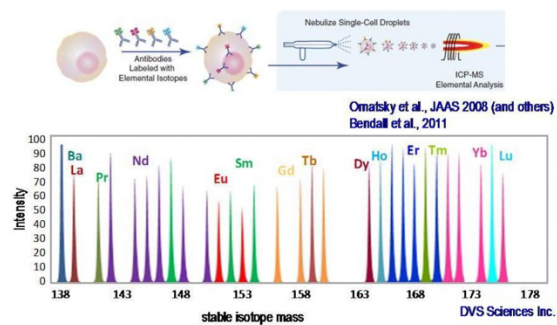
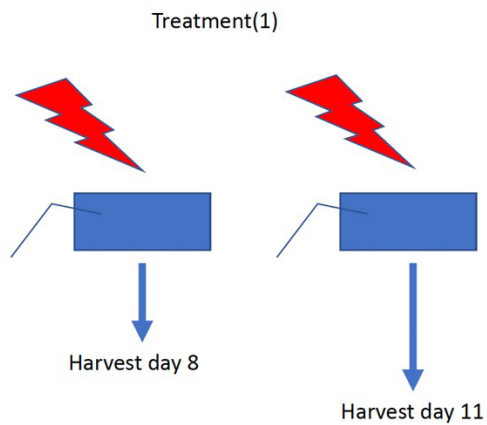


Data info:

- (1) Samples: 3 treatment groups, compared between 2 days.
- (2) Data was acquired on a mass spec cytometer "cyTOF"
- (3) Data exports were raw expression levels of 62 cellular antigen markers from splenocytes.



