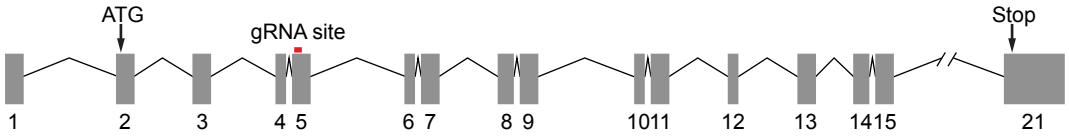
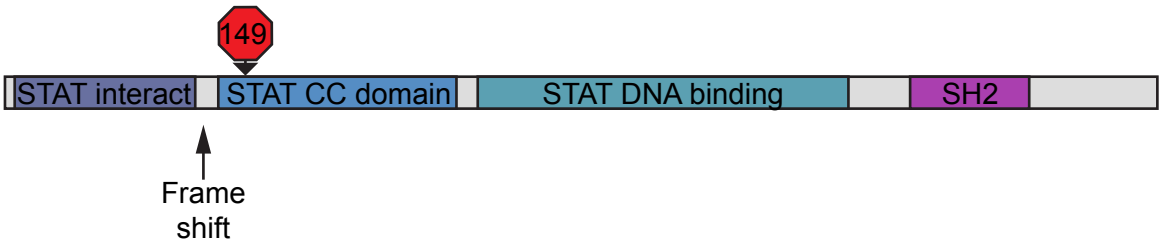
A



B

```
     N  M  Q  F  P  P  F  A  P  P  P  T  N  G  I  P  Q
361  AATATGCAGTTCCCTCCCTTCGCACCTCCACCTACCAACGGCATCCCACA 410
     ||||||||||||||||||||||||||          |||||||||||||||
361  AATATGCAGTTCCCTCCCTTCGCACC----------AACGGCATCCCACA 400
     N  M  Q  F  P  P  F  A  P            T  A  S  H  N

      Q  A  W  E  E  K  K  T  D  V  E  N  L  M  S  K  V
411  ACAAGCATGGGAGGAGAAAAAGACTGATGTGGAGAATCTAATGTCCAAAG 460
     ||||||||||||||||||||||||||||||||||||||||||||||||||
401  ACAAGCATGGGAGGAGAAAAAGACTGATGTGGAGAATCTAATGTCCAAAG 450
       K  H  G  R  R  K  R  L  M  W  R  I  *  C  P  K
```

C