

Supplementary Figure 1. Effect of ST1926 on total levels of 17dhCer, and ratio of total 17Cer to 17dhCer species in HTLV-1 positive and negative malignant $T$ cells.
(A) HTLV-1 positive (HuT-102) cells has lower basal levels of 17C-dihydroceramide (17dhCer species) than Molt-4 cells, and treatment with ST1926 increased 17dhCer levels in both HuT-102 and Molt-4 cells, while (B) there was no significant change in the ratio of 17Cer/17dhCer species upon ST1926 treatment in both cell lines, or in the basal level ratios of 17Cer/17dhCer species between the two cell lines. Cells were seeded at a density of $3 \times 105$ cells $/ \mathrm{ml}$, labeled with $4 \mu \mathrm{M}$ of unnatural 17C-sphinganine and treated with either $0.1 \%$ DMSO as control or $1 \mu \mathrm{M}$ ST1926 for 24 h . 17 dhCer species levels ( pmol ) were measured in lyophilized samples as duplicates by LC/MS as described in Methods and normalized to total cellular lipid phosphate levels ( $\mu \mathrm{mol}$ ). Data points represent the mean $\pm$ range ( $\mathrm{n}=2$ ). Results are representative of two independent experiments.

## Supplementary Table 1

Percent increase in individual 17dhCer species in HuT-102 and Molt-4 cells

|  | Total 17dhCer | 17dhC16 <br> $(\%$ Total 17dhCer $)$ | 17dhC22 <br> $(\%$ Total 17dhCer) | 17dhC24:1 <br> (\% Total dhCer) |
| :---: | :---: | :---: | :---: | :---: |
| HuT-102 | 327 | 48 | 7 | 12 |
| Molt-4 | 483 | 71 | 2 | 6 |

Supplementary Table 1. Percent accumulation of 17 dhC 16 , 17 dhC 22 , and $17 \mathrm{dhC} 24: 1$ in HTLV-1 positive (HuT-102) and negative (Molt-4) malignant $T$ cells, representing most prominent increase in their respective categories of medium long chain (MLC), long chain (LC), and very long chain (VLC) 17dhCer.
Cells were seeded at a density of $3 \times 10^{5}$ cells $/ \mathrm{ml}$ and treated with $0.1 \%$ DMSO as control or $1 \mu \mathrm{M}$ ST1926 for 24 h . 17dhCer species levels ( pmol ) were measured in lyophilized samples as duplicates by LC/MS as described in Methods and normalized to total cellular lipid phosphate levels ( $\mu \mathrm{mol}$ ). Data points represent the mean $\pm$ range ( $\mathrm{n}=2$ ). Results are representative of two independent experiments.

