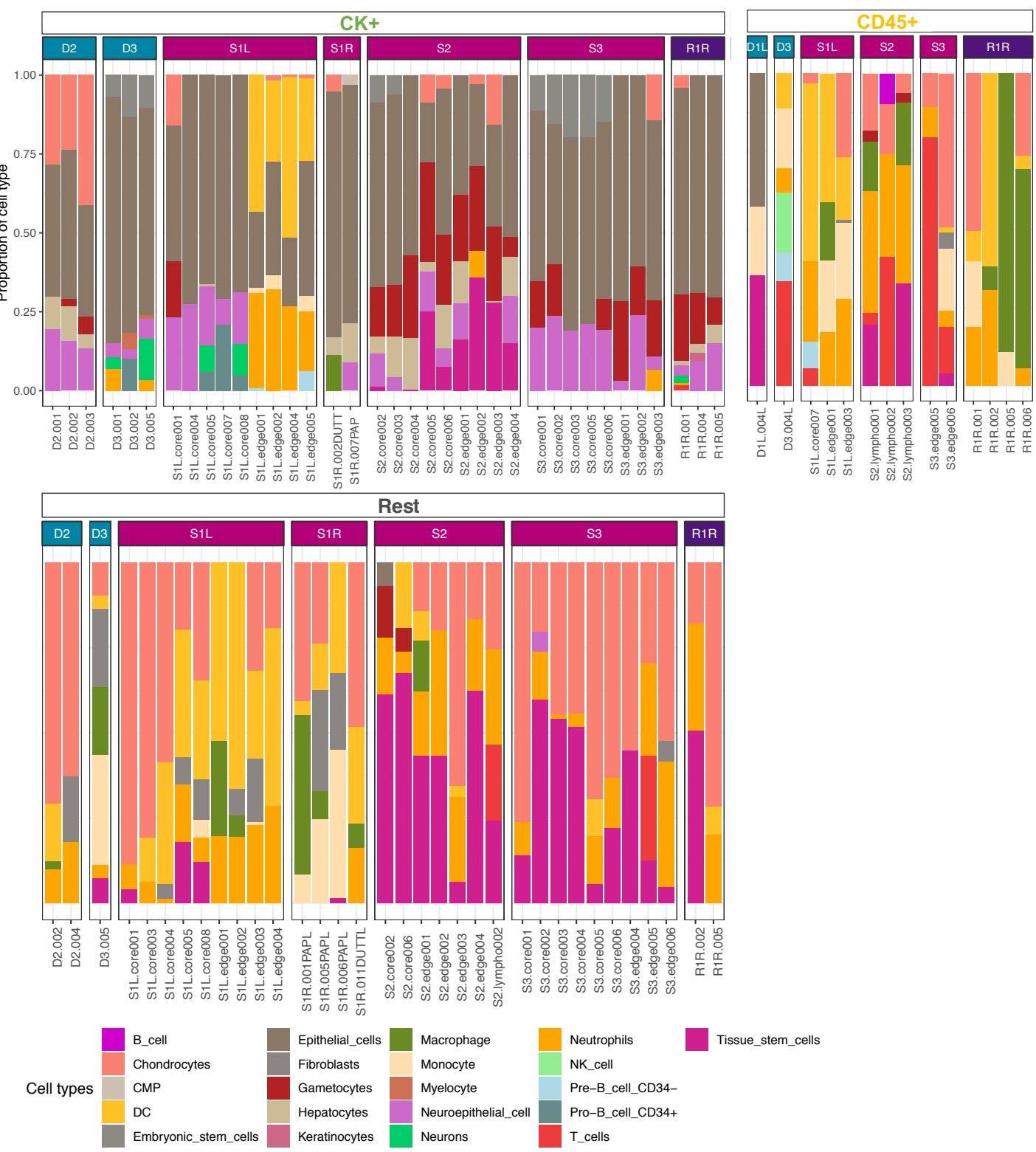


# Supplementary Figure S6. Spatially resolved changes in the transcriptional profile of tumour-associated immune cells and stroma



**Supplementary Figure S6. Spatially resolved changes in the transcriptional profile of tumour-associated immune cells and stroma.** Deconvolution of cell types estimated from gene expression profiles in each region of interest according to relevant staining (CK+: tumour cells, CD45+: immune cells, rest: stroma) using signatures from human protein cell atlas (HPCA) as reference. Patient1 (D1L: diagnostic biopsy left, S1L: surgery biopsy left, S1R: surgery biopsy right, R1R: relapse biopsy right) Patient2 (D2: diagnostic biopsy, S2: surgery biopsy), Patient3 (D3: diagnostic biopsy, S3: surgery biopsy).