



Supplementary Figure S2. Kaplan-Meier plots displaying correlations of processing mutations to OS in the sarcomatoid NSCLC cohort. Only patients with advanced-stage tumour disease (IIIB-IV) were included. A high mutation count constitutes more than one mutation, while low is equivalent to just one mutation. Dotted lines represent the medial OS of each group. The p-value was calculated by score-logrank test.