



Supplementary Figure S-2: Electron ionising mass spectrometry of duramycin-porphyrin conjugate.

The peak for the conjugate was found to be ESI-MS (+): 675.66 $[(M+4H^+)/4]^{4+}$ where the expected was 675.67. Conjugate retention time; $R_t = 7.34$ minutes. The peaks at <300.00 were unidentified but were presumed to be a discrepancy within the machine most likely due to an impurity within the mass spectrometer.