

Supplementary Materials

Figure S1. Linearity of each standard lipid used to spike the cell suspension. A suspension containing 1×10^6 WR/SMS1 cells was spiked with lipid standards and analyzed by the present method, without the use of any internal standards during MS. Data are presented as the \log_e - \log_e plot of the MS peak area for each eluted lipid *versus* the amount of standard added.

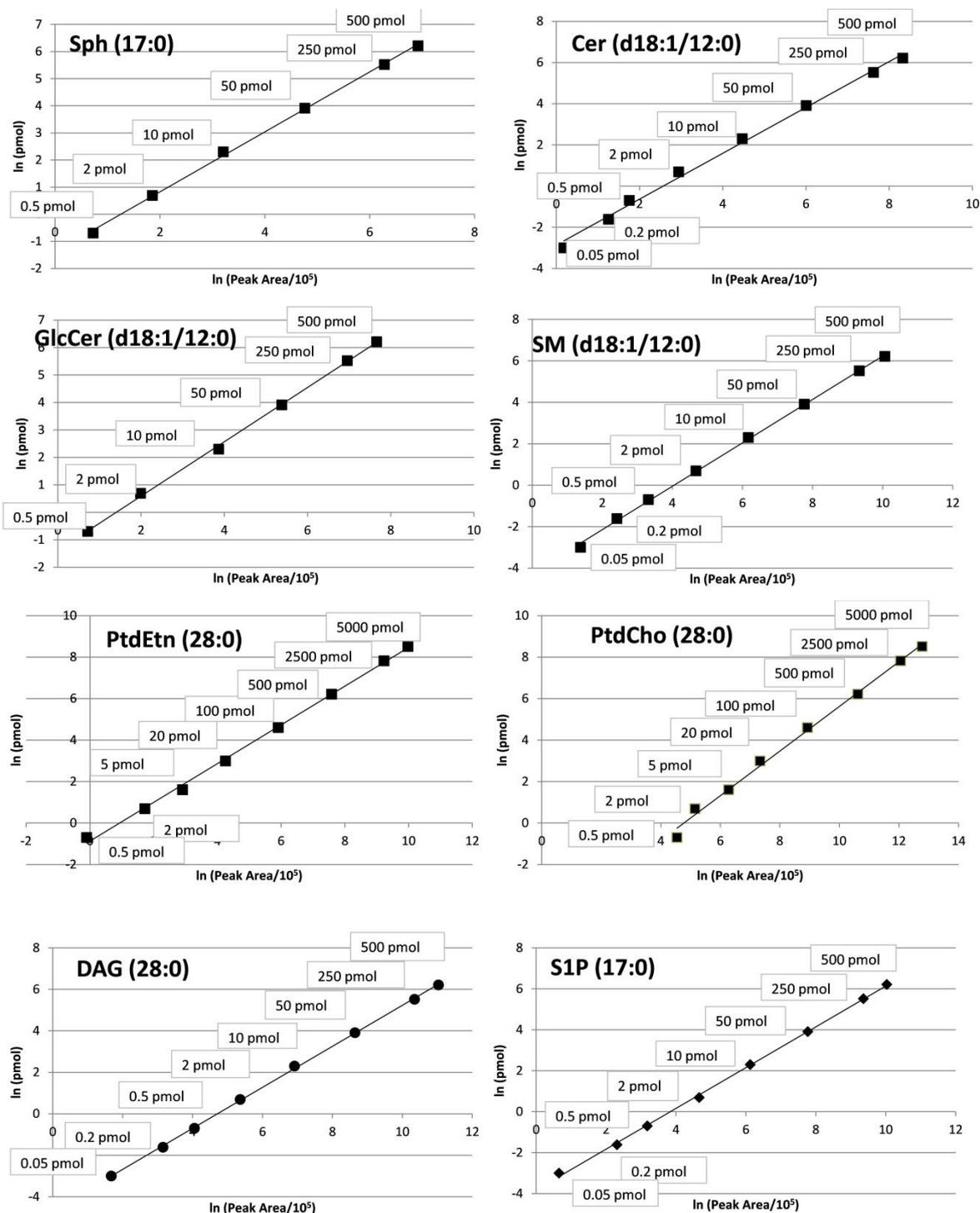


Figure S1. Cont.

