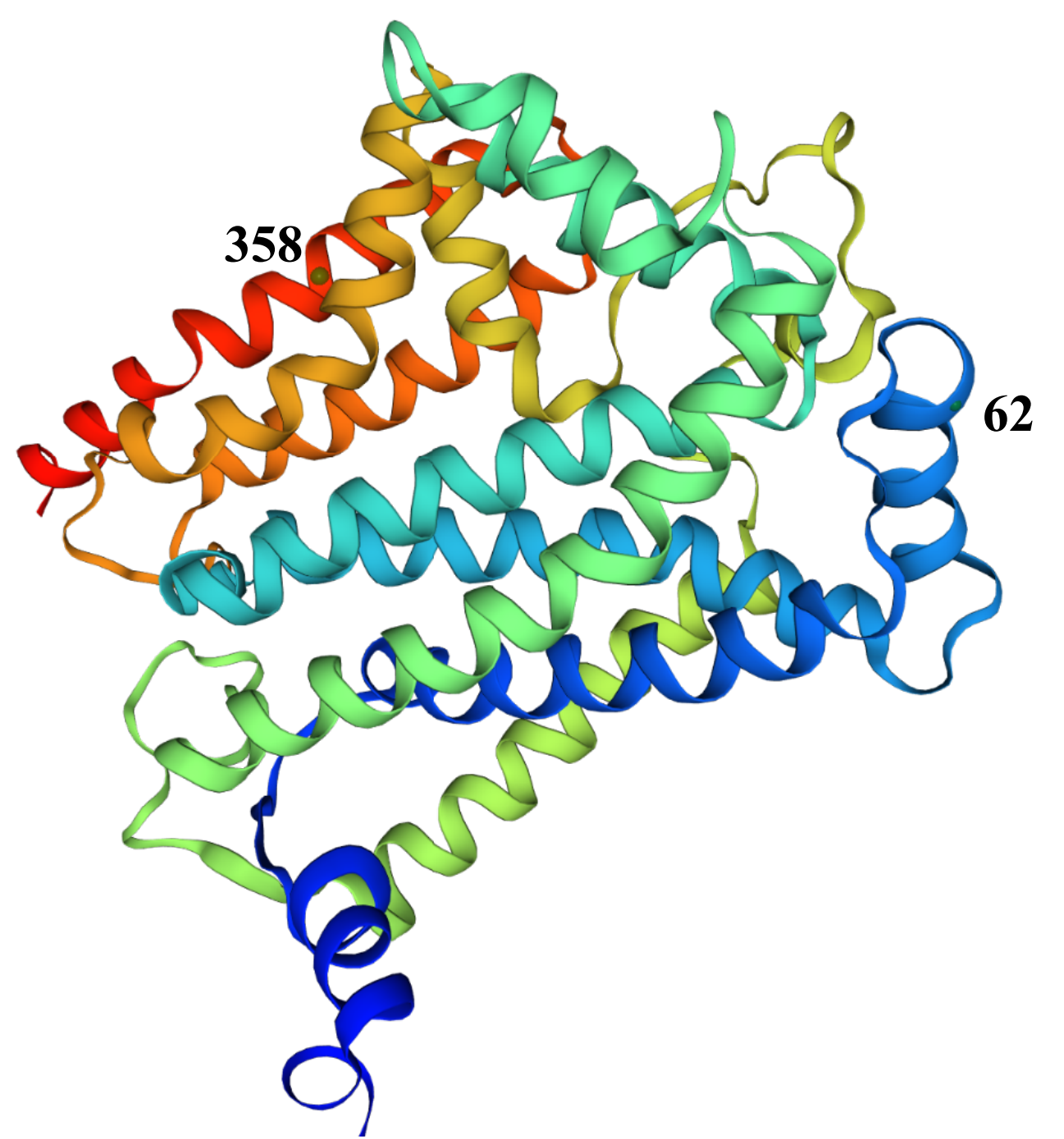
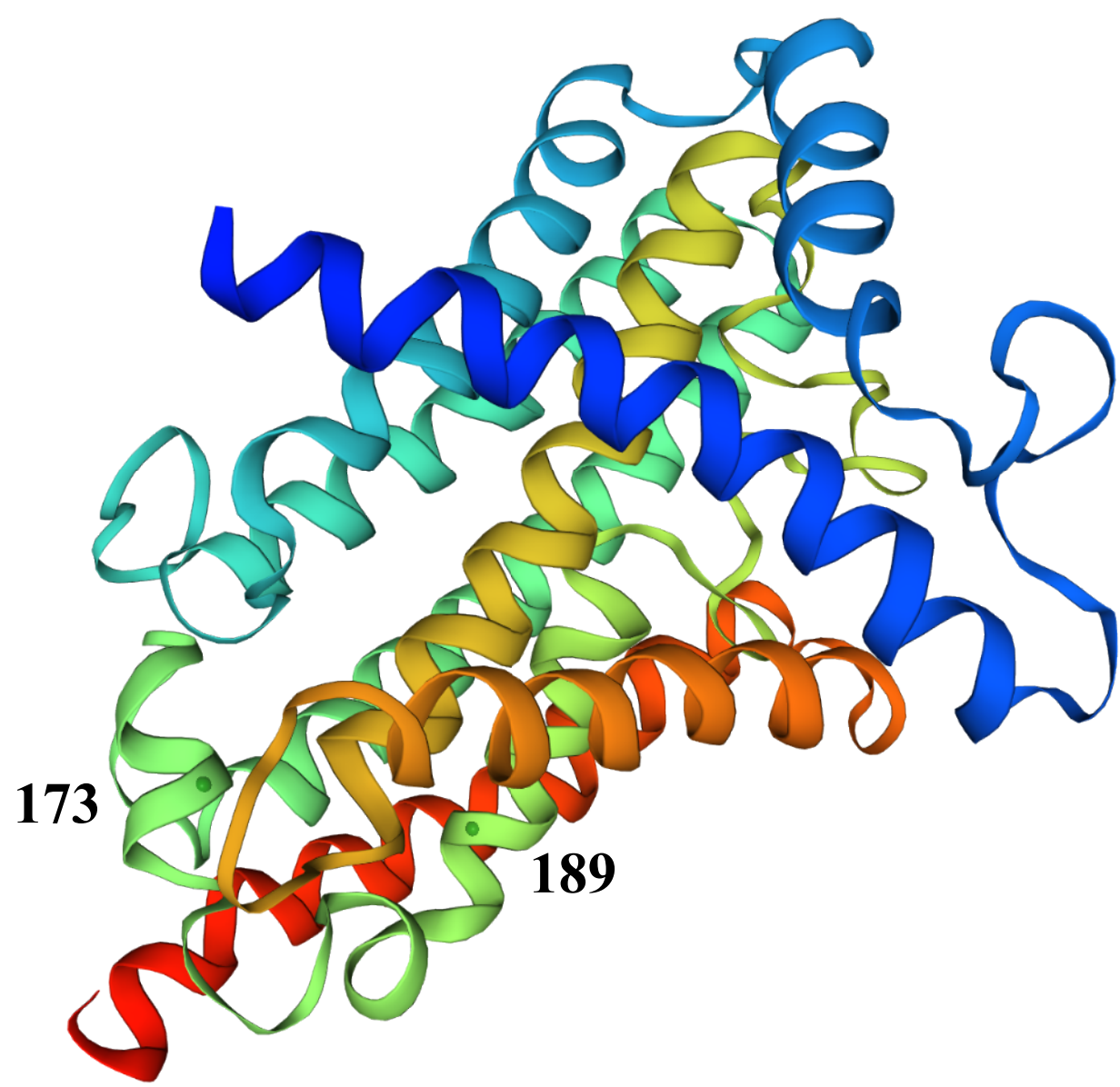