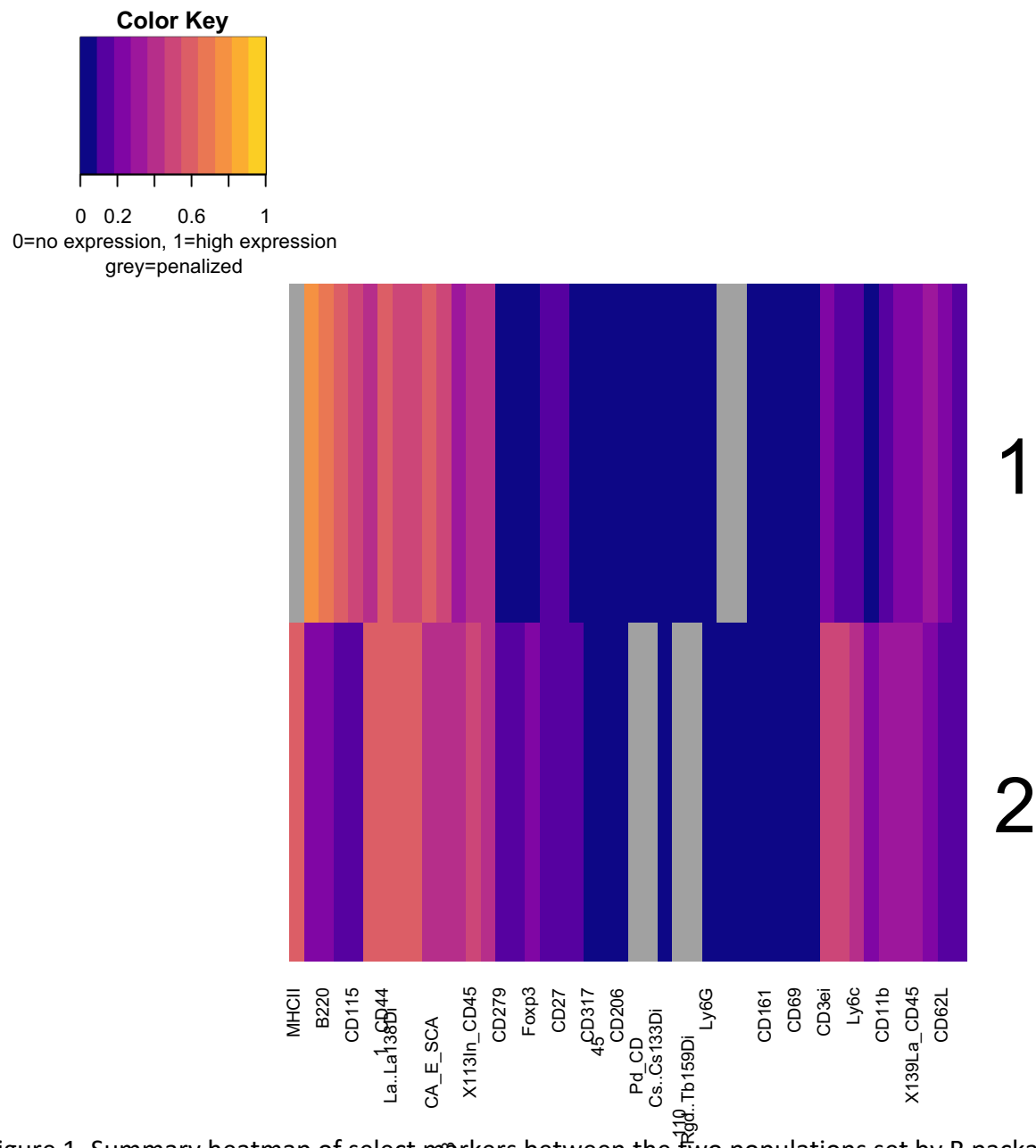


Figure 1. Summary heatmap of select markers between the two populations set by R package: Depeche.

