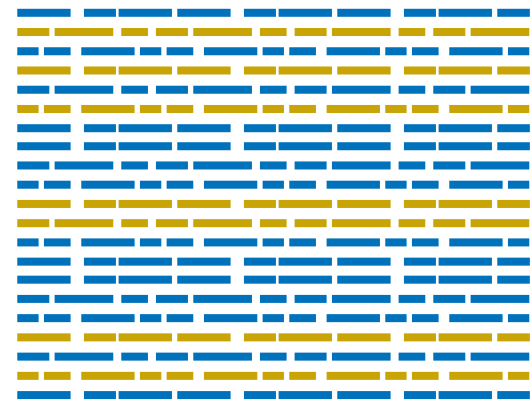
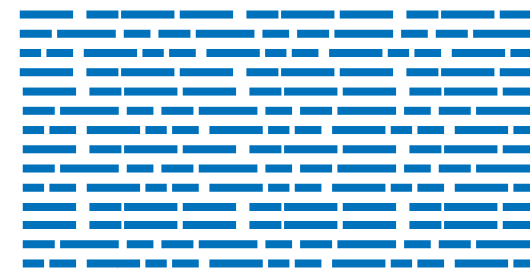


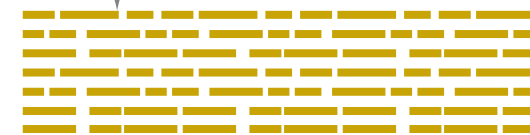
trusted “gold standard”  
alignment (152 sequences):



seed alignment  
(101 sequences):



no seed/test sequence  
pair is > 80% identical



test alignment  
(51 sequences)

