Supplementary Materials

Deciphering structural alterations associated with activity reductions of genetic polymorphisms in cytochrome P450 2A6 using molecular dynamics simulations

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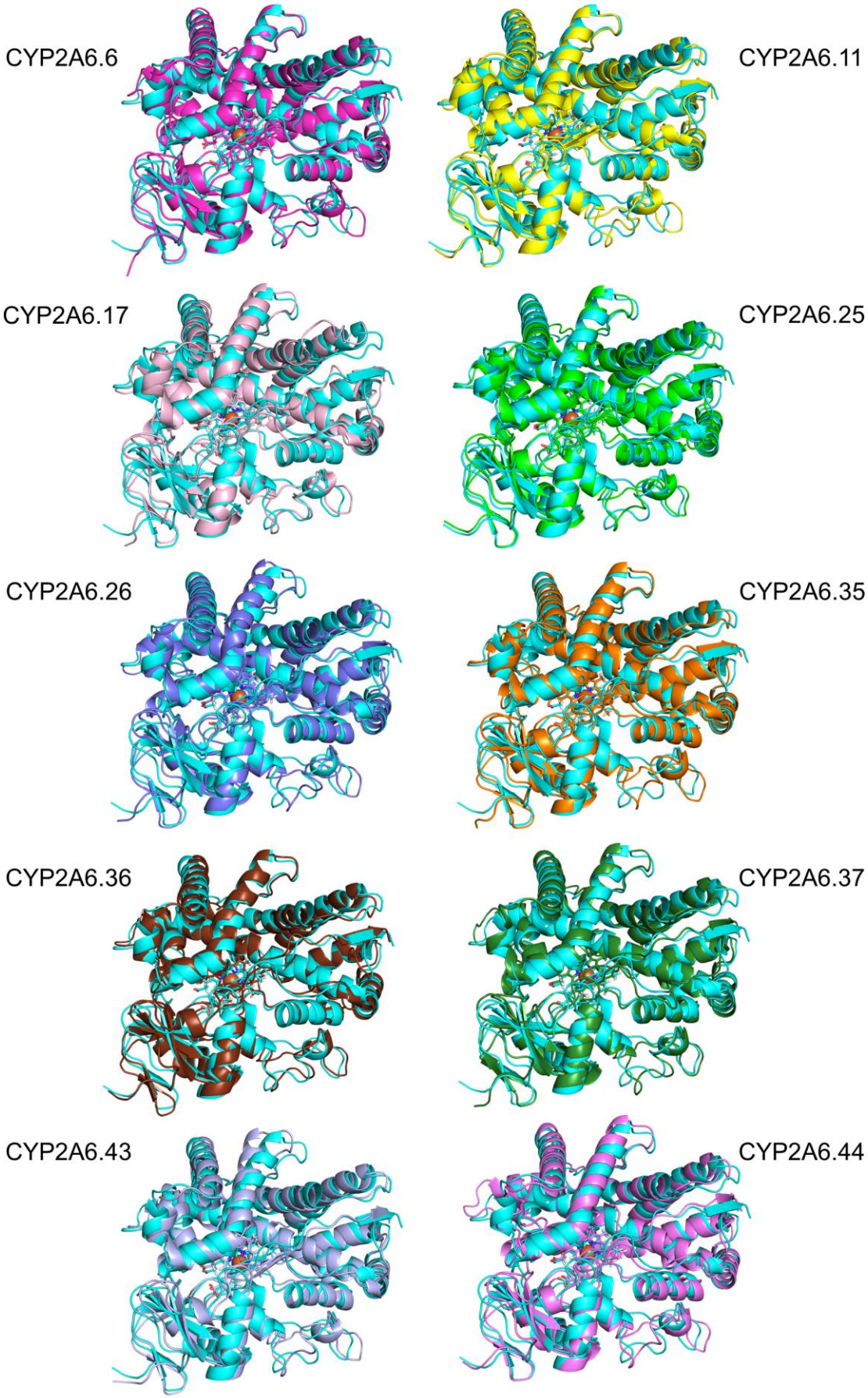


Figure S1. Final structures obtained by MD simulations. All mutant structures are superposed with the simulated structure of the wild type (cyan). Heme and iron are shown in stick and ball models, respectively.