

ID: 1-72; FC*: significant t-test of two consecutive segments;

Term: termination (2), NS: new start (0), PS: pausing site (0), iTSS_l: internal starting site (1), (*): p-value below 0.05; TI: transcription interference.



ID: 73–153; Term: termination (1), NS: new start (2), PS: pausing site (2), iTSS_I: internal starting site (1)



Term: termination (2), NS: new start (1), PS: pausing site (2), iTSS_I: internal starting site (2)

ID: 154–162; Term: termination (0), NS: new start (0), PS: pausing site (0), iTSS_I: internal starting site (0)

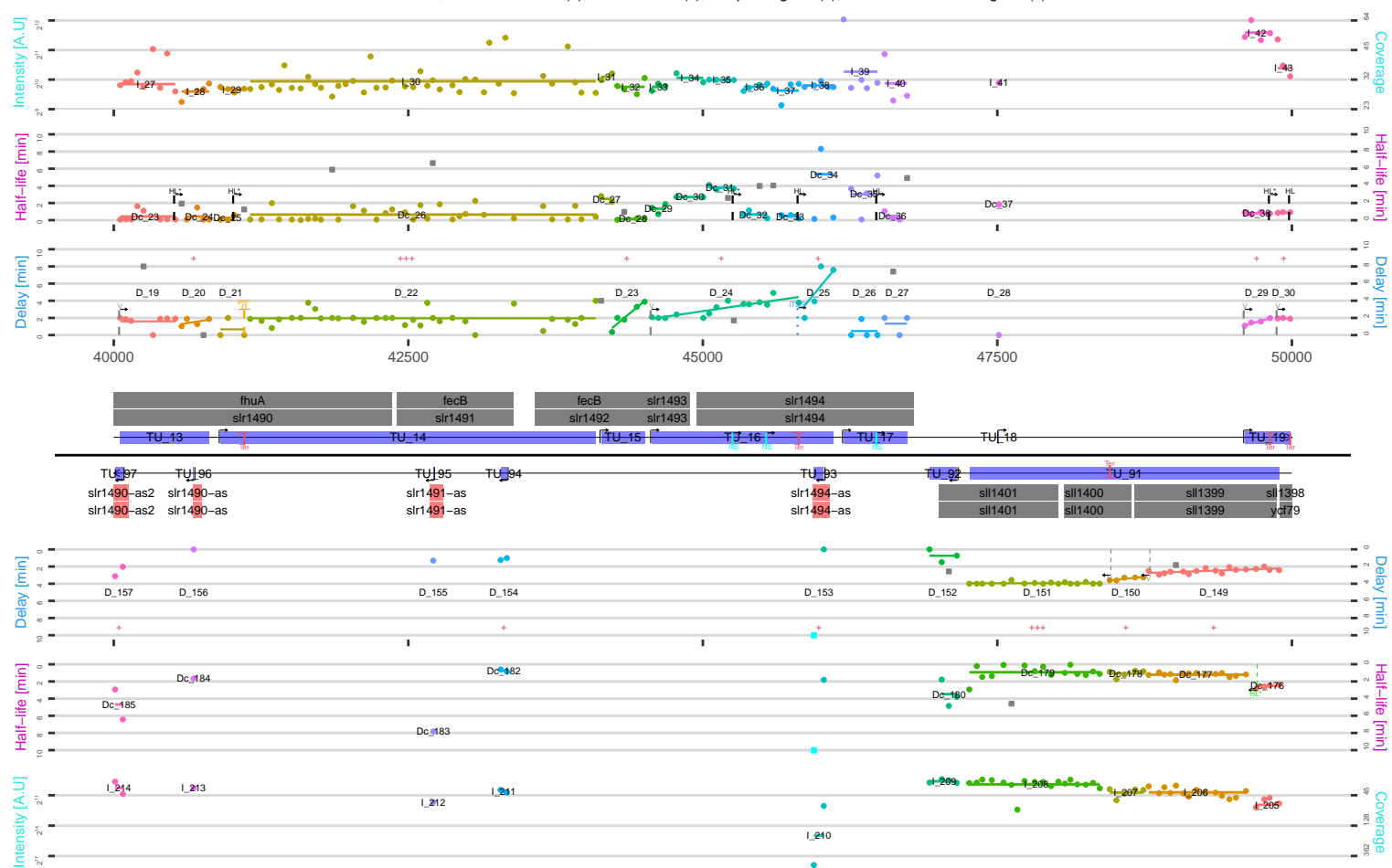


ID: 163–258; Term: termination (2), NS: new start (1), PS: pausing site (0), iTSS_I: internal starting site (2)

