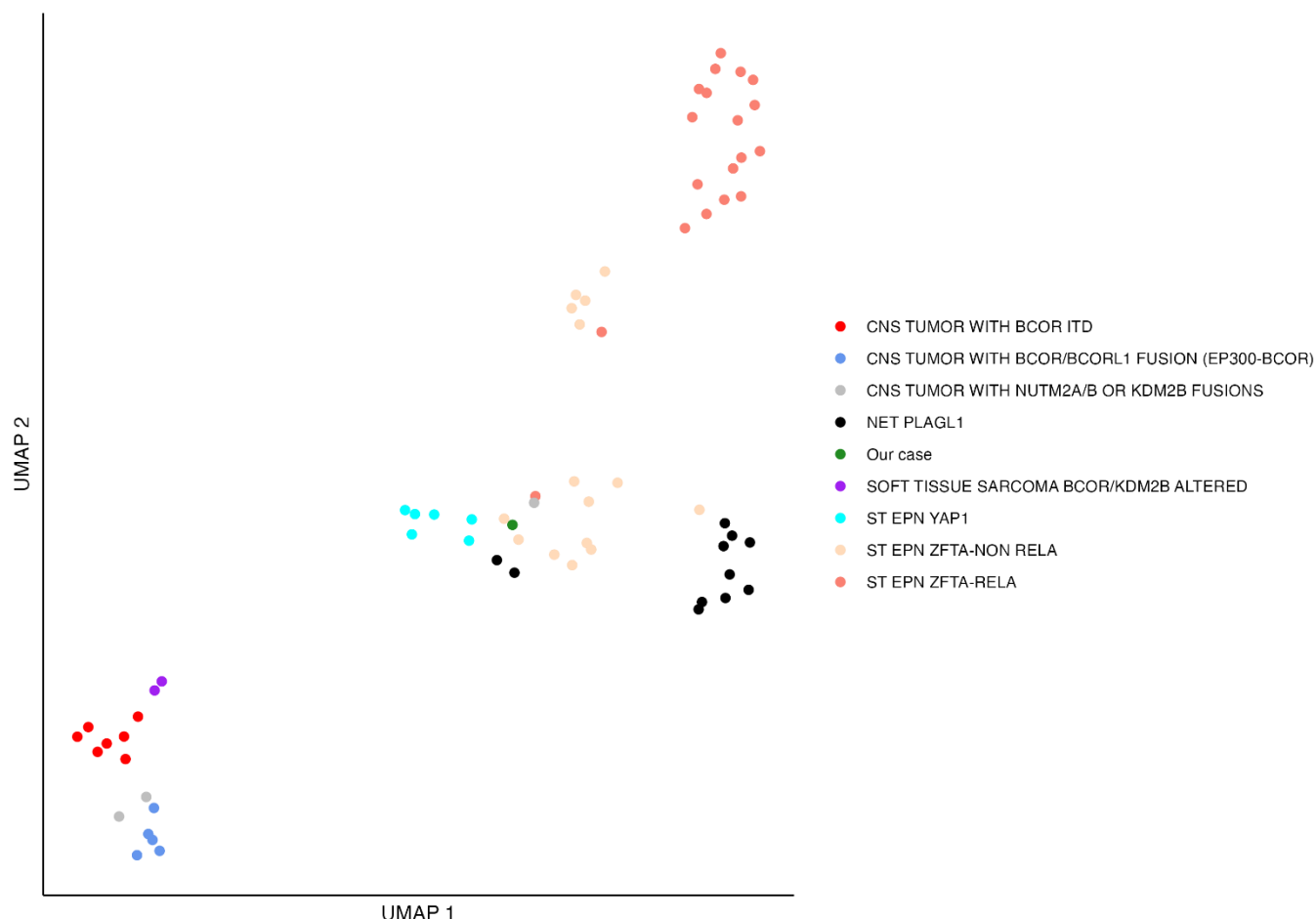


Supplementary Figure 01

Supplementary Figure 1. DNA methylation-based uniform manifold approximation and projection



The current case did not cluster with a methylation class when compared to previous published cohorts with well-known fusions belonging to the CNS TUMOR WITH BCOR ITD (n = 7) (PMID: [35174661](https://pubmed.ncbi.nlm.nih.gov/35174661/)), CNS TUMOR WITH BCOR/BCORL1 FUSION (EP300-BCOR) (n = 5) (PMID: [36782314](https://pubmed.ncbi.nlm.nih.gov/36782314/), PMID: [33138864](https://pubmed.ncbi.nlm.nih.gov/33138864/)), CNS TUMOR WITH NUTM2A/B OR KDM2B FUSIONS (n = 3) (PMID: [34717763](https://pubmed.ncbi.nlm.nih.gov/34717763/), PMID: [35174661](https://pubmed.ncbi.nlm.nih.gov/35174661/)), SOFT TISSUE SARCOMA BCOR/KDM2B-ALTERED (n = 2), CNS NET PLAGL1 (n = 11) (PMID: [38581034](https://pubmed.ncbi.nlm.nih.gov/38581034/)), ST EPN YAP1 (n = 6) (PMID: [30246434](https://pubmed.ncbi.nlm.nih.gov/30246434/)), ST EPN ZFTA-NON RELA (n = 15) (PMID: [34389065](https://pubmed.ncbi.nlm.nih.gov/34389065/)) and ST EPN ZFTA-RELA (n = 19) (PMID: [30325077](https://pubmed.ncbi.nlm.nih.gov/30325077/)).