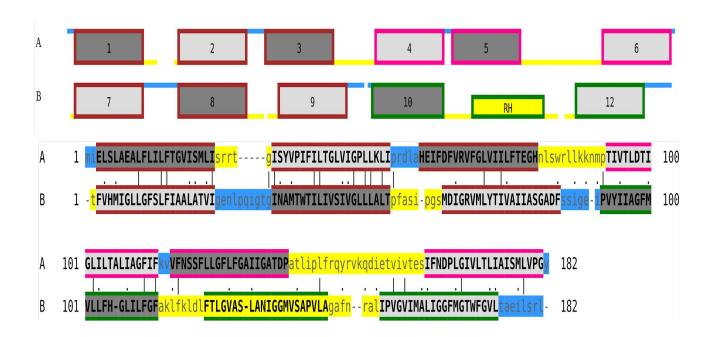
(a) A: Na_H_Exchanger_1-NR, B: DUF819-CR, Aligned helices: A: 1-6, B:7-12, E-value: 0.061



(b) A: Na_H_Exchanger_1-NR, B: DUF819-NR, Aligned helices: A:1-6, B:1-6, E-value: 1.7e-09

