Figure S2. Related to Figure 2. MRcd wild-type, L802W, and R805E proteins bind to ATP and form the open and closed states at low temperature. SAXS analyses of MRcd variants, as indicated by the key, were compared to wild-type complexes in the absence (left) and presence (right) of ATP. Scattering curves overlay across the whole scattering profile. Inset are models of the MR complexes adopted in the different states.