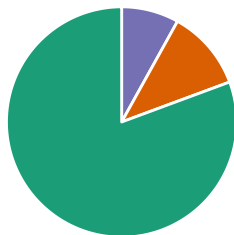
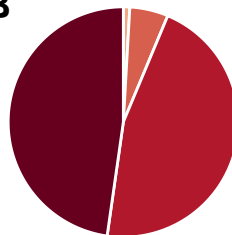
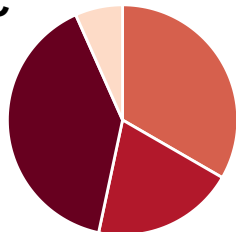


A**Group**

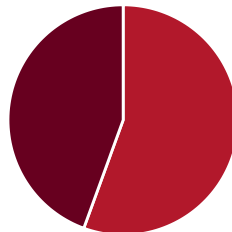
- All conserved motifs (80.65)%
- At least one conserved motif (11.29)%
- No conserved motifs (8.06)%

B**Group**

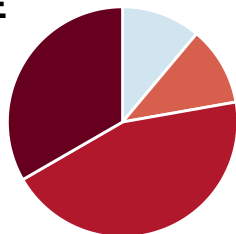
- Conserved-Region (47.75)%
- DNA-Binding-Region (45.95)%
- Nucleotide-Phosphate-Binding-Region (5.41)%
- Zn-Finger-Region (0.9)%

C**Group**

- Catalytic-Domain (6.67)%
- Conserved-Region (40)%
- DNA-Binding-Region (20)%
- Nucleotide-Phosphate-Binding-Region (33.33)%

D**Group**

- Conserved-Region (44.44)%
- DNA-Binding-Region (55.56)%

E**Group**

- Conserved-Region (33.33)%
- DNA-Binding-Region (44.44)%
- Nucleotide-Phosphate-Binding-Region (11.11)%
- Transmembrane-Region (11.11)%