

## MutationFinder Annotation in GATE

By combining random point mutagenesis, saturation mutagenesis, and DNA shuffling, a thermostable variant, Xylst, was identified which contained three amino acid substitutions: Q7H, N8F, and S179C.

Type	Set	Start	End	Id
Mutation		190	195	340 {normalized=S179C}

1 Annotations (0 selected) Select:

☒ Mutation  
☐ MutationFinderIn  
☐ MutationFinderOut  
☐ Sentence  
☐ SpaceToken  
☐ Split  
☐ Token  
☐ Original markups



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