

SUPPLEMENTAL FIGURE LEGEND

Figure S.1. Analytical ultracentrifugation sedimentation equilibrium of the p300_{CH1}/viral oncoprotein complexes. A. Analytical ultracentrifugation sedimentation equilibrium curves and residuals of the p300_{CH1}/Ad5-E1A₁₋₇₇ complex. The experiment was performed using three protein complex concentrations and three speeds, as indicated. All of the data was fit globally to a single species model, as well as to a 1:1 binding model: $A + B \rightleftharpoons AB$. B. Analytical ultracentrifugation sedimentation equilibrium curves and residuals of the p300_{CH1}/HPV16-E7₁₋₅₁ complex. The experiment was performed using three protein complex concentrations and three speeds, as indicated. All of the data was fit globally to a single species model, as well as to a 1:1 stoichiometry binding model: $A + B \rightleftharpoons AB$.

Figure S.1A

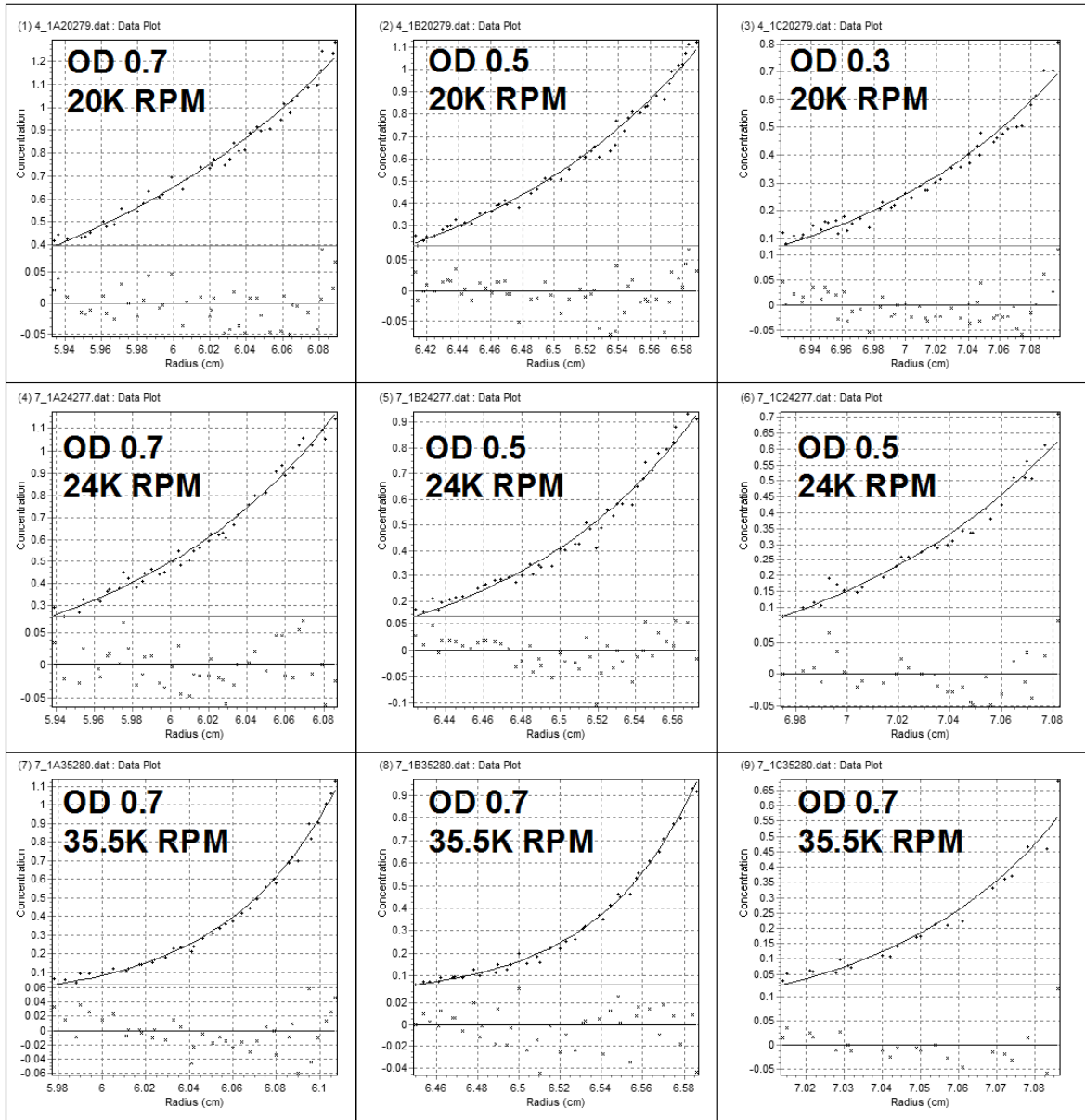


Figure S.1B

