

6 Appendix

Table 3. Clinical characteristics of patients in the dataset.

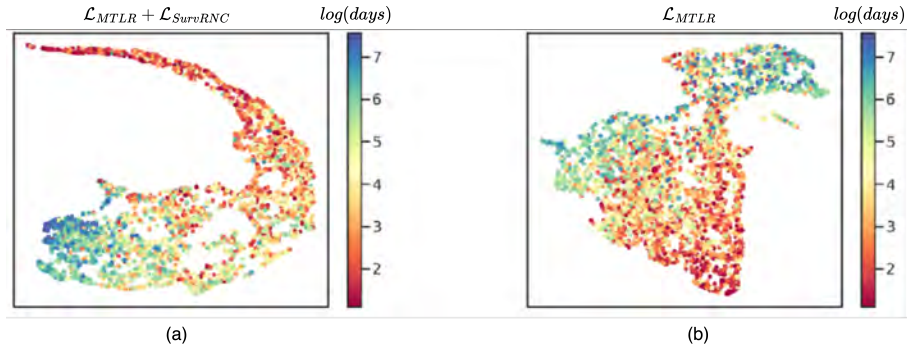
| Characteristics | Dataset (n = 488) |
|----------------------------|--------------------------|
| RFS | |
| Uncensored | 96 (19.7) |
| Censored | 392 (80.3) |
| Age (year) | |
| Weight (kg) | |
| Gender | |
| Male | 402 (82.4) |
| Female | 86 (17.6) |
| Alcohol consumption | |
| Yes | 95 (19.5) |
| No | 59 (12.1) |
| Unknown | 334 (68.4) |
| Tobacco consumption | |
| Yes | 85 (17.4) |
| No | 105 (21.5) |
| Unknown | 298 (61.1) |
| HPV status | |
| Positive | 274 (56.1) |
| Negative | 43 (8.8) |
| Unknown | 171 (35.1) |
| Performance status | |
| 0 | 86 (17.7) |
| 1 | 114 (23.3) |
| 2 | 11 (2.3) |
| 3 | 3 (0.6) |
| 4 | 1 (0.2) |
| Unknown | 273 (55.9) |
| Surgery | |
| Yes | 50 (10.3) |
| No | 248 (50.8) |
| Unknown | 190 (38.9) |
| Chemotherapy | |
| Yes | 422 (86.5) |
| No | 66 (13.5) |

Table 4. Preprocessing and augmentation details.

| Augmentations | Axis | Probability | Size |
|-------------------------|---------|-------------|--------------|
| Orientation | PLS | - | - |
| CT/PET Concatenation | 1 | - | - |
| Normalization | - | - | - |
| Random crop | - | 0.5 | 96 x 96 x 96 |
| Random flip | x, y, z | 0.1 | - |
| Rotate by 90 (up to 3x) | x, y | 0.1 | - |

Table 5. Ablation to study the effect of the parameter λ values.

| Models | λ | | | |
|--|-----------|--------|--------|--------|
| | 0.3 | 0.5 | 0.7 | 1.0 |
| DeepMTLR [8] $\mathcal{L}_{MTLR} + \mathcal{L}_{SurvRNC}$ | 0.6870 | 0.7009 | 0.6690 | 0.6742 |
| DeepHit [14] $\mathcal{L}_{Hit} + \mathcal{L}_{SurvRNC}$ | 0.6802 | 0.7233 | 0.6950 | 0.6808 |

**Fig. 3. UMAP Visualization:** $\mathcal{L}_{SurvRNC}$ with the native loss function of a DeepMTLR survival model on the SUPPORT dataset [13] shows a better continuous latent feature representation (a) as compared to using only the native loss function (b).