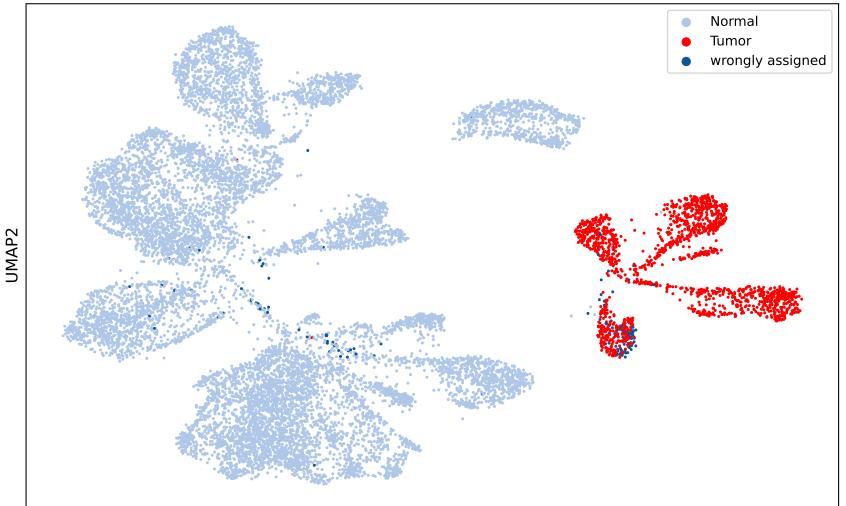kildisiute_celseq2: prediction

Legend:
- Normal
- Tumor
- wrongly assigned

Axis labels: UMAP1 (x-axis), UMAP2 (y-axis)