

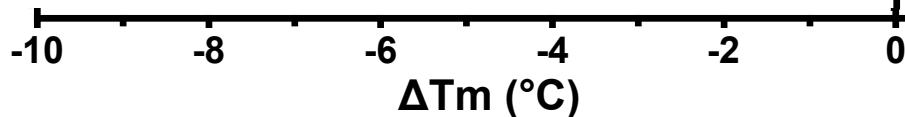
D14 (13 μ M)

GR24

Br-PMF

I-PMF

Br-BMF



DAD2 (33 μ M)

GR24

Br-PMF

I-PMF

Br-BMF



Supplementary information, Figure S11. Thermal destabilization of D14 and DAD2 by SL analogs.
 Colored bars represent the change in T_m (midpoint temperature between native and unfolded protein) relative to apo receptor, as determined by thermostability shift assay (TSA). Top: D14, Bottom: DAD2. The Y-axes indicate the molar ratios of ligand to receptor (0:1 [no ligand=apo receptor], 1:1, 2.5:1, 5:1, 10:1, 20:1 [20-fold ligand excess over receptor])