



### Supplementary Figure 7

**To Figure 4:** Distribution of variant allele frequencies in EATL biopsies. **(A)** Box and whisker plots illustrating the number of mutations or **(B)** mean variant allele frequency (VAF) per sample. Indicated p-values were calculated by ANOVA with Tukey correction for multiple comparisons. **(C)** Scatter graph depicts VAF detected per patient with each dot representing the mean VAF for every single gene called from various runs and techniques for *de novo* EATL and RCDII-EATL as indicated. Black dots indicate VAF from autologous samples without EATL (no-EATL) and red dots from EATL samples. Box for 75th percentile and whiskers ranging from min to max, horizontal lines are median and grey dots indicating mean.