Figure S1

Overview of our approach to identifying variants in SPTs

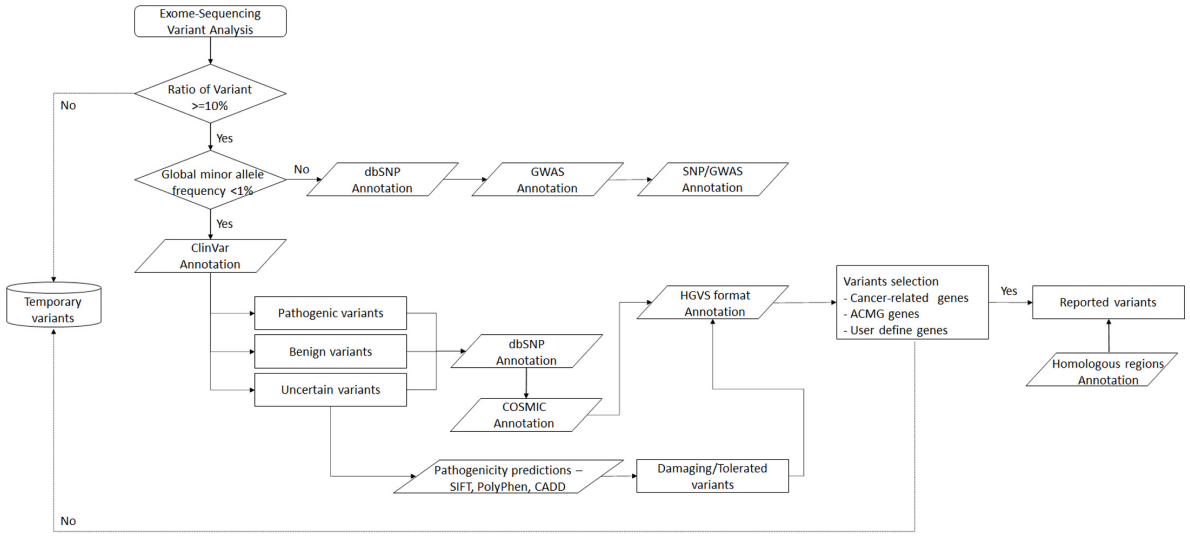


Figure S1. Overview of our approach to identifying variants in SPTs.

Figure S2

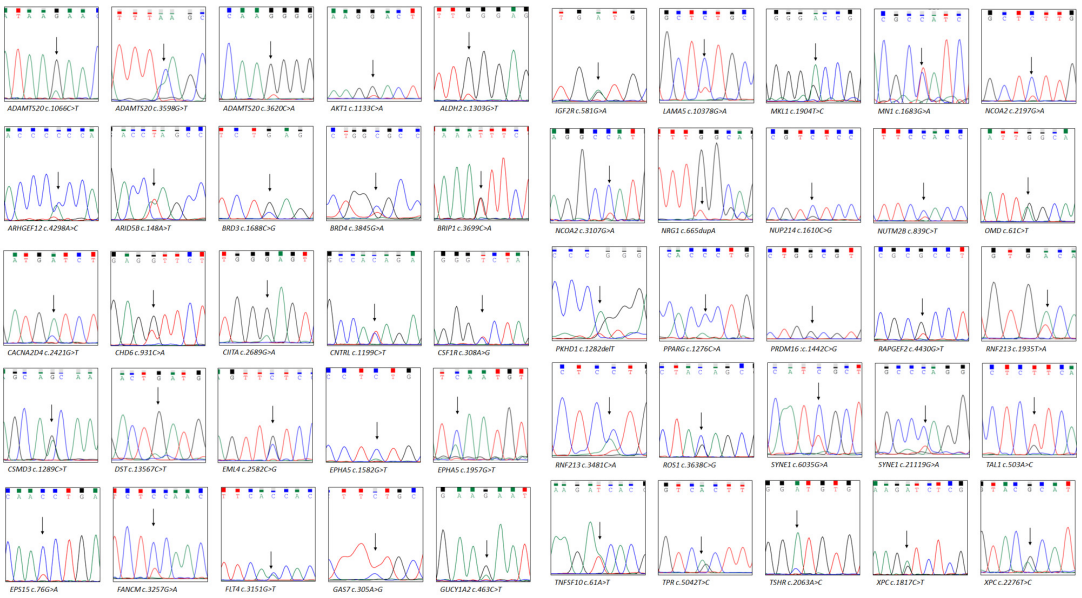


Figure S2. Overview of our approach to identifying variants in SPTs.

