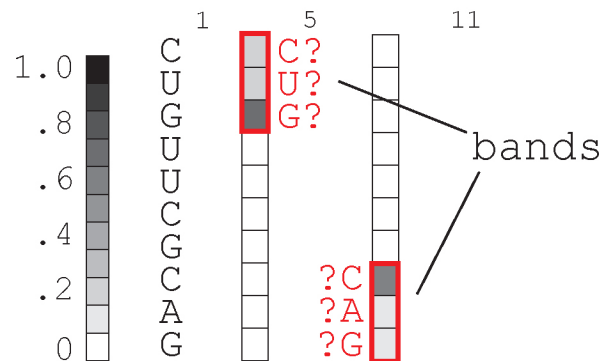


HMMs – Each column of seed alignment corresponds to a column of matrix.

yeast GUGUUCGCUAC
human -UCUUCGGCG-
fly AGAUU-GUACU



struct <<<--->-->>
yeast GUGUUCGCUAC
human -UCUUCGGCG-
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 • G	G • C ^A
G • C		A • U