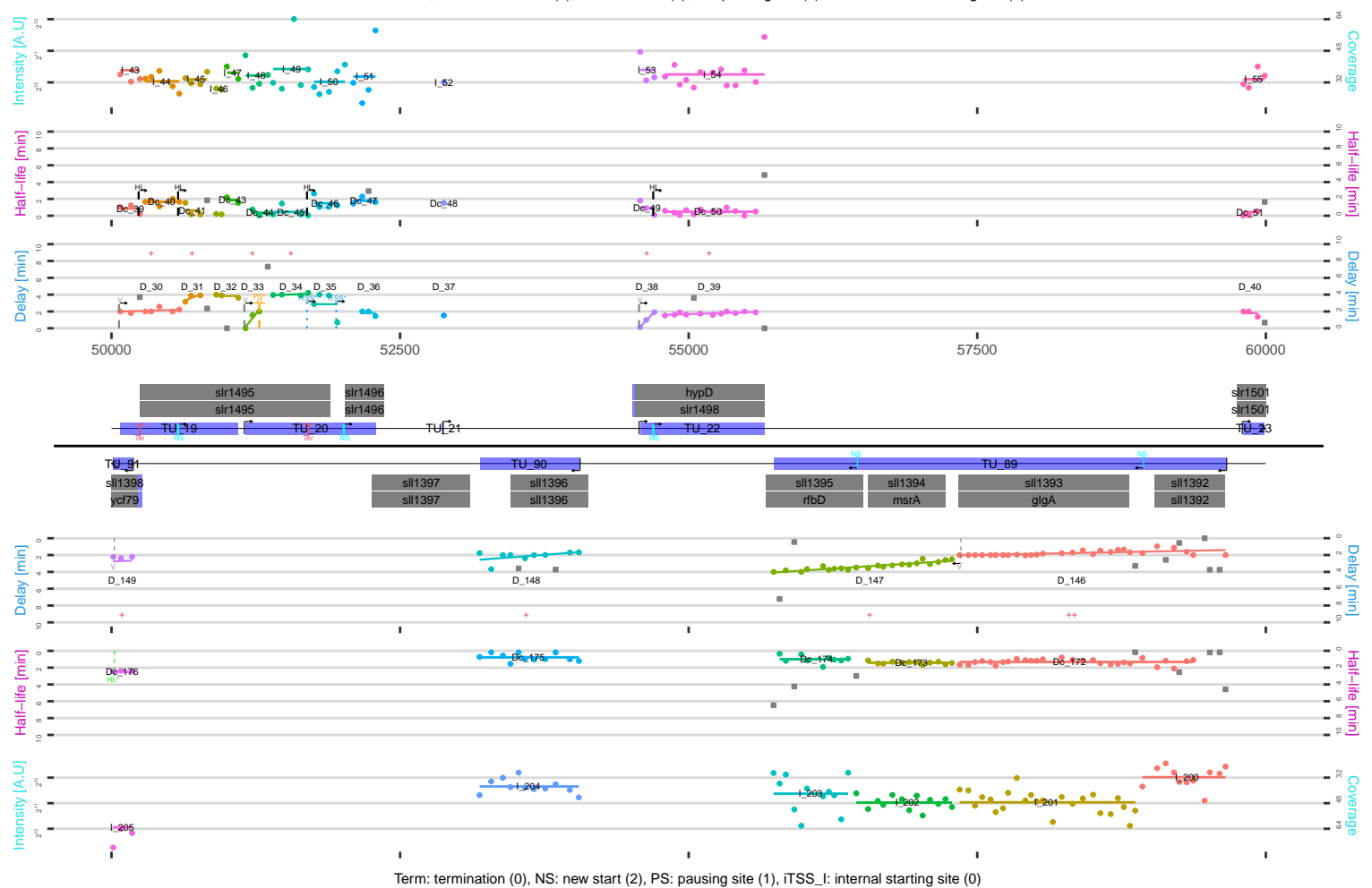


Term: termination (1), NS: new start (1), PS: pausing site (0), iTSS_I: internal starting site (0)

