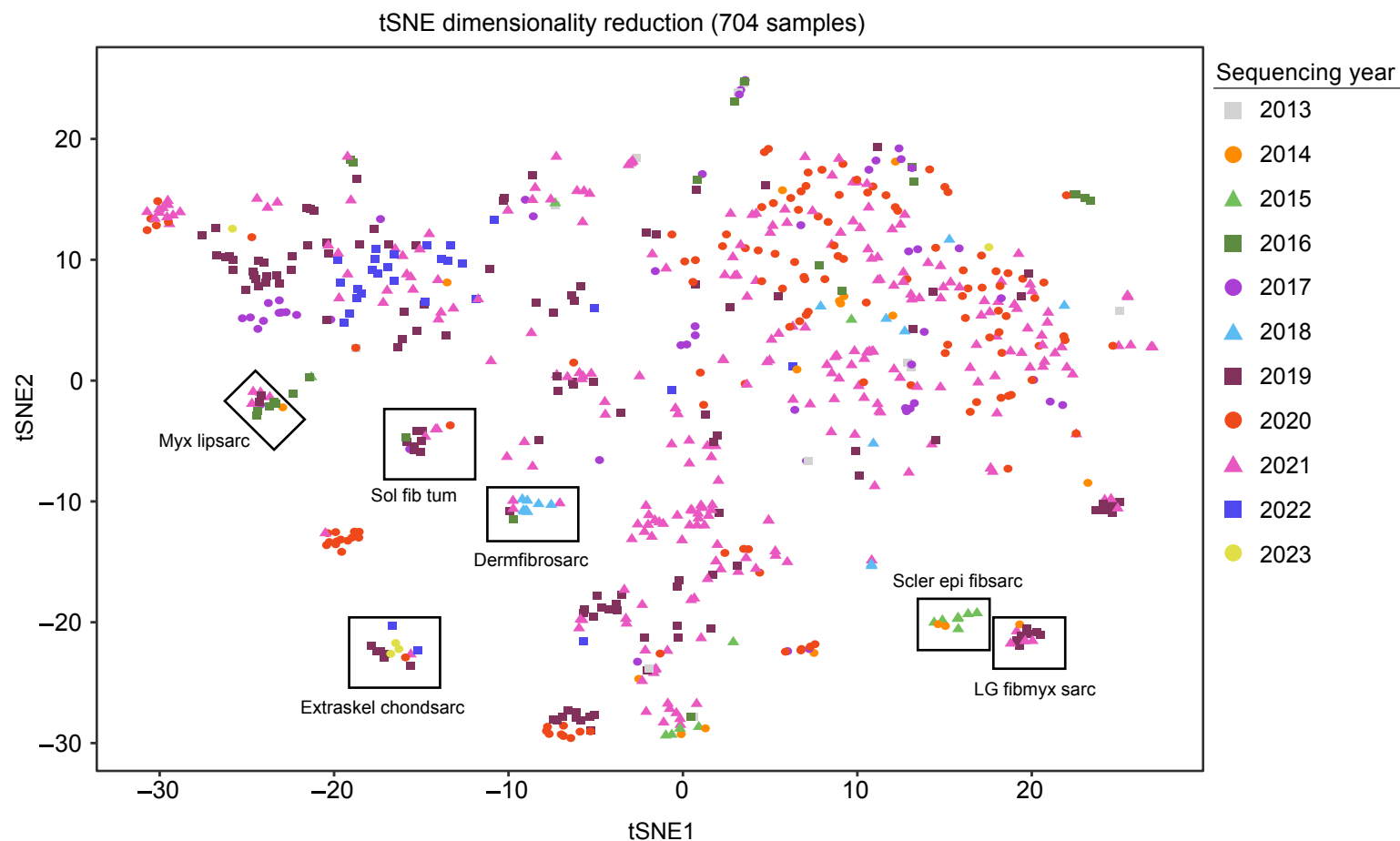




Supplementary Figure S1: Batch effects on transcriptomic clustering

The RNA-seq analyses were performed during an 11-year period, which potentially could have caused batch effects. However, no significant effect of year of sequencing was seen, here exemplified by six tumor types (black frames) that had been sequenced at 2-5 separate occasions.