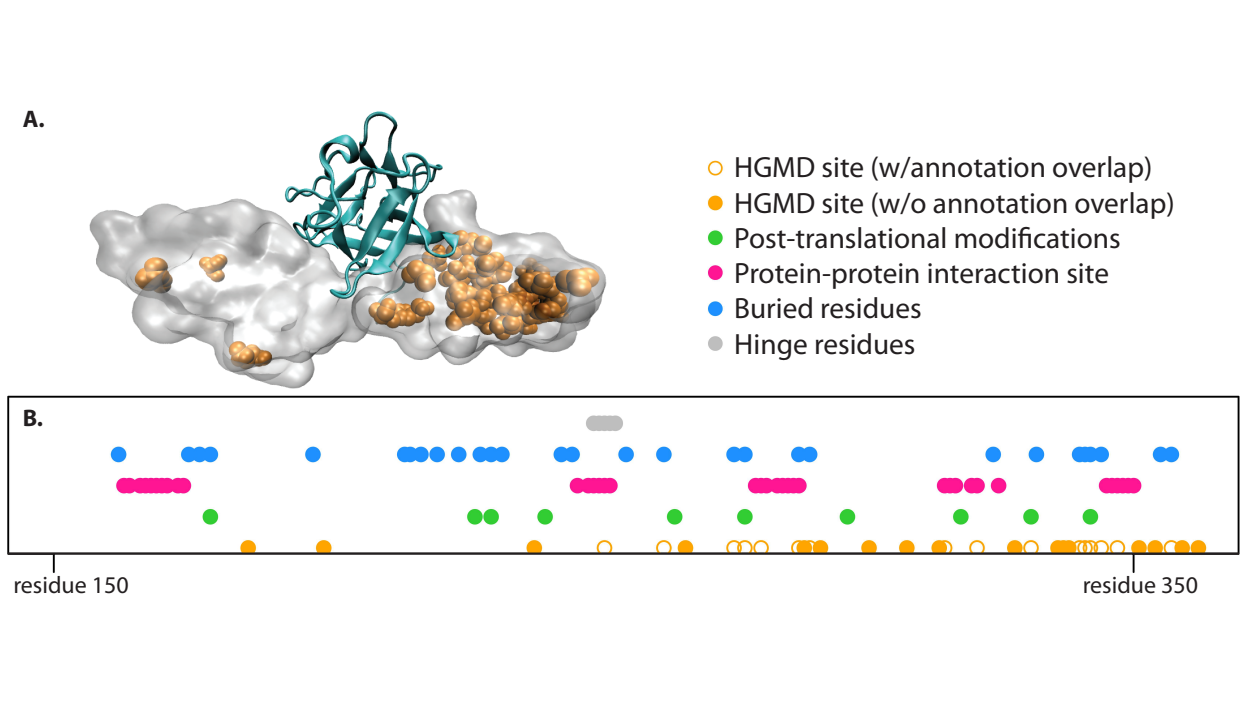
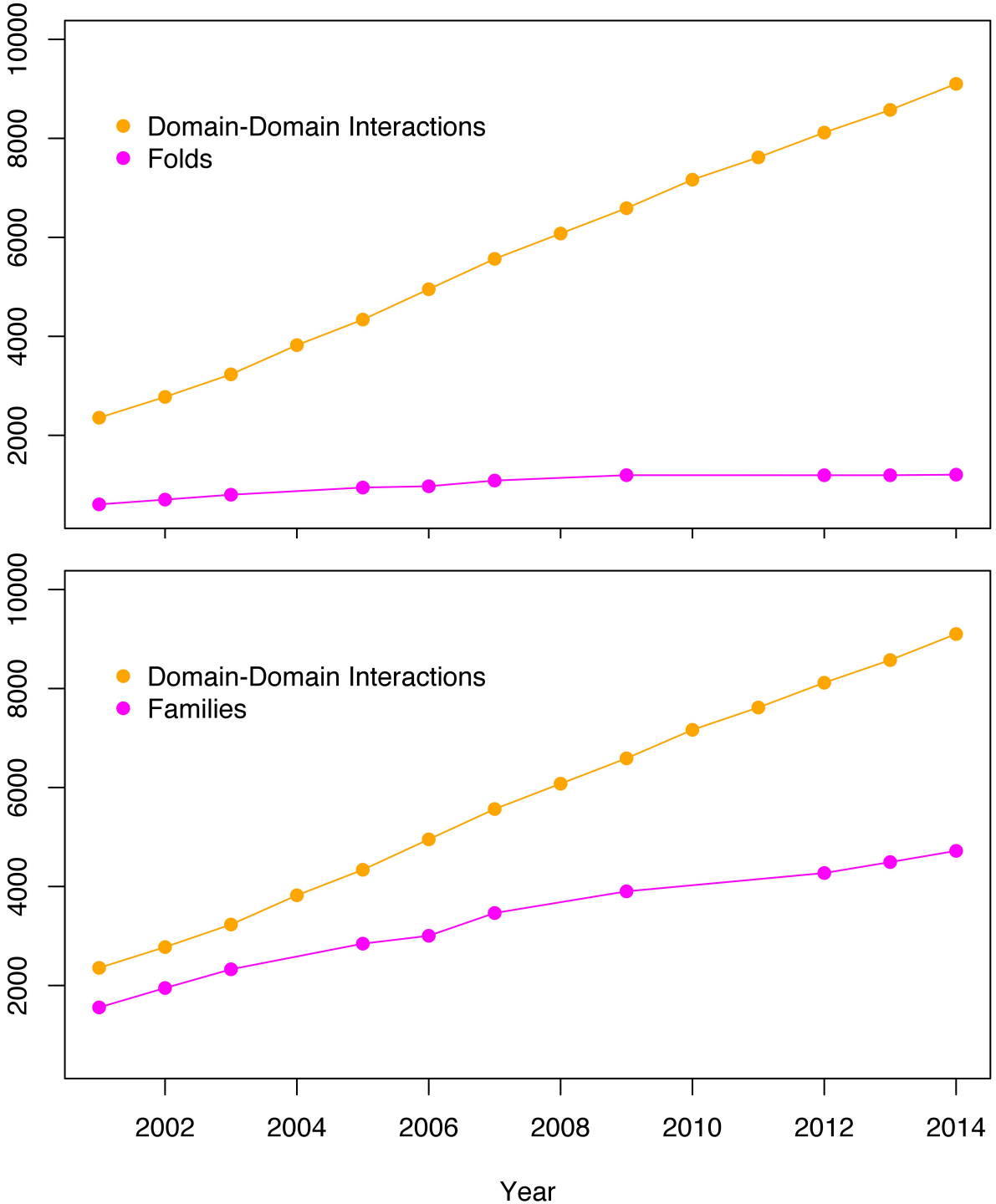
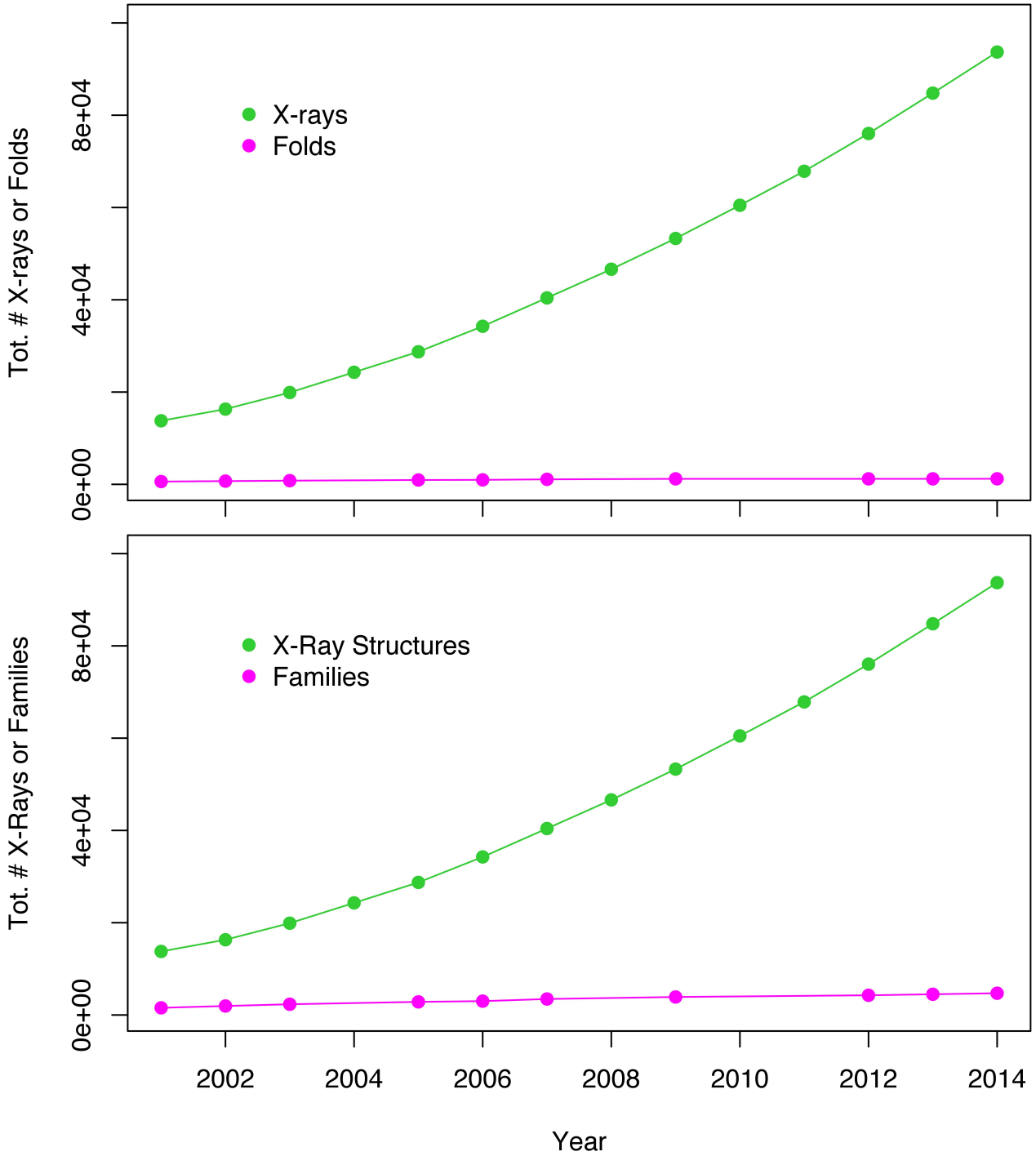

A.
HGMD site (w/annotation overlap)
HGMD site (w/o annotation overlap)
Post-translational modifications
Protein-protein interaction site
Buried residues
Hinge residues
B.
residue 150
residue 350


Domain-Domain Interactions
Folds
Domain-Domain Interactions
Families
2000
4000
6000
8000
10000
2002
2004
2006
2008
2010
2012
2014
Year


Tot. # X-rays or Folds
0e+00
4e+04
8e+04
X-rays
Folds
Tot. # X-Rays or Families
0e+00
4e+04
8e+04
X-Ray Structures
Families
2002
2004
2006
2008
2010
2012
2014
Year