





Variable                      Average                      Conserved

- e** - An exposed residue according to the NACSES algorithm.
- b** - A buried residue according to the NACSES algorithm.
- f** - A predicted functional residue (highly conserved and exposed).
- s** - A predicted structural residue (highly conserved and buried).
- x** - Insufficient data - the calculation for this site was performed on less than 10% of the sequences.