## 6EQDA PETase

Depolymerase Acidovorax delafieldii hypothet protein Rhizobacter gummiphilus alpha/beta hydrolase Polyangium brachysporum lipase Streptomyces coelicolor A3(2) consensus

## logo

#### 6EQDA PETase

Depolymerase Acidovorax delafieldii hypothet protein Rhizobacter gummiphilus alpha/beta hydrolase Polyangium brachysporum lipase Streptomyces coelicolor A3(2) consensus

## logo

#### 6EQDA PETase

Depolymerase Acidovorax delafieldii hypothet protein Rhizobacter gummiphilus alpha/beta hydrolase Polyangium brachysporum lipase Streptomyces coelicolor A3(2) consensus

## logo

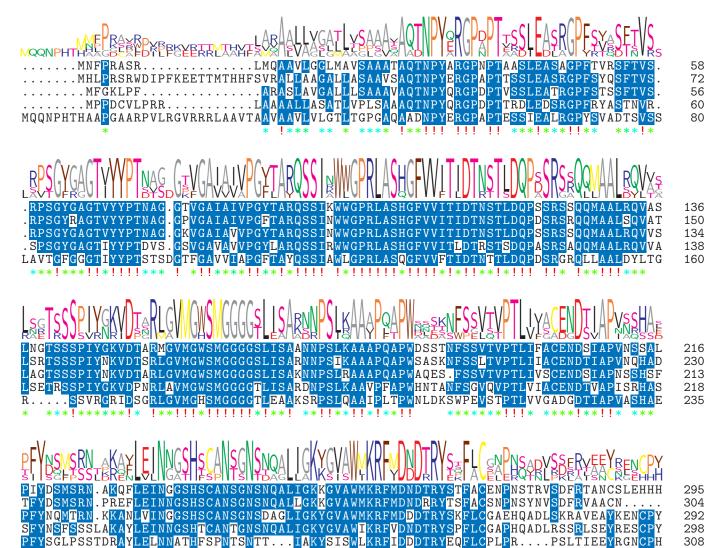
### 6EQDA PETase

Depolymerase Acidovorax delafieldii hypothet protein Rhizobacter gummiphilus alpha/beta hydrolase Polyangium brachysporum lipase Streptomyces coelicolor A3(2) consensus

## logo

#### 6EQDA PETase

Depolymerase Acidovorax delafieldii hypothet protein Rhizobacter gummiphilus alpha/beta hydrolase Polyangium brachysporum lipase Streptomyces coelicolor A3(2) consensus



# HSH

HHH. 298 .... 304 .... 292 .... 298 GS.. 310

non conserved

 $\geq$  50% conserved