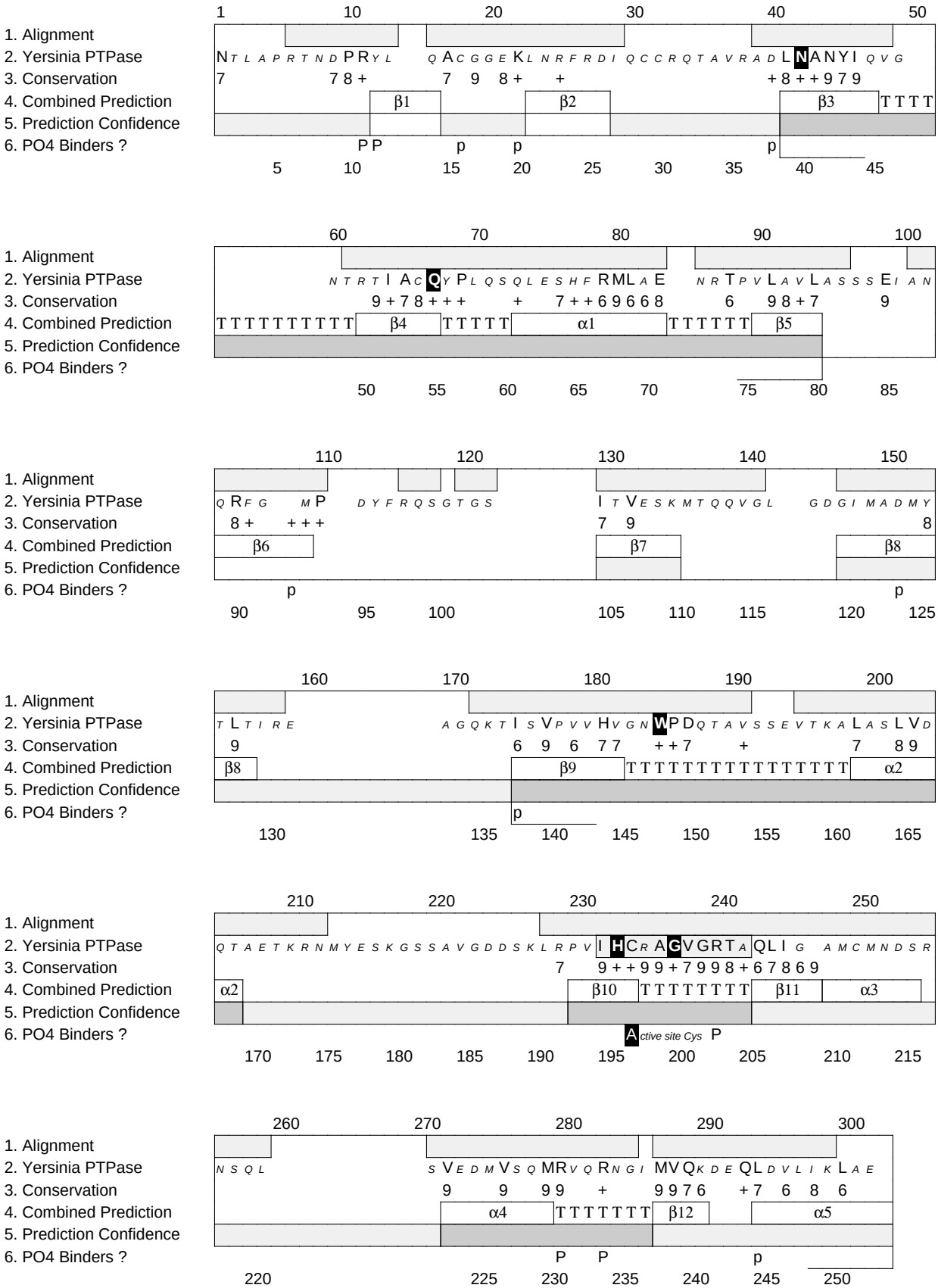
1. Alignment
2. Yersinia PTPase
3. Conservation
4. Combined Prediction
5. Prediction Confidence
6. PO4 Binders ?
β1
β2
β3
β4
α1
β5
β6
β7
β8
β9
α2
β10
β11
α3
α4
β12
α5
Active site Cys