Fig S3

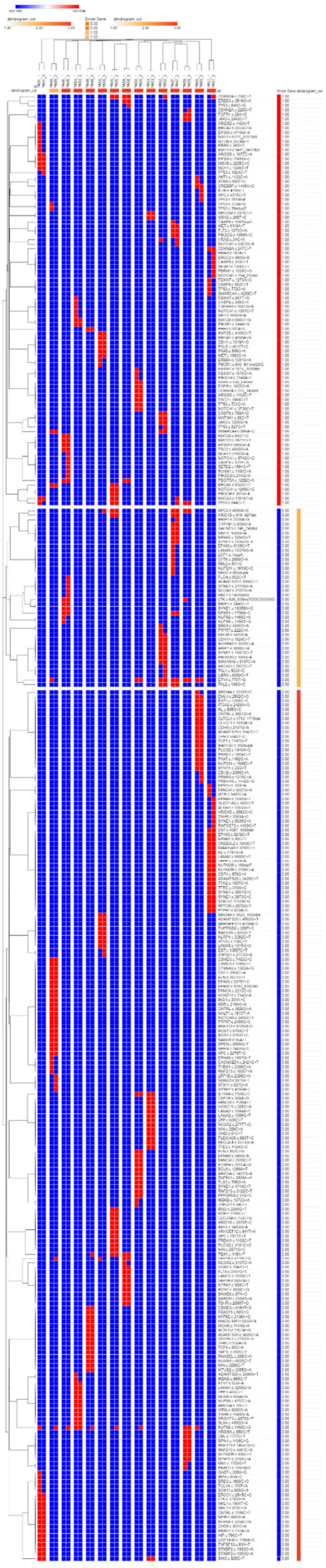


Figure S3. A total of 297 genes with verified mutations were classified into driver and nondriver categories.

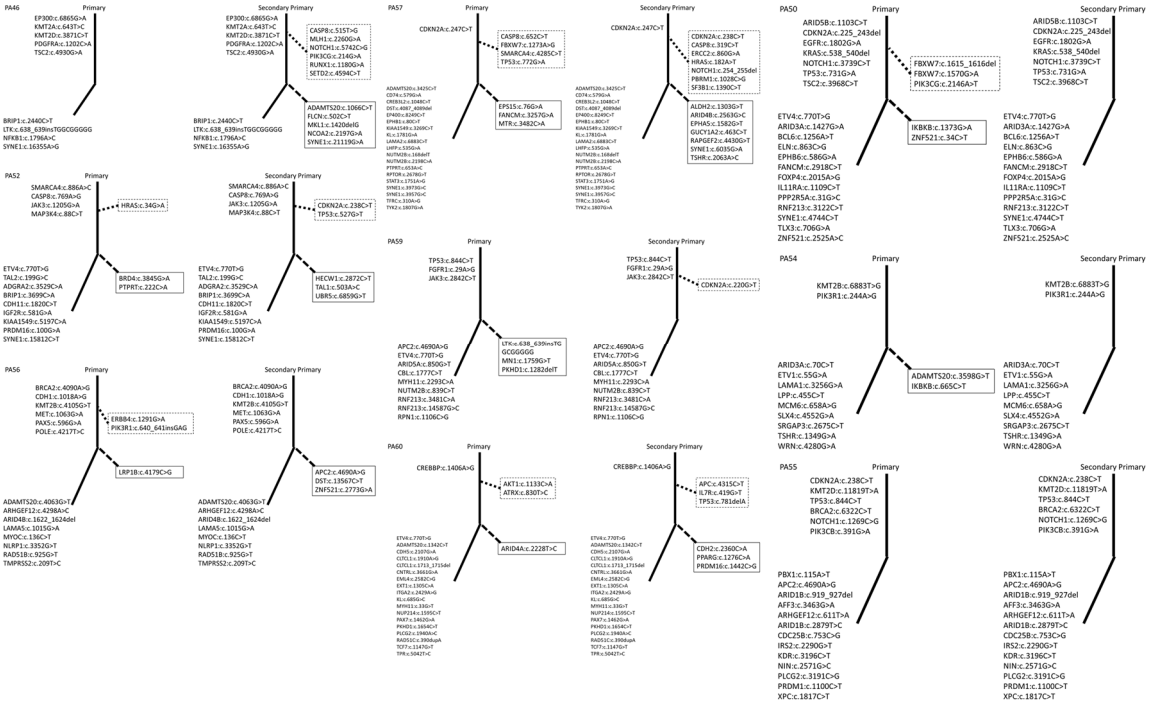


Figure S4. Analysis of driver gene and trunk mutations in first and second primary tumors.