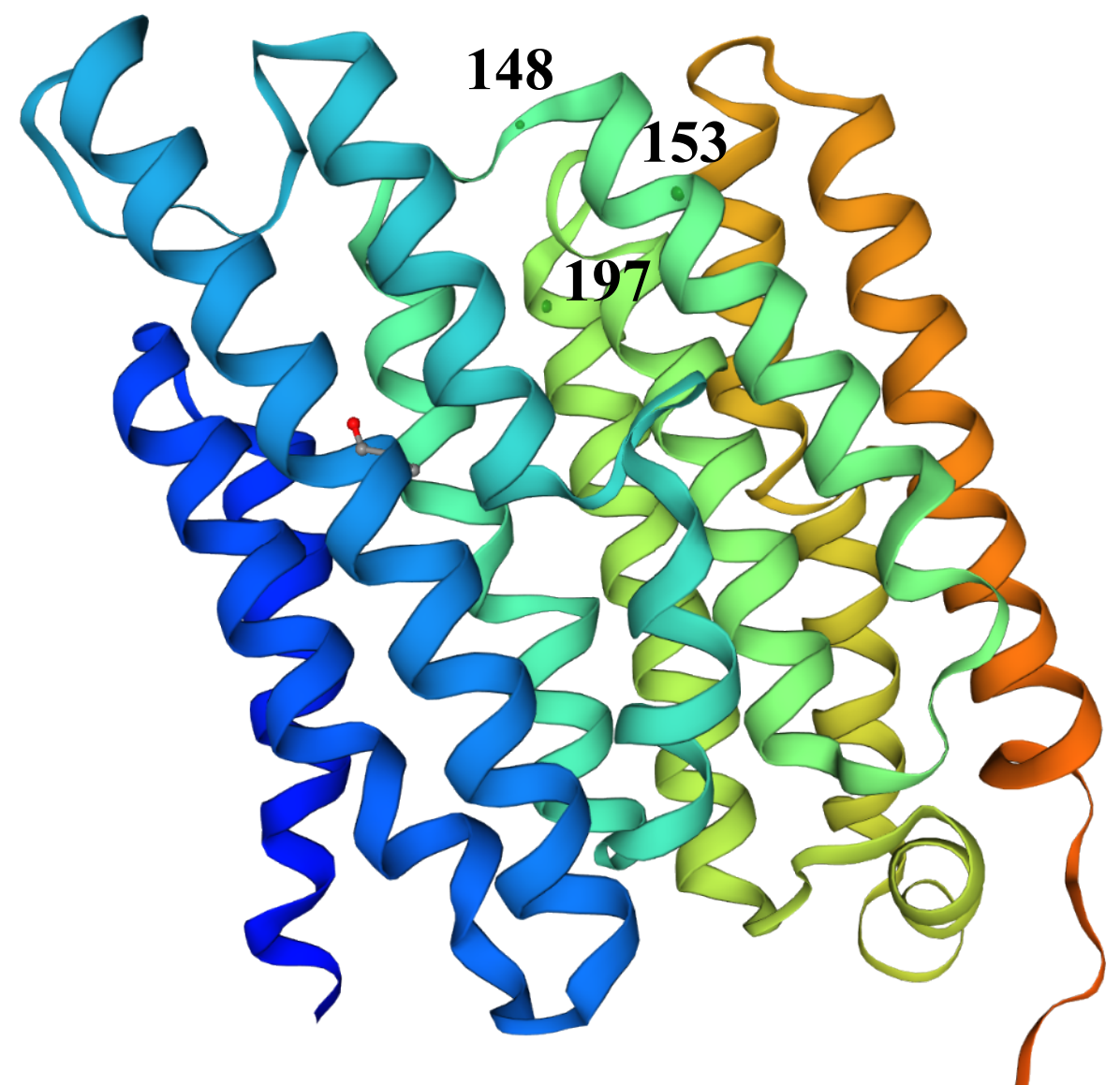
A. *COX1*



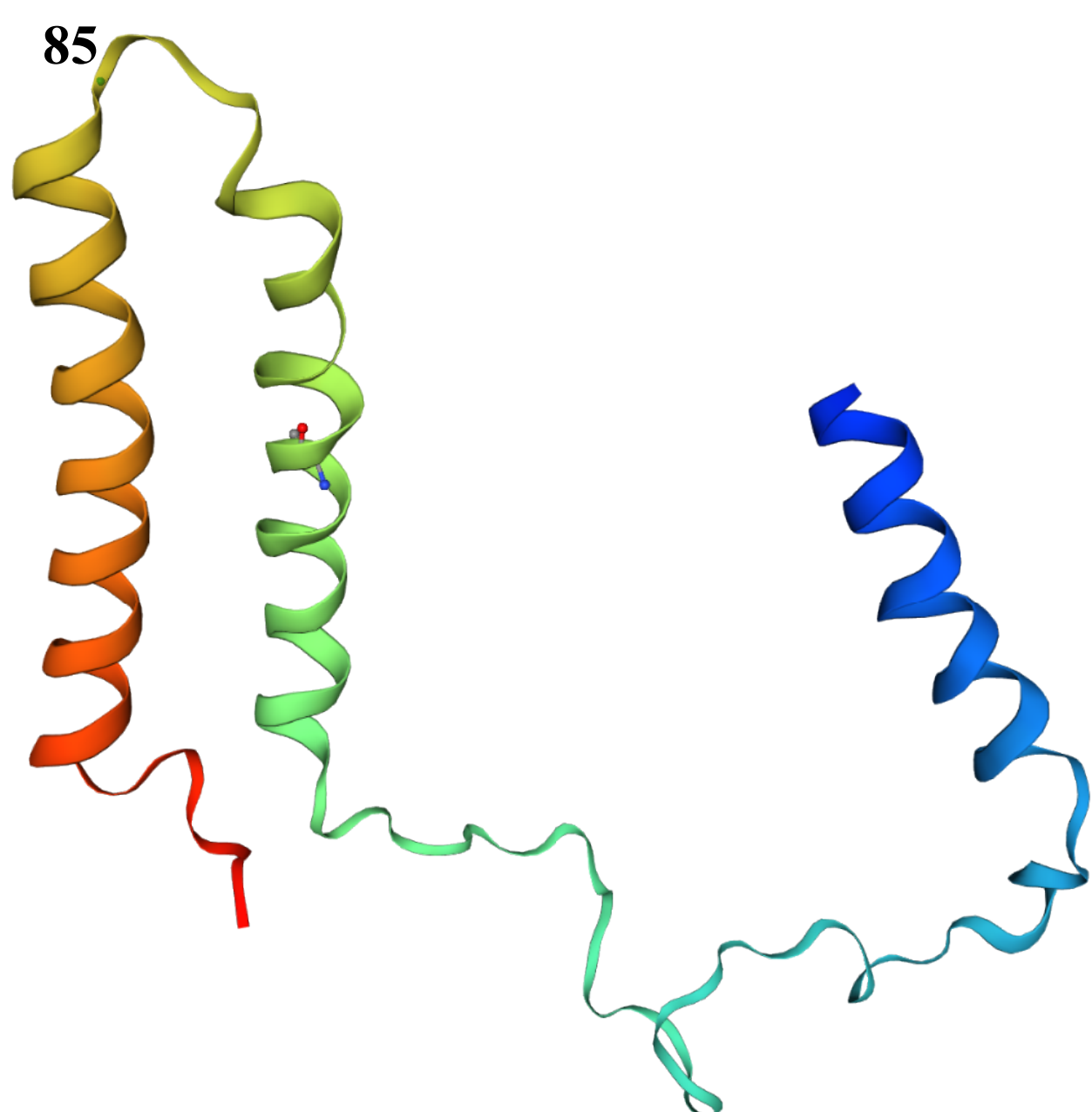
