

[illegible]

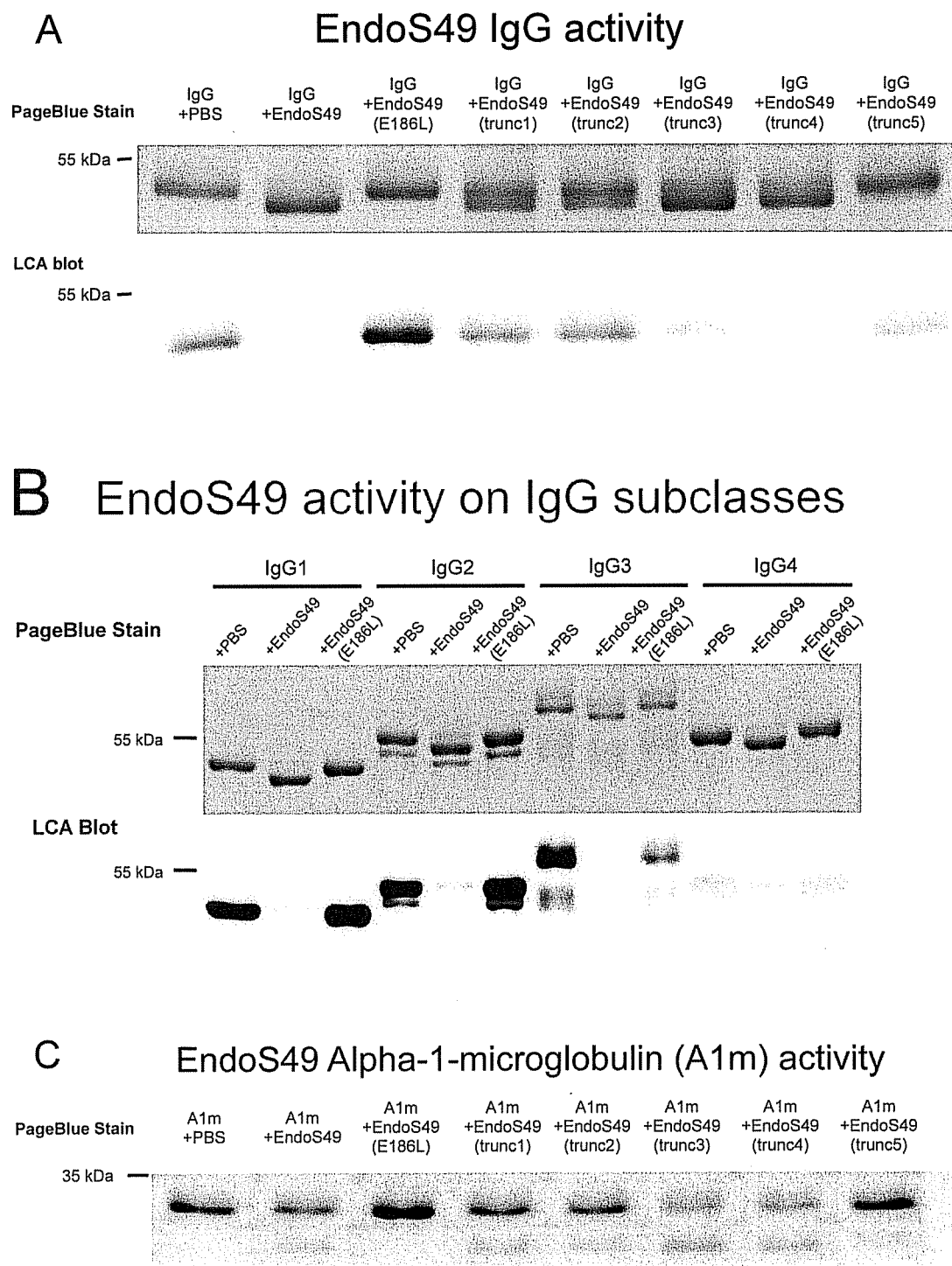
Figure 2.

