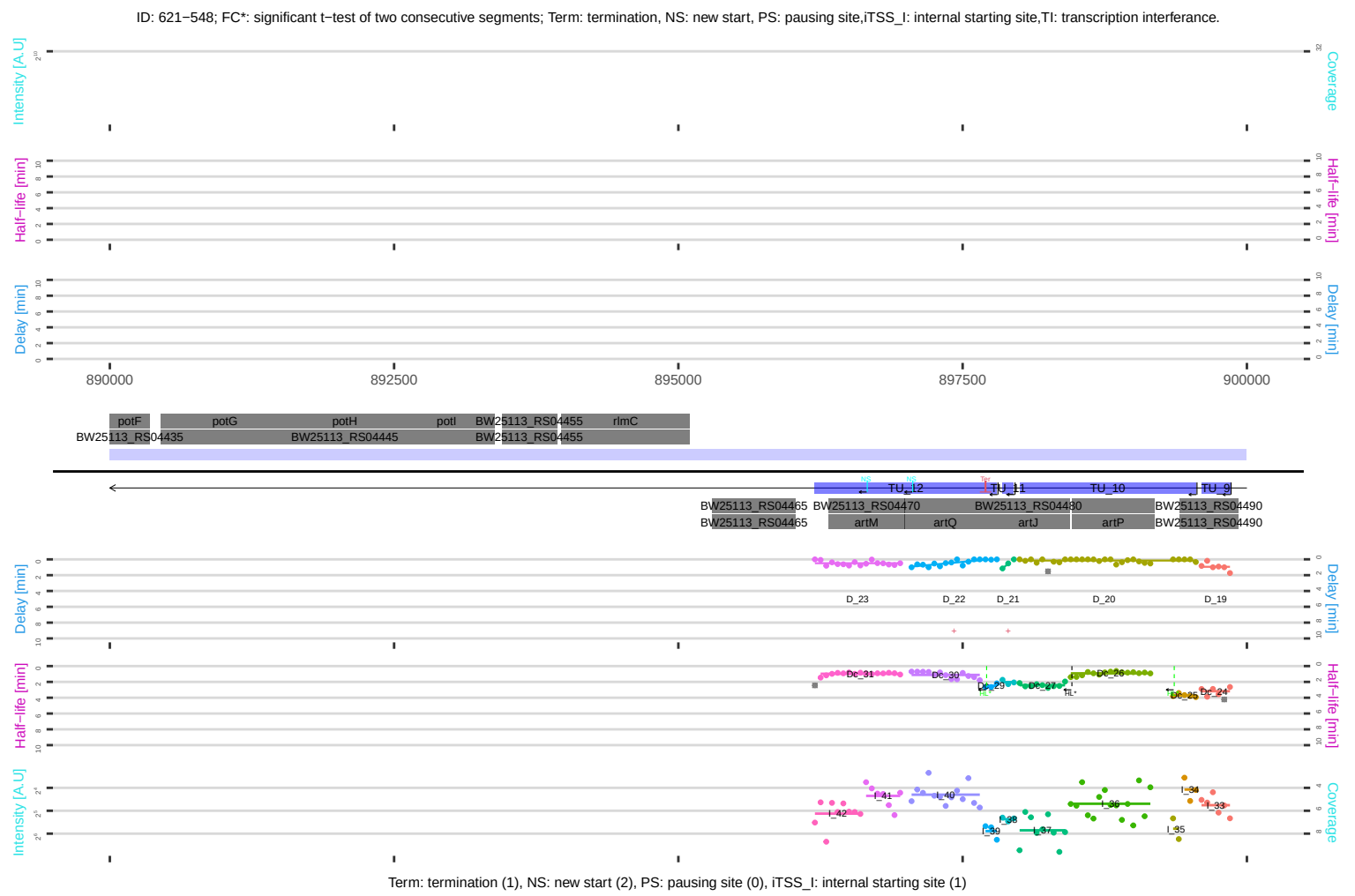
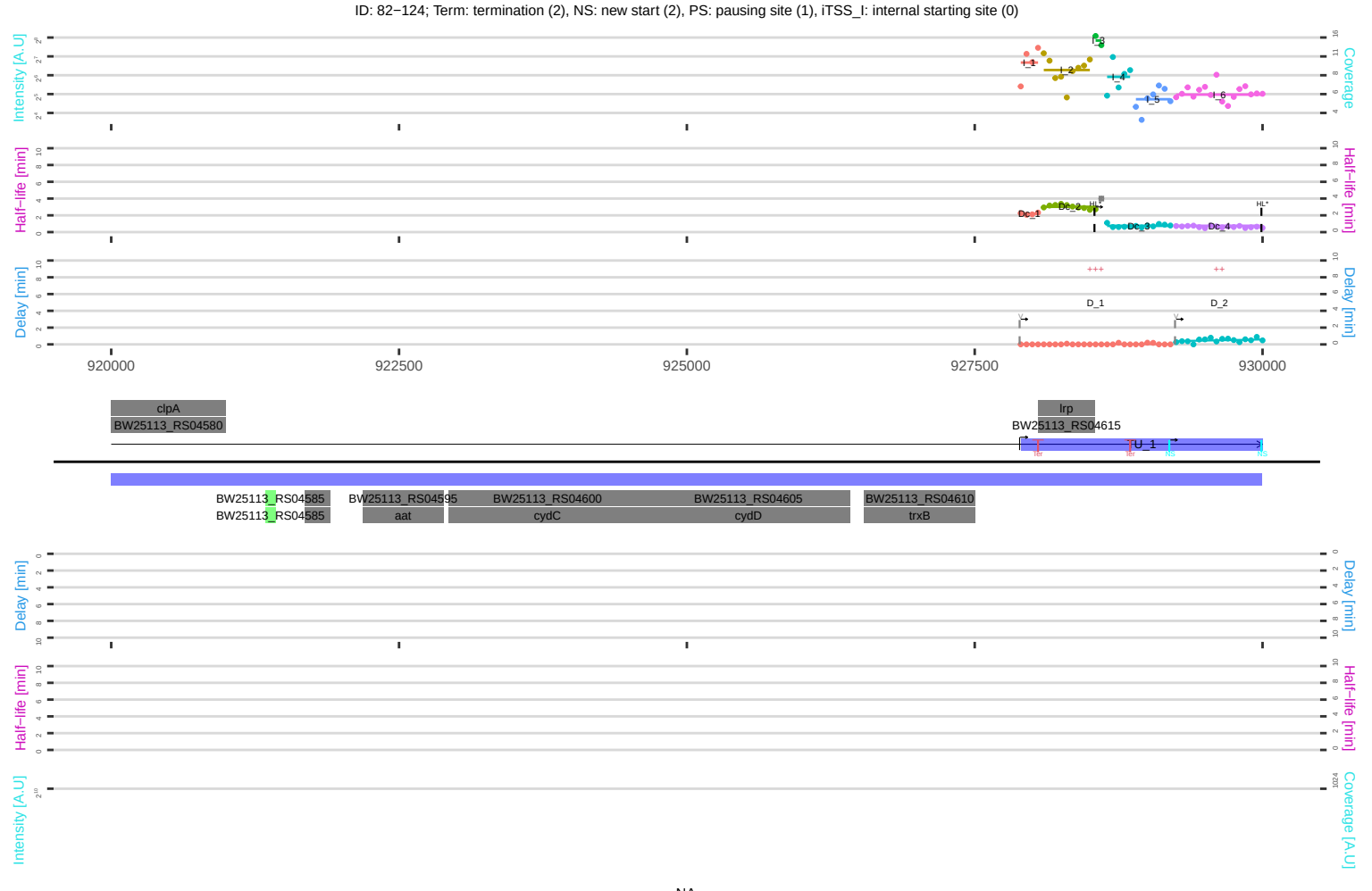
