

I:

 $I0_i$: phase-0 insertion of state i $I1_i$: phase-1 insertion of state i $I2_i$: phase-2 insertion of state i

... : continuing states

 i_0 : phase-0 insertion opening weight i_1 : phase-1 insertion opening weight i_2 : phase-2 insertion opening weight

e: insertion extension weight in the unit of codon

 \emptyset :NNN: one codon insertion

N:N: nucleotide to nucleotide match