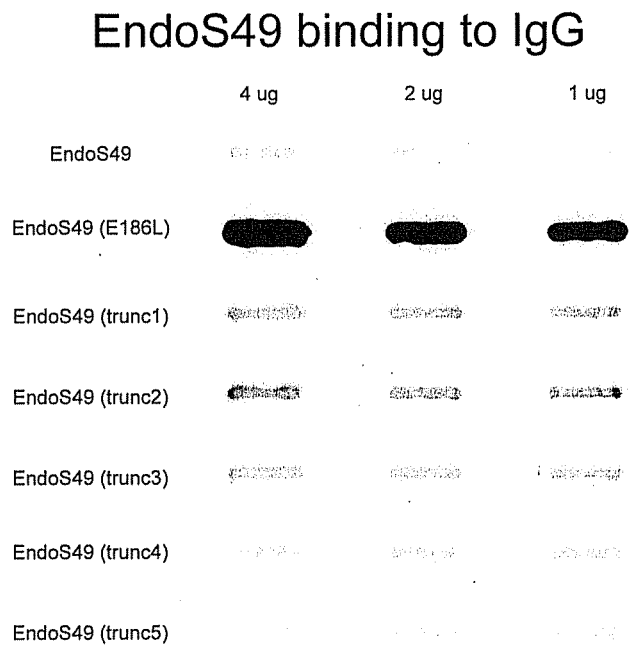
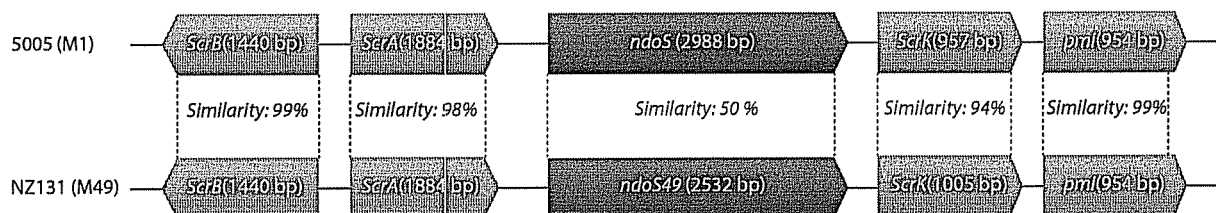
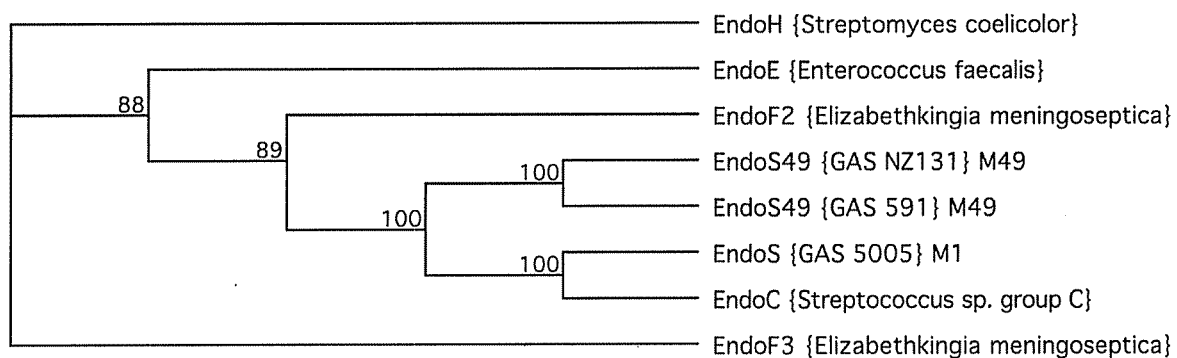
Figure 3.**Figure 4.**

Figure 5. Phylogenetic tree of endoglycosidases

Method: Neighbor Joining; Bootstrap (10000 reps); tie breaking = Systematic
 Distance: Uncorrected ("p")
 Gaps distributed proportionally

**Figure 6.**