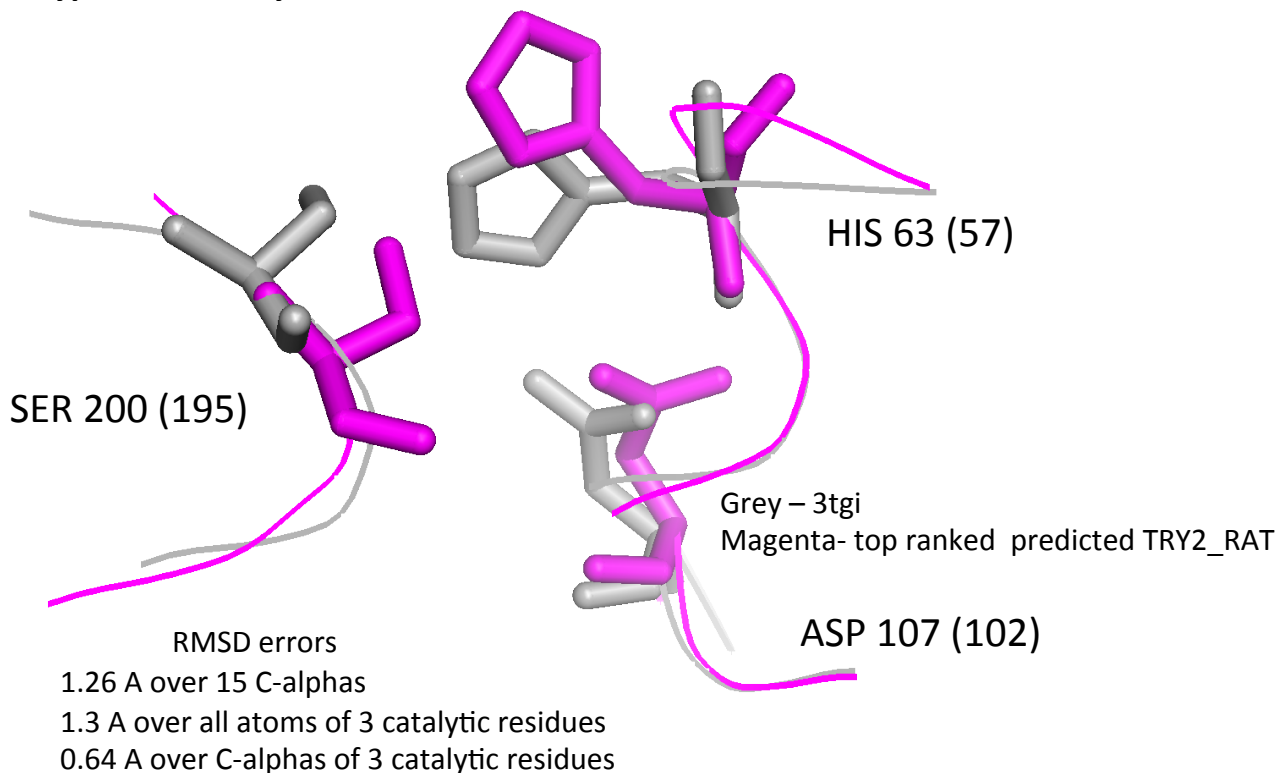
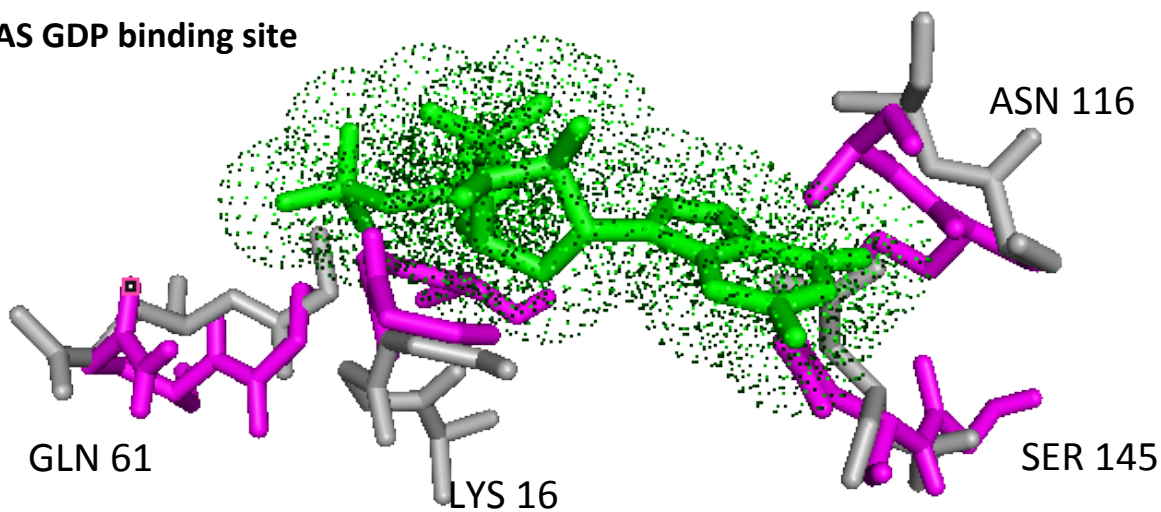


Figure S4. Active sites of top-ranked predicted Trypsin and Ras structures

A. Trypsin – 3 catalytic residues



B. RAS GDP binding site



Grey: 5p21, green: bound GDP, magenta: top ranked predicted Ras structure