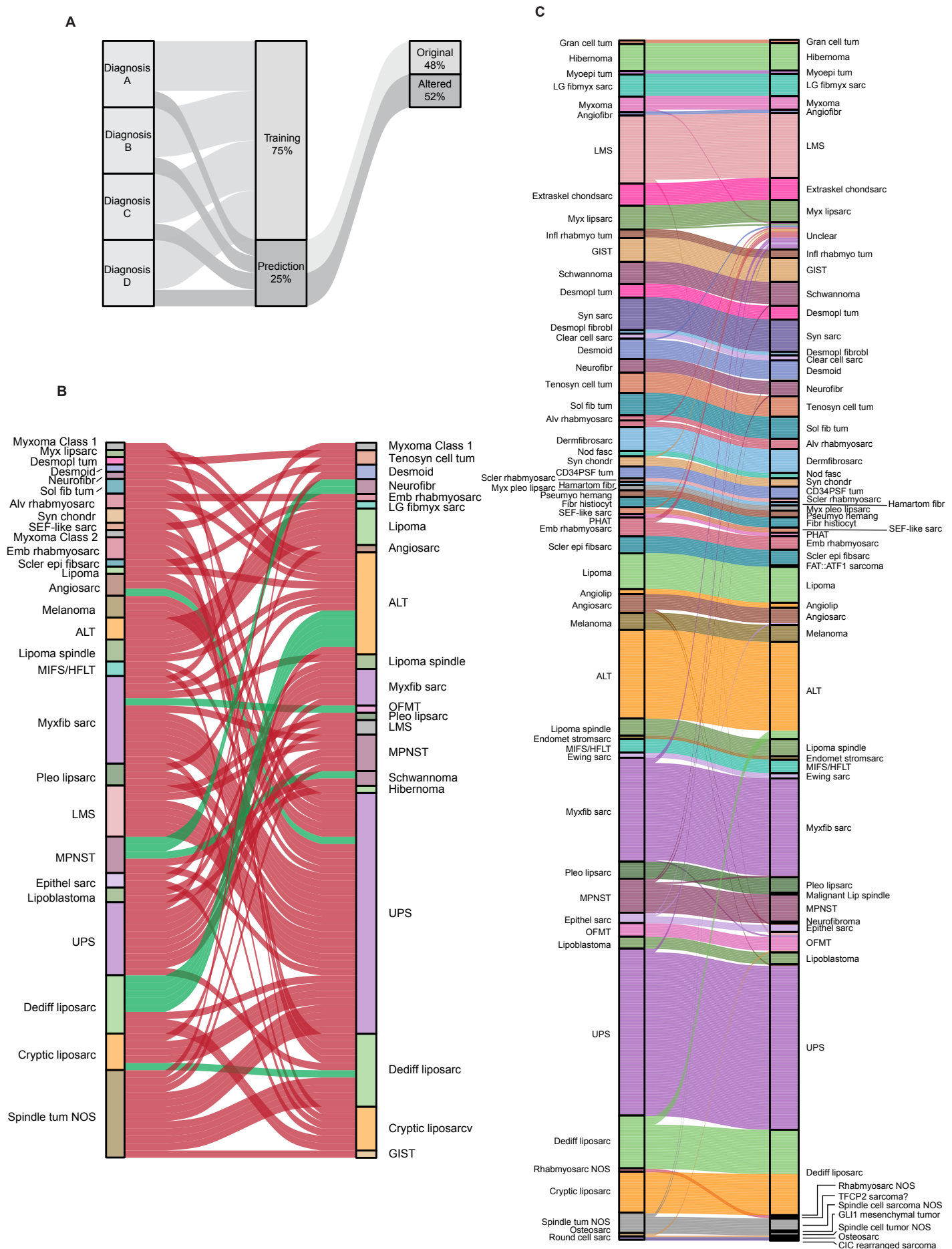




Supplementary Figure S7: Gene expression based classification of soft tissue tumors

(A) Schematic overview of the random forest classifier approach to outlier identification. (B) Alluvial plot illustrating the 98 samples where the classifier-predicted diagnosis differed from the institutional diagnosis. Green lines indicate samples where the predicted diagnosis was in agreement with the revised diagnosis and red lines indicate samples where the classifier was unsupported. (C) Alluvial plot of all 704 samples with the original institutional diagnosis on the left and the revised diagnosis on the right. For abbreviations of tumor types, see legend to Fig. 1.