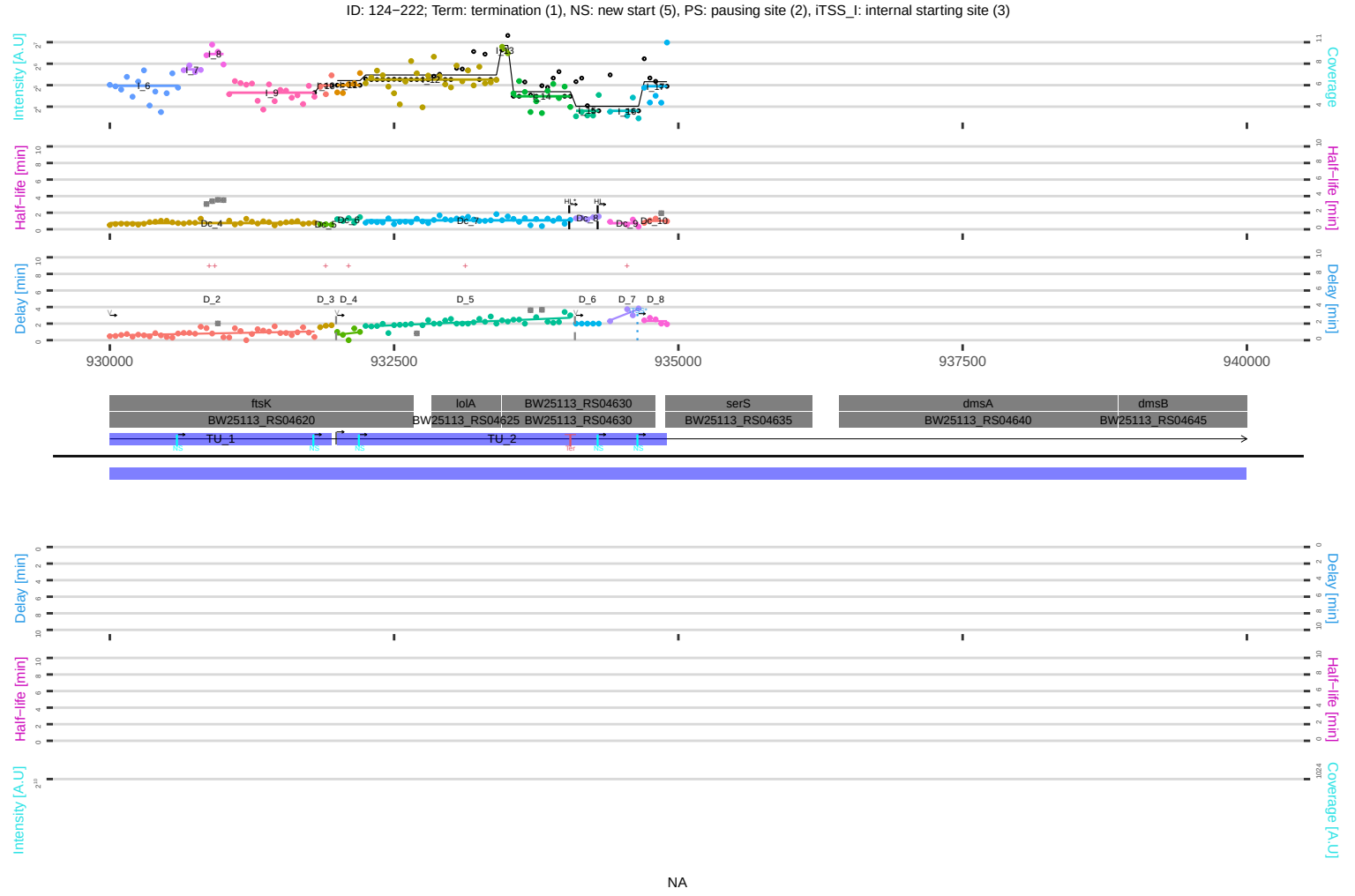
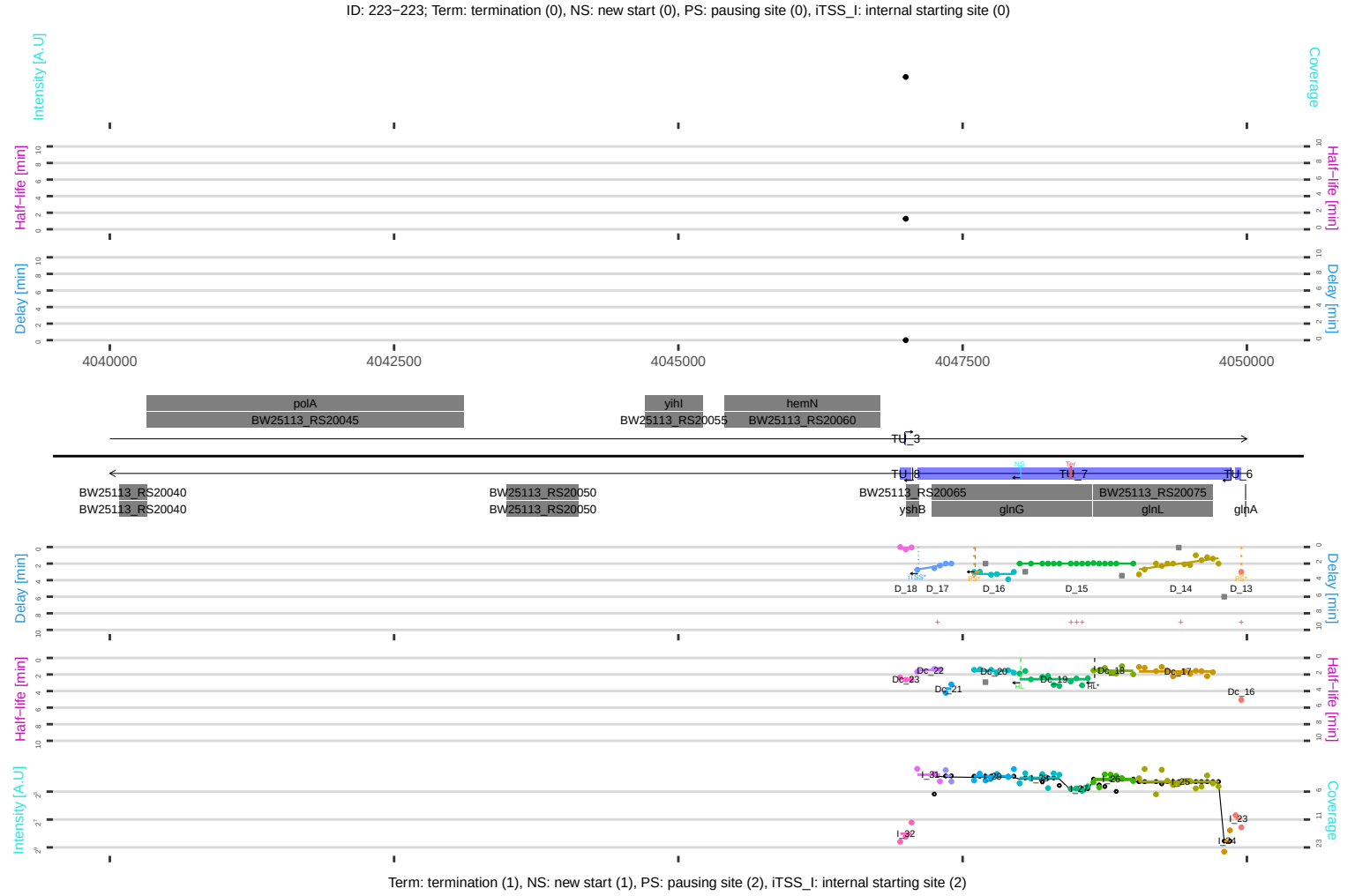

ID: 621−548; FC*: significant t−test of two consecutive segments; Term: termination, NS: new start, PS: pausing site,iTSS_I: internal starting site,TI: transcription interferance.
