Toward the Discovery of a Novel Class of YAP–TEAD Interaction Inhibitors by Virtual Screening Approach Targeting YAP–TEAD Protein–Protein Interface

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Supplementary Material

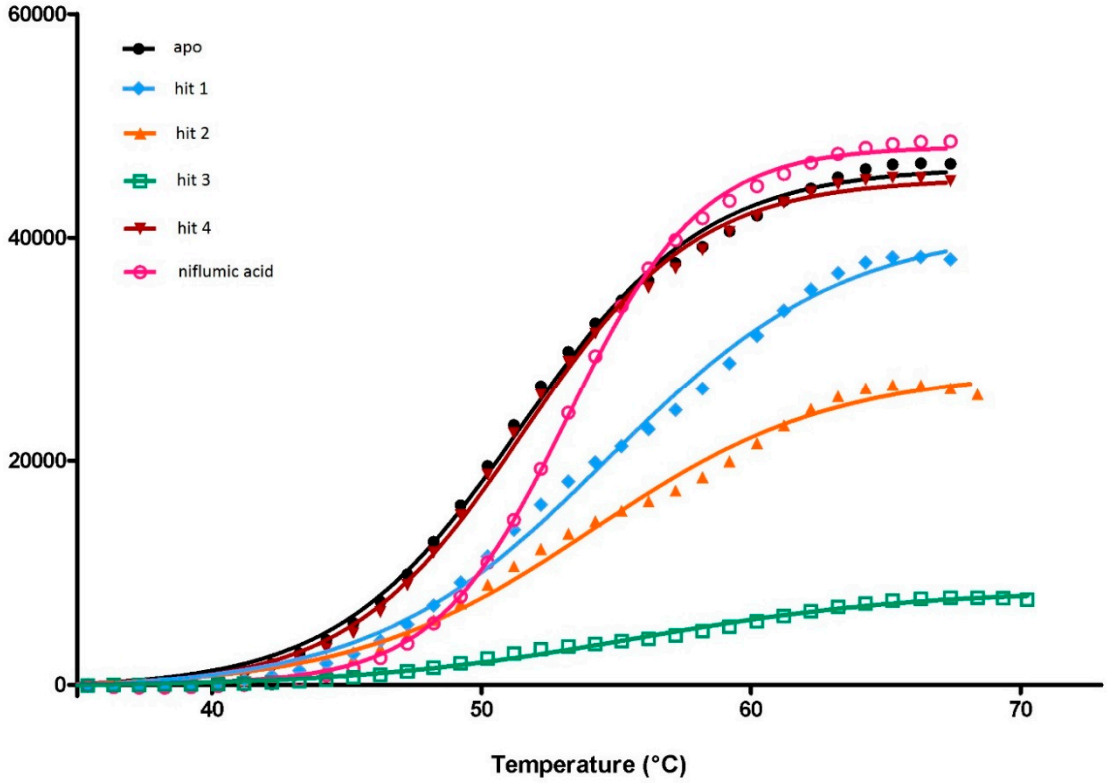

Figure S1. Representative normalized graphs showing a shift in thermal stability of hTEAD2$_{217–447}$ in presence of the potential TEAD2 binders **1-4** and niflumic acid. Apo refers to the melting curve of the protein alone.

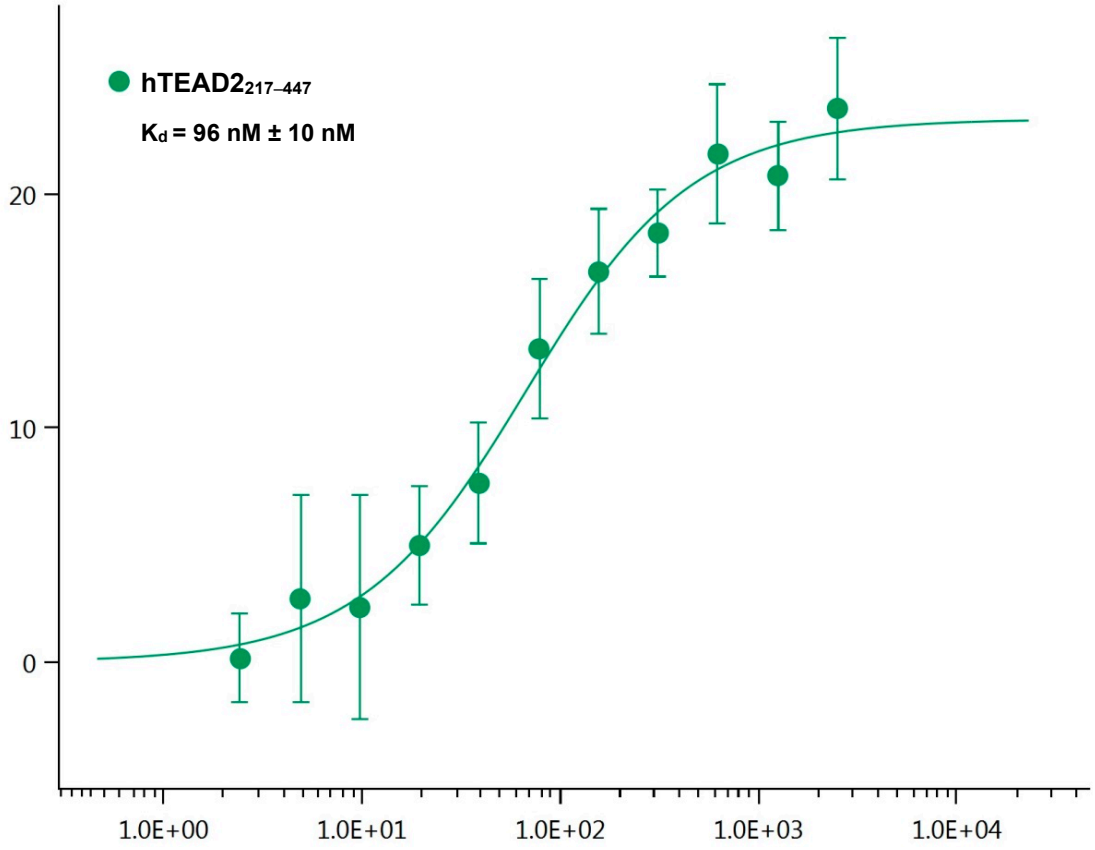

Figure S2. Dose-response curve for the binding interaction between hTEAD2$_{217–447}$ and YAP. The concentration of hTEAD2$_{217–447}$ is kept constant at 10 nM while the YAP$_{50–102}$ concentration varies from 10 μM to 4.8 nM.