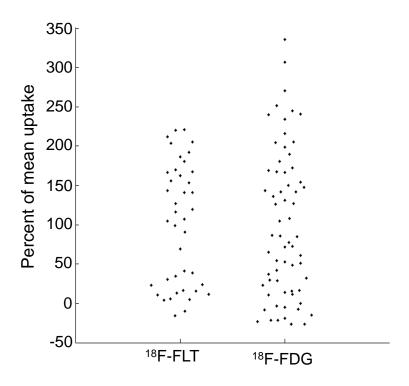


**Supplemental Figure 1:** (A) Automated clustering applied to single-cell <sup>18</sup>F-FLT measurements (20% serum), using a gaussian mixture model with 2 clusters. The symbols  $\times$  and  $\Diamond$  represent single cells in the <sup>18</sup>F-FLT-low and <sup>18</sup>F-FLT-high clusters, as determined by clustering. (B) Cumulative distribution function (CDF) for the experimentally measured <sup>18</sup>F-FLT uptake (*blue*), and maximum-likelihood curve fitting using 1-gaussian model (*red*) and 2-gaussian mixture model (*orange*).



**Supplemental Figure 2:** Typical single-cell distribution of <sup>18</sup>F-FLT and <sup>18</sup>F-FDG, normalized to average population uptake.