

Comparative oncology DNA sequencing of canine T cell lymphoma via human hotspot panel

SUPPLEMENTARY MATERIALS

Supplementary Table 1: All mutations screened and identified. See [Supplementary_Table_1](#)

Supplementary Table 2: Mutations categorized based on canine breed. See [Supplementary_Table_2](#)

Supplementary Table 3: Targeted sequencing data for 346 called variants. See [Supplementary_Table_3](#)