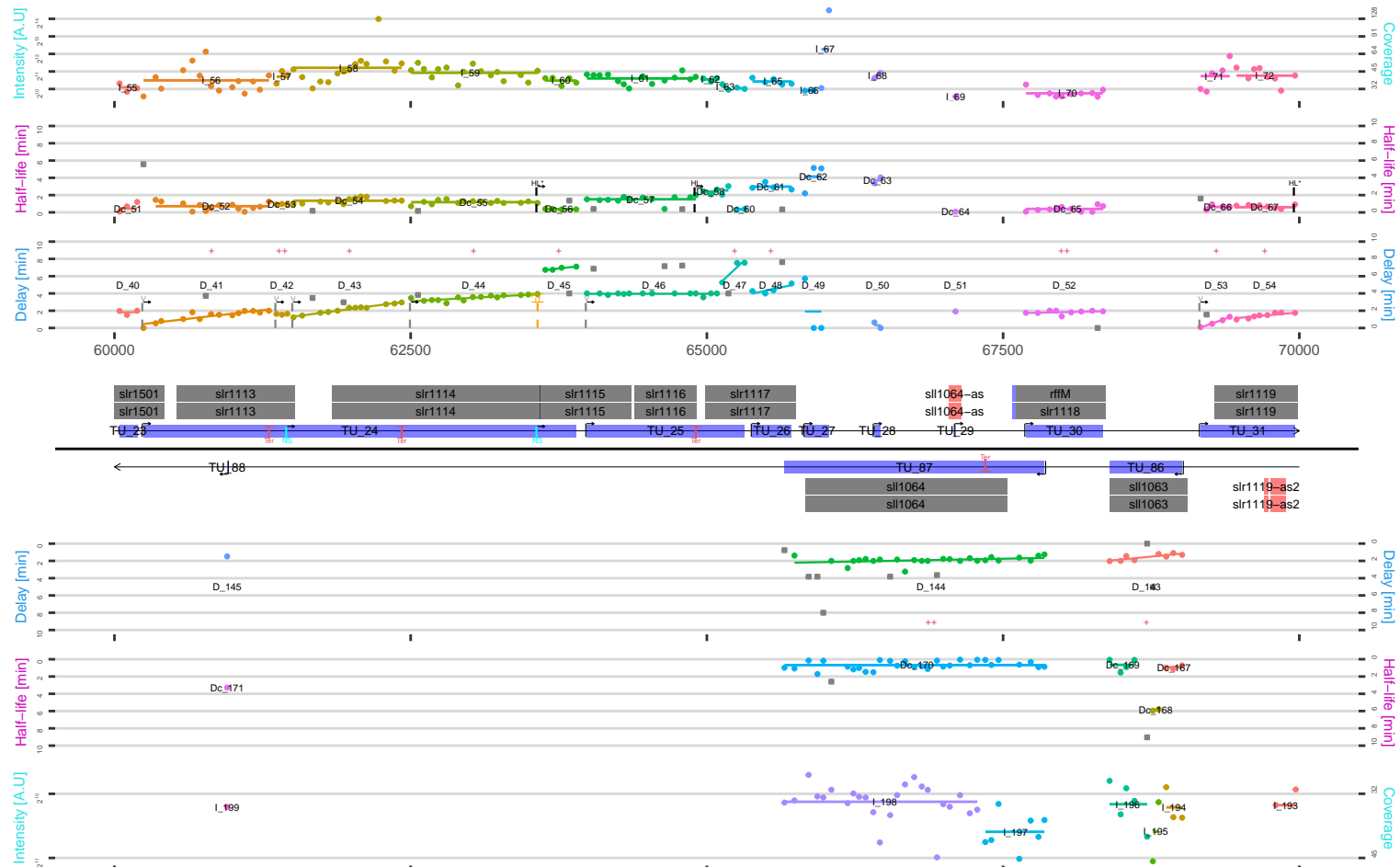
ID: 259-361; Term: termination (4), NS: new start (3), PS: pausing site (2), iTSS_L: internal starting site (3)



