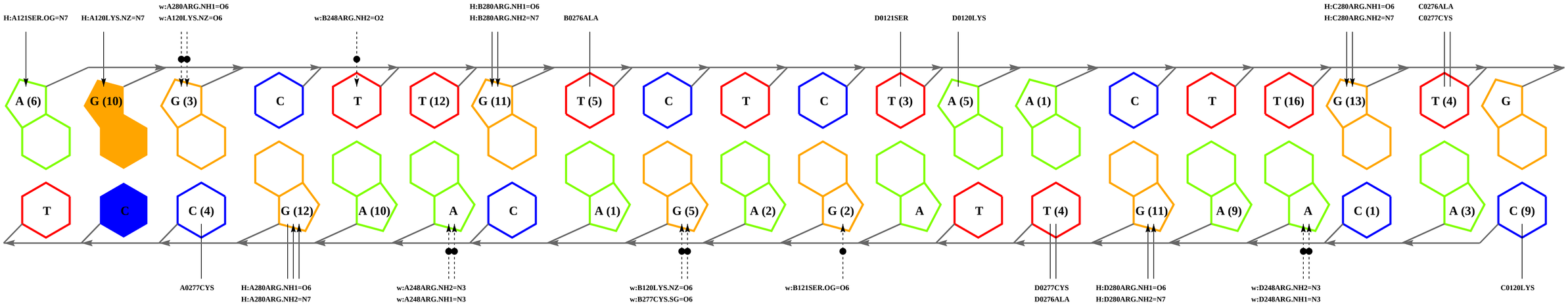
H:A121SER.OG=N7
H:A120LYS.NZ=N7
w:A280ARG.NH1=O6
w:A120LYS.NZ=O6
w:B248ARG.NH2=O2
H:B280ARG.NH1=O6
H:B280ARG.NH2=N7
B0276ALA
D0121SER
D0120LYS
H:C280ARG.NH1=O6
H:C280ARG.NH2=N7
C0276ALA
C0277CYS
A (6)
G (10)
G (3)
C
T
T (12)
G (11)
T (5)
C
T
C
T (3)
A (5)
A (1)
C
T
T (16)
G (13)
T (4)
G
T
C
C (4)
G (12)
A (10)
A
C
A (1)
G (5)
A (2)
G (2)
A
T
T (4)
G (11)
A (9)
A
C (1)
A (3)
C (9)
A0277CYS
H:A280ARG.NH1=O6
H:A280ARG.NH2=N7
A0276ALA
w:A248ARG.NH2=N3
w:A248ARG.NH1=N3
w:B120LYS.NZ=O6
w:B277CYS.SG=O6
w:B121SER.OG=O6
D0277CYS
D0276ALA
H:D280ARG.NH1=O6
H:D280ARG.NH2=N7
w:D248ARG.NH2=N3
w:D248ARG.NH1=N3
C0120LYS