B. *Cyt b*



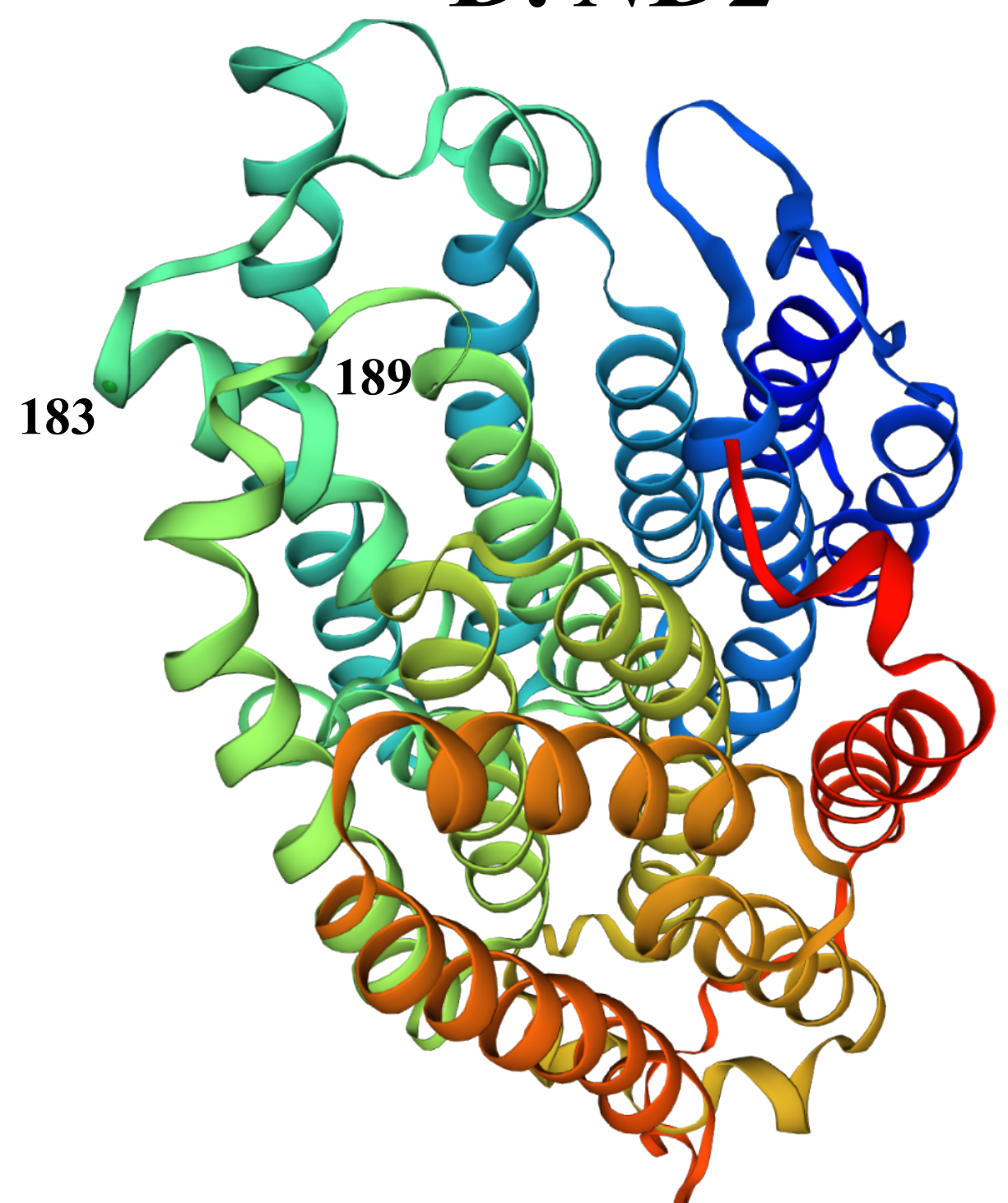
C. *ND1*



