



Figure 11.31. Schematic of output from a chromatography column. The relative t_{\max} values are used to calculate resolution, while the standard deviation of a peak ($\sigma_{t_{\max,i}}$) is used to estimate yields, purity, and capacity and is used in scale-up.

The yield is defined as the amount of a solute, such as i , collected between times t_1 and t_2 divided by the total amount of i injected into the column. Thus,

$$Y_{\Delta t,i} = \frac{\int_{t_1}^{t_2} y_i F dt}{\int_0^{\infty} y_i F dt} \quad (11.86)$$

where y_i is the concentration of solute, i , and F is the solvent flow rate.

The value of y_i can be estimated from the peak height and shape if the peak is assumed to be Gaussian. This assumption leads to

$$y_i = y_{\max,i} \exp \left[-\frac{(t - t_{\max,i})^2}{2(\sigma_{t_{\max,i}})^2} \right] \quad (11.87)$$

where $t_{\max,i}$ is the time at which the maximum amount of i elutes, t is elution time, and $\sigma_{t_{\max,i}}$ is the standard deviation for peak i . Then the yield can be calculated as

$$Y_{\Delta t,i} = \frac{1}{2} \left\{ \operatorname{erf} \left[\frac{(t_2 - t_{\max,i})}{\sqrt{2} \sigma_{t_{\max,i}}} \right] - \operatorname{erf} \left[\frac{(t_1 - t_{\max,i})}{\sqrt{2} \sigma_{t_{\max,i}}} \right] \right\} \quad (11.88)$$

where erf is the error function as defined by