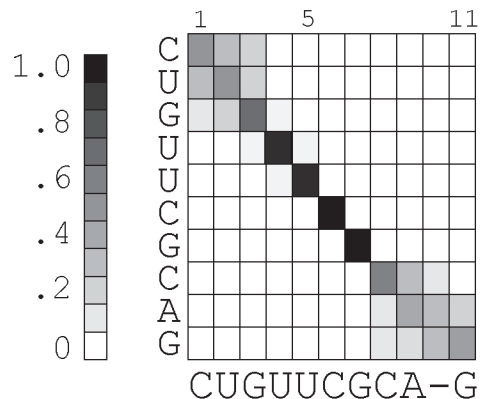
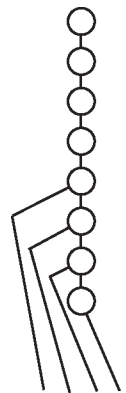


HMMs – Each column of seed alignment corresponds to a column of matrix.

yeast GUGUUCGCUAC
human -UCUUCGCG-
fly AGAUU-GUACU



states



struct <<<----->-->>
yeast GUGUUCGCUAC
human -UCUUCGCG-
fly AGAUU-GUACU

CMs – Each column of seed alignment corresponds to a state.

yeast human fly

U C U C U

U G U G U G

G•C C•G A•U

U•A^U U•G^C G•C^A

G•C A•U