Intensity [A.U]
Coverage
Half−life [min]
Delay [min]
890000
892500
895000
897500
900000
potF
potG
potH
potI
BW25113_RS04455
rlmC
BW25113_RS04435
BW25113_RS04445
BW25113_RS04455
TU_12
TU_11
TU_10
TU_9
BW25113_RS04465
BW25113_RS04470
BW25113_RS04480
BW25113_RS04490
artM
artQ
artJ
artP
D_23
D_22
D_21
D_20
D_19
Dc_31
Dc_30
Dc_29
Dc_27
Dc_26
Dc_25
Dc_24
I_41
I_40
I_42
I_38
I_39
I_37
I_36
I_34
I_33
I_35
Term: termination (1), NS: new start (2), PS: pausing site (0), iTSS_I: internal starting site (1)


ID: 82−124; Term: termination (2), NS: new start (2), PS: pausing site (1), iTSS_I: internal starting site (0)
Intensity [A.U]
Coverage
Half−life [min]
Delay [min]
I_1
I_2
I_3
I_4
I_5
I_6
Dc_1
Dc_2
Dc_3
Dc_4
HL*
D_1
D_2
+++
++
920000
922500
925000
927500
930000
clpA
BW25113_RS04580
lrp
BW25113_RS04615
U_1
Ter
NS
BW25113_RS04585
BW25113_RS04595
BW25113_RS04600
BW25113_RS04605
BW25113_RS04610
aat
cydC
cydD
trxB
NA


ID: 124−222; Term: termination (1), NS: new start (5), PS: pausing site (2), iTSS_I: internal starting site (3)
Intensity [A.U]
Coverage
Half−life [min]
Delay [min]
I_1
I_2
I_3
I_4
I_5
I_6
I_7
I_8
I_9
I_10
I_11
I_12
I_13
I_14
I_15
I_16
I_17
Dc_1
Dc_2
Dc_3
Dc_4
Dc_5
Dc_6
Dc_7
Dc_8
Dc_9
Dc_10
HL
D_1
D_2
D_3
D_4
D_5
D_6
D_7
D_8
930000
932500
935000
937500
940000
ftsK
BW25113_RS04620
lolA
BW25113_RS04625
BW25113_RS04630
BW25113_RS04630
serS
BW25113_RS04635
dmsA
BW25113_RS04640
dmsB
BW25113_RS04645
TU_1
TU_2
NS
Ter
NA


ID: 223−223; Term: termination (0), NS: new start (0), PS: pausing site (0), iTSS_I: internal starting site (0)
Intensity [A.U]
Coverage
Half−life [min]
Delay [min]
4040000
4042500
4045000
4047500
4050000
polA
BW25113_RS20045
yihI
BW25113_RS20055
hemN
BW25113_RS20060
TU_3
TU_8
TU_7
TU_6
BW25113_RS20040
BW25113_RS20040
BW25113_RS20050
BW25113_RS20050
BW25113_RS20065
yshB
glnG
BW25113_RS20075
glnL
glnA
D_18
D_17
D_16
D_15
D_14
D_13
Dc_22
Dc_23
Dc_21
Dc_20
Dc_19
Dc_18
Dc_17
Dc_16
I_31
I_32
I_23
Term: termination (1), NS: new start (1), PS: pausing site (2), iTSS_I: internal starting site (2)

ID: 343−314; FC*: significant t−test of two consecutive segments; Term: termination, NS: new start, PS: pausing site,iTSS_I: internal starting site,TI: transcription interferance.
Intensity [A.U]
Coverage
Half−life [min]
Delay [min]
4050000
4052500
4055000
4057500
4060000
typA
BW25113_RS20090
BW25113_RS20095
BW25113_RS20095
BW25113_RS20105
BW25113_RS20105
TU_6
TU_5
BW25113_RS20080
glnA
BW25113_RS20110
ompL
BW25113_RS20115
BW25113_RS20115
BW25113_RS20120
BW25113_RS20120
D_13
D_12
D_11
Dc_16
Dc_15
Dc_14
Dc_13
Dc_12
I_19
I_20
Term: termination (0), NS: new start (0), PS: pausing site (0), iTSS_I: internal starting site (0)