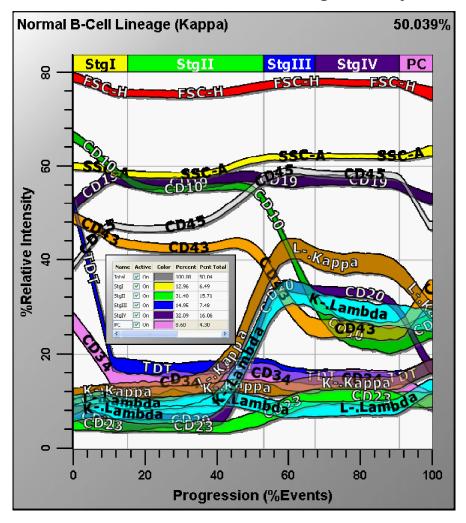
12-Parameter B-Cell Lineage Analysis



Data kindly provided by Dr. Frederic Preffer, Boston, MA

The above graph summarizes 12 parameter correlations associated with the lineage of B-cells in a normal marrow specimen (CD19, SSC, TdT, CD10, CD20, CD45, CD43, CD34, CD23, FSC, Kappa, and Lambda). The x-axis delineates the lineage progression where the percentages of all the sub-categories are directly read from the axis. The y-axis represents the relative intensity of all the parameters. The width of the bands is set to 0.2 SD's and therefore corresponds to the variability of the data as the cells differentiate. The tightly coordinated parameter transitions associated with the stages of B-cell maturation are clearly shown. The percentages of each of the B-cell states are summarized in the table. These percentages account for population overlap due to measurement error. It should be noted that this one graph represents all the correlations that would be present in 66 two-parameter histograms.

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- Novel analysis approach defines populations without gates.
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