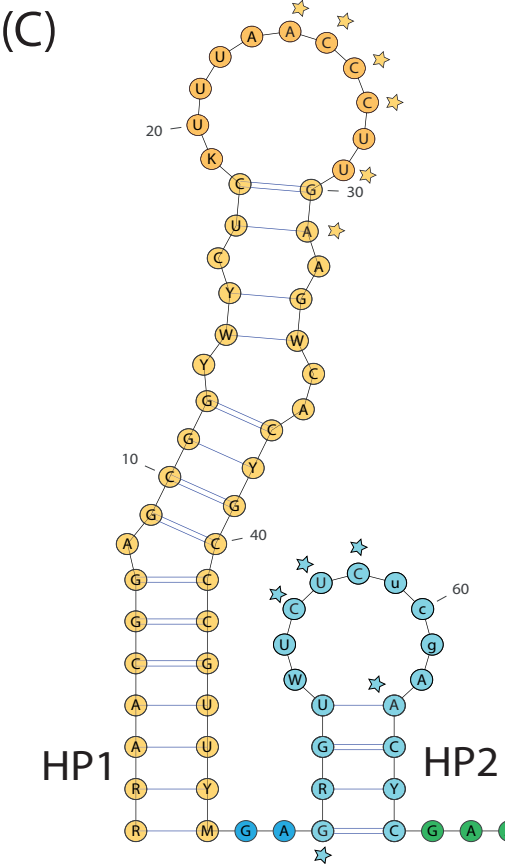


(A)

```
>RSSP1|maxrightloopextent=1|maxleftloopextent=1|maxmispair=6|weight=1
NNNNNNNNNNNNNNNNNNNNNNNACCCNUNANNNNNNNNNNNNNNNNN
(((((((.(.(((.(.(.(.....)).)).)).))))))))))
>RSSP2|maxrightloopextent=5|weight=1
GNNNNNCUCACNN
((((.....)))
>RSSP3|maxmispair=2|maxrightloopextent=2|weight=1
NNGGANCUNNNNNNNNNNNN
(((((((.(.....)))))))
```



(B)

```
parms
wc +=gu;
descr
h5(len=7)
ss(len=1)
h5(len=4)
ss(len=1)
h5(len=2)
ss(len=1)
h5(len=2)
ss(seq="N\{0,1\}NNNNACCCNUN\{0,1\}",minlen=10,maxlen=12)
h3(seq="NA",len=2)
ss(len=1)
h3(len=2)
ss(len=2)
h3(len=4)
h3(len=7)
ss(minlen=2,maxlen=3) #single strand between HP1 and HP2
h5(len=4)
ss(seq="NNCUCN\{0,5\}",minlen=5,maxlen=10)
h3(len=4)
ss(minlen=27,maxlen=31)#single strand between HP2 and HP3
h5(len=7, seq="NNGGANC",mispair=2,ends='mm')
ss(seq="UN\{4,6\}",minlen=5,maxlen=7)
h3(len=7)
```