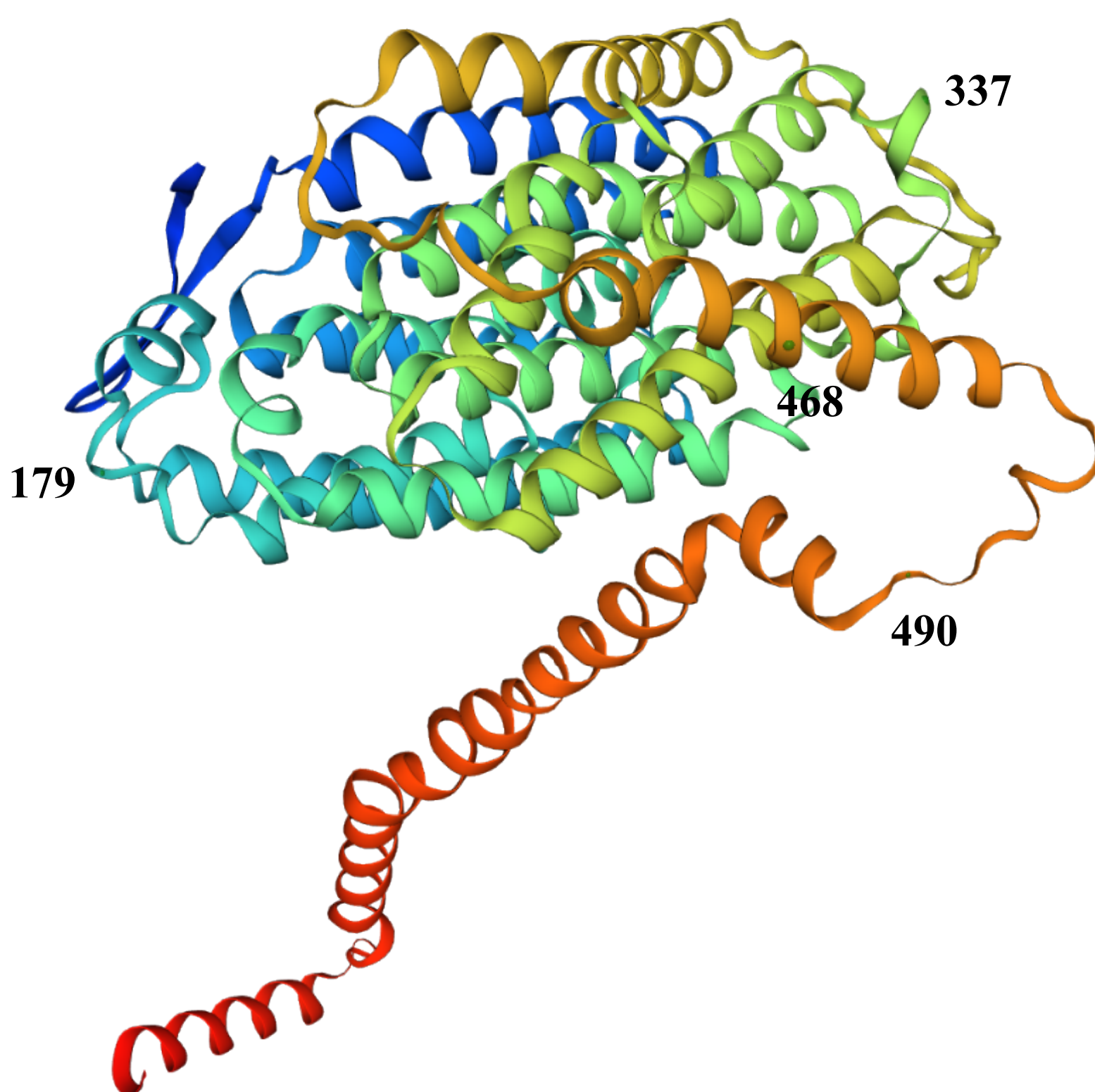
D. *ND2*



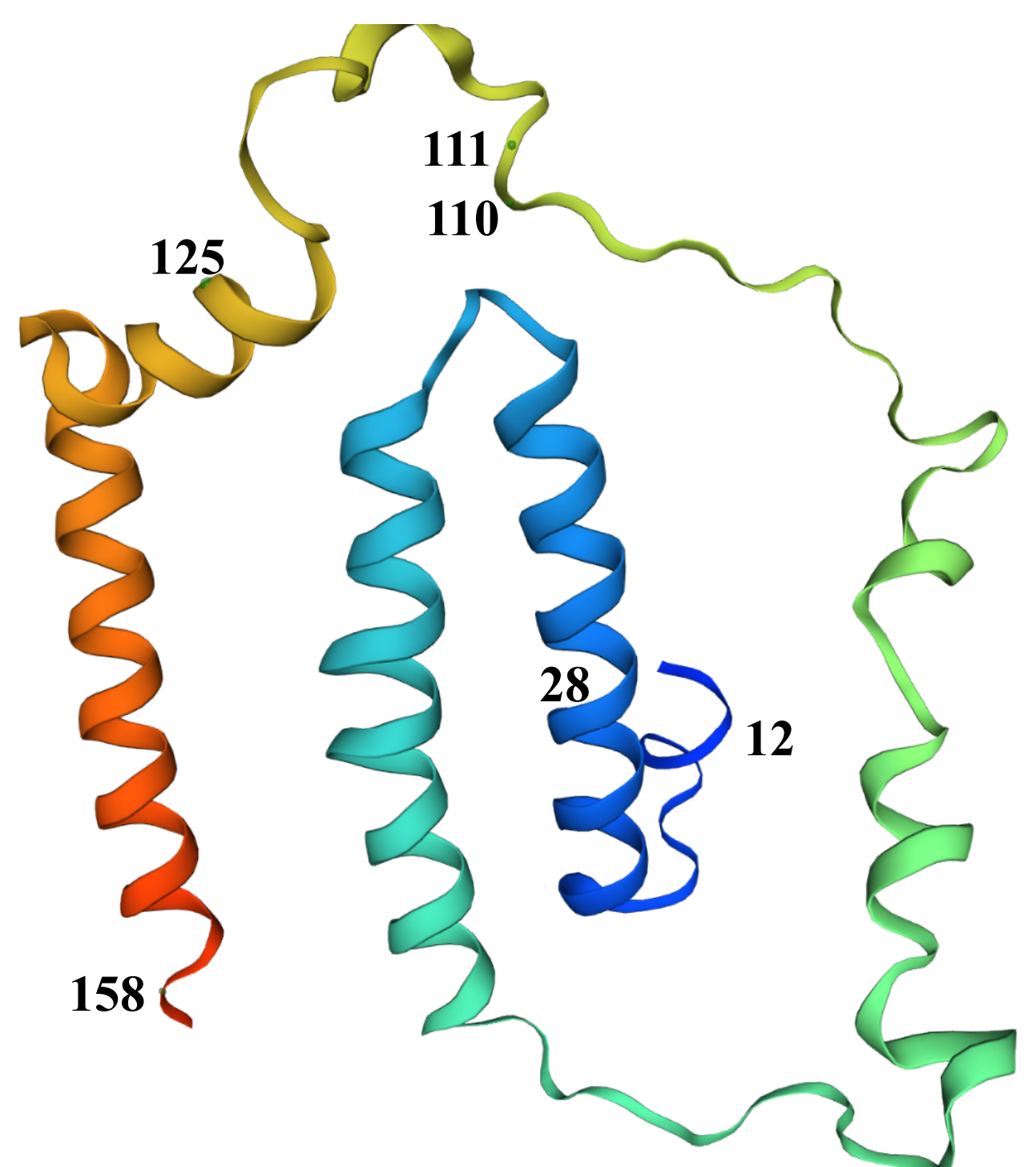
E. *ND3*



F. *ND4*



G. *ND5*



H. *ND6*

Note. The 318 and 341 amino acid sites at *ND2*, the 25 and 29 amino acid sites at *ND5*, along with the 3 amino acid site at *ND6* were not predicted due to the lack of corresponding template sequences.