



Figure S4. Another view of the ribbon model of hADAR2-D. The protein is rotated 90° about the vertical axis from Fig. 1B of the main text. IP₆ is shown as sticks. The active site zinc atom is obscured by β11. The color scheme is the same as Figure 1B of the main text. Ends of the disordered segment (residues 462-473) are indicated with asterisks (the beginning of the disordered region is obscured by β10).