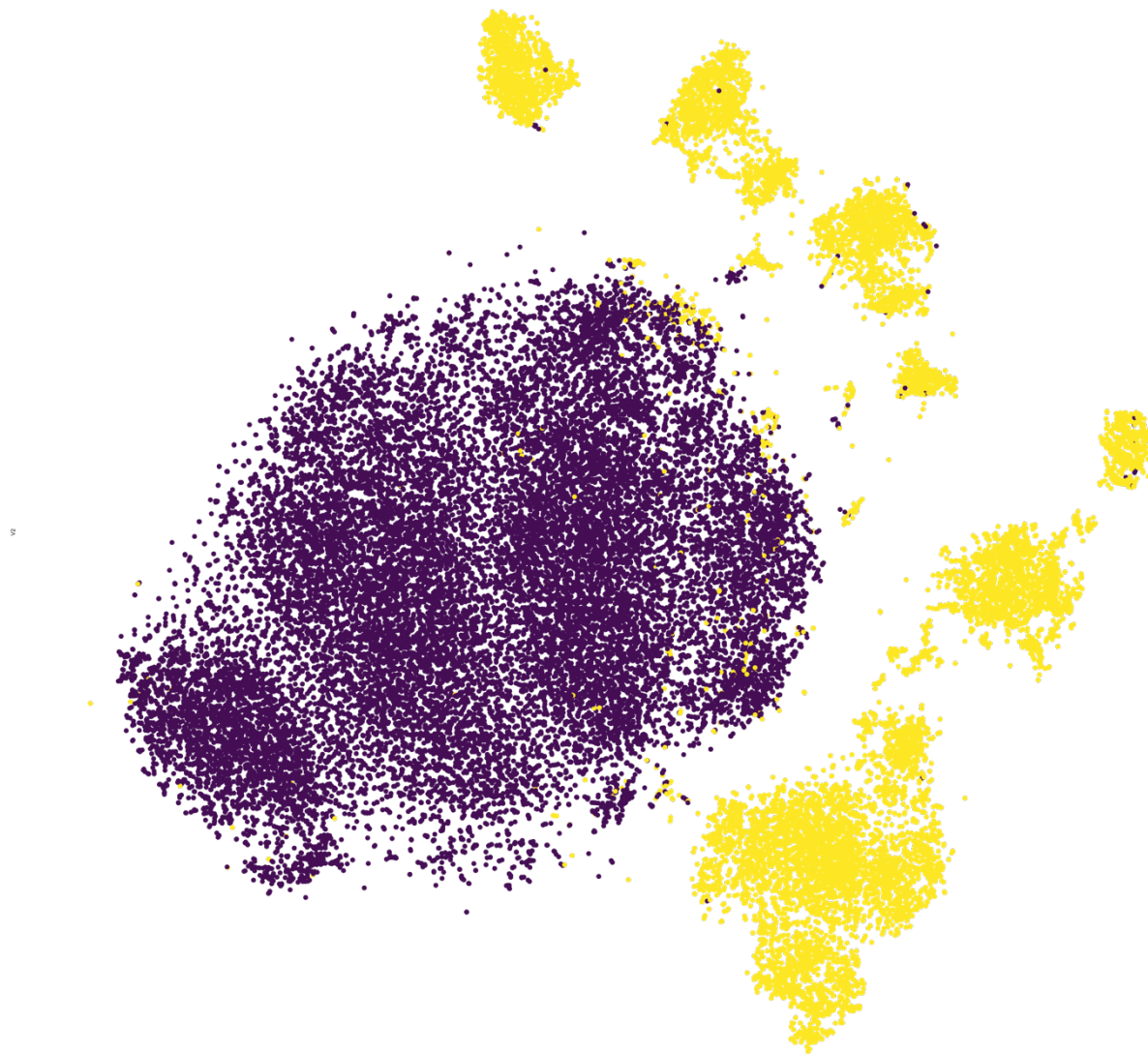


Figure 2. tSNE plot of all cells. R package: rTSNE plot of two populations set by R package: Depeche. 2000 iterations & 30 perplexity.

