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dbSNP: the NCBI database of genetic variation

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Whole exome sequencing analysis of canine urothelial carcinomas without BRAF V595E mutation:
Short in-frame deletions in BRAF and MAP2K1 suggest alternative mechanisms for MAPK pathway disruption. **2023**, 19, e1010575

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Evaluating The Infinium Human MethylationEPIC v2 BeadChip.

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