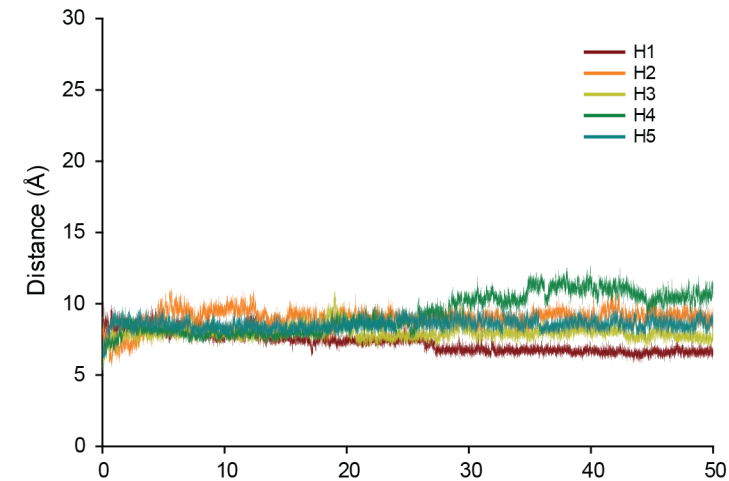
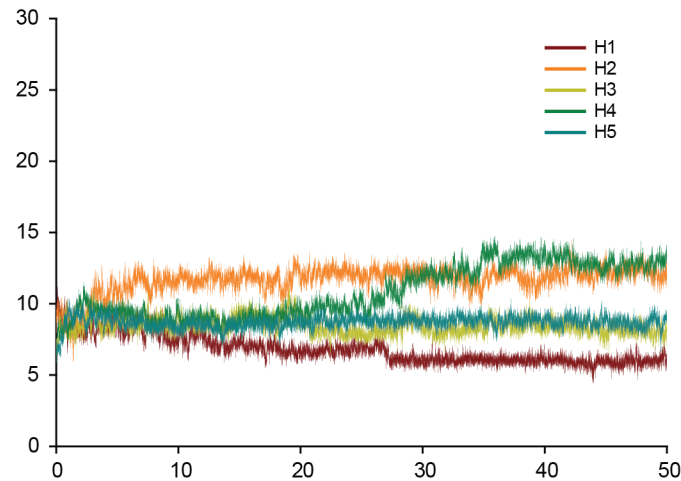
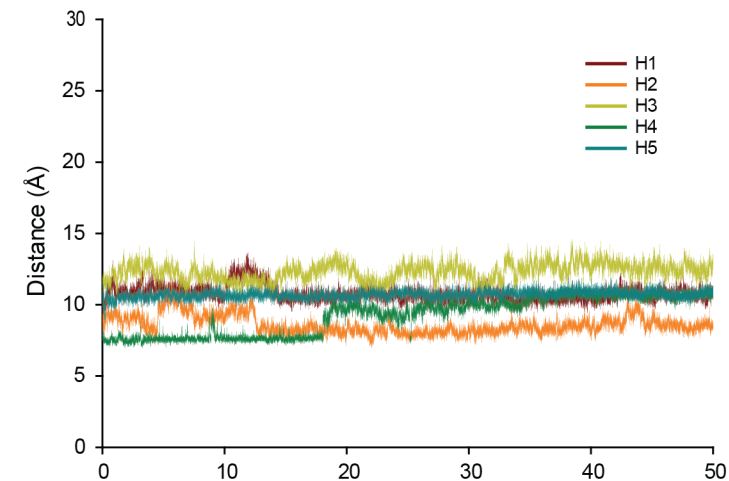
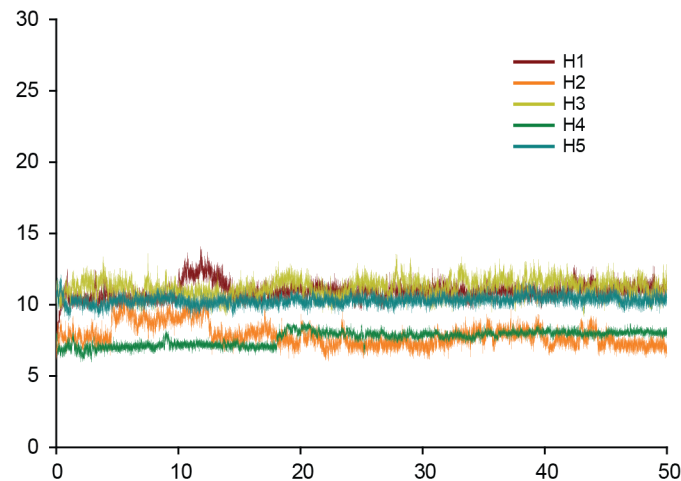
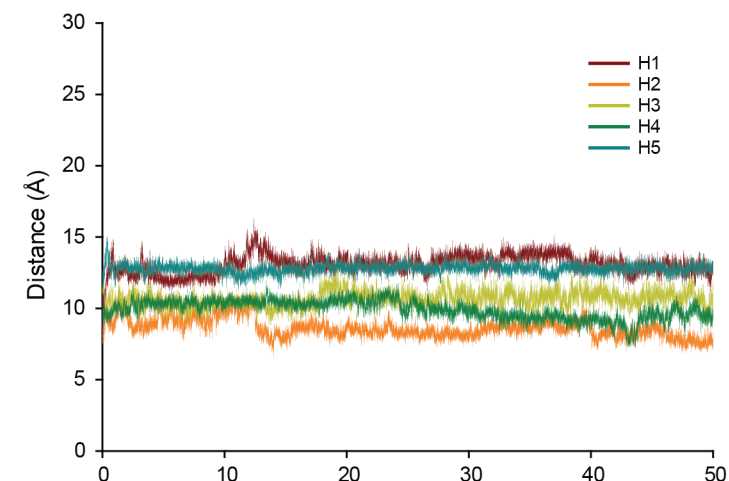


**A** Distance from residue K288**B** Distance from residue S289**C** Distance from residue D374**D** Distance from residue E375**E** Distance from residue Q404**F** Distance from residue R567