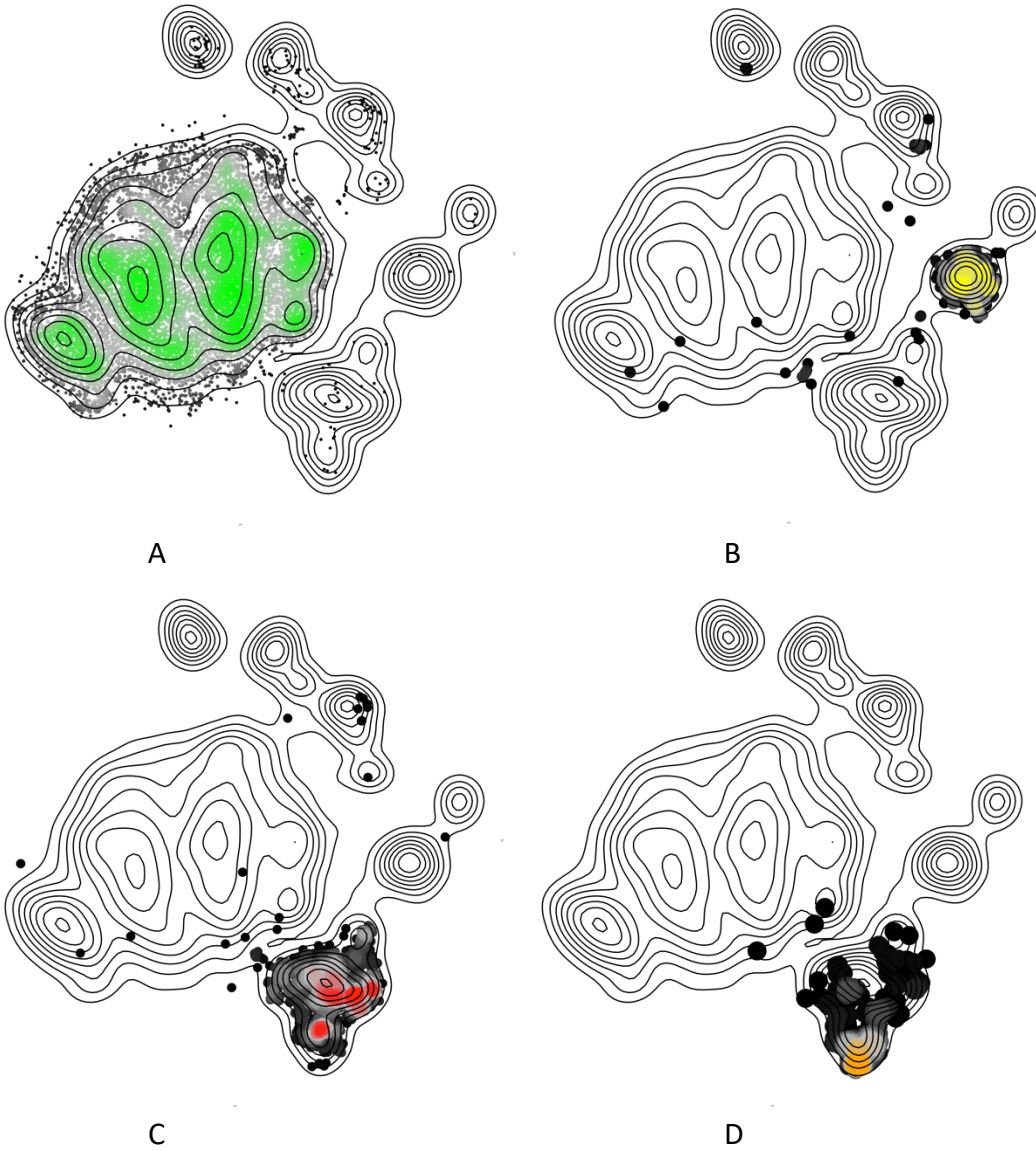


Figure 3. tSNE plots highlighting: B-cells (A), cytotoxic T-lymphocytes (B), T-helper cells (C), and T-regulatory cells (D).

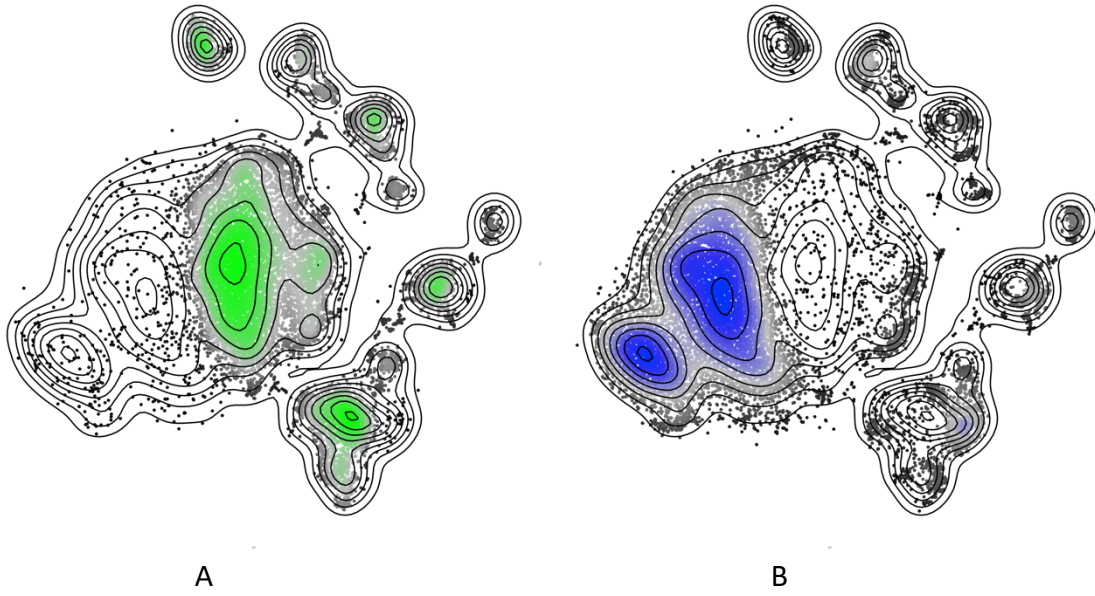


Figure 4. tSNE plots highlighting: Day 8 (A), and Day 11 (B).