ID: 362-416; Term: termination (2), NS: new start (3), PS: pausing site (2), iTSS_L: internal starting site (4)

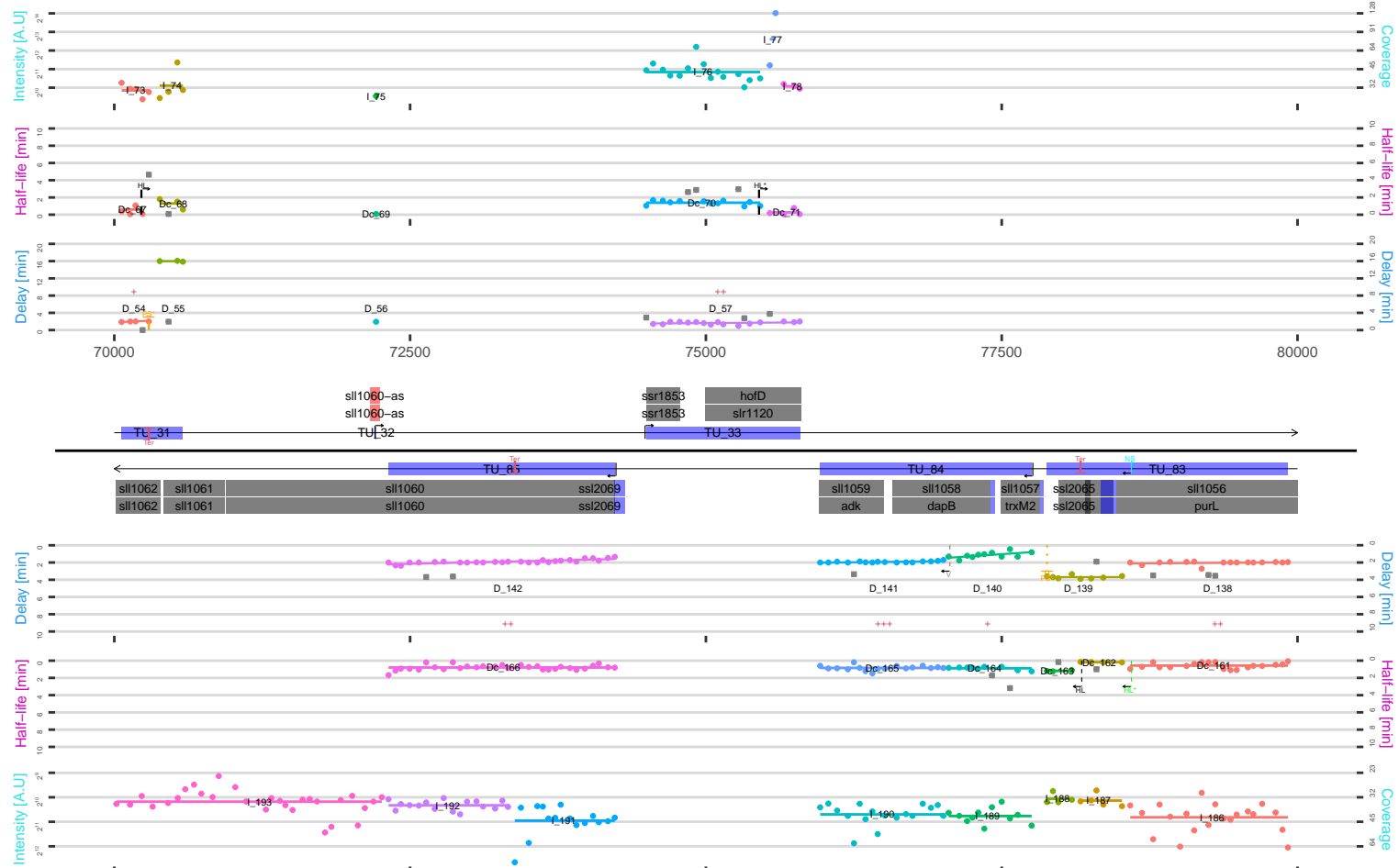


ID: 417-529; Term: termination (3), NS: new start (2), PS: pausing site (0), iTSS_L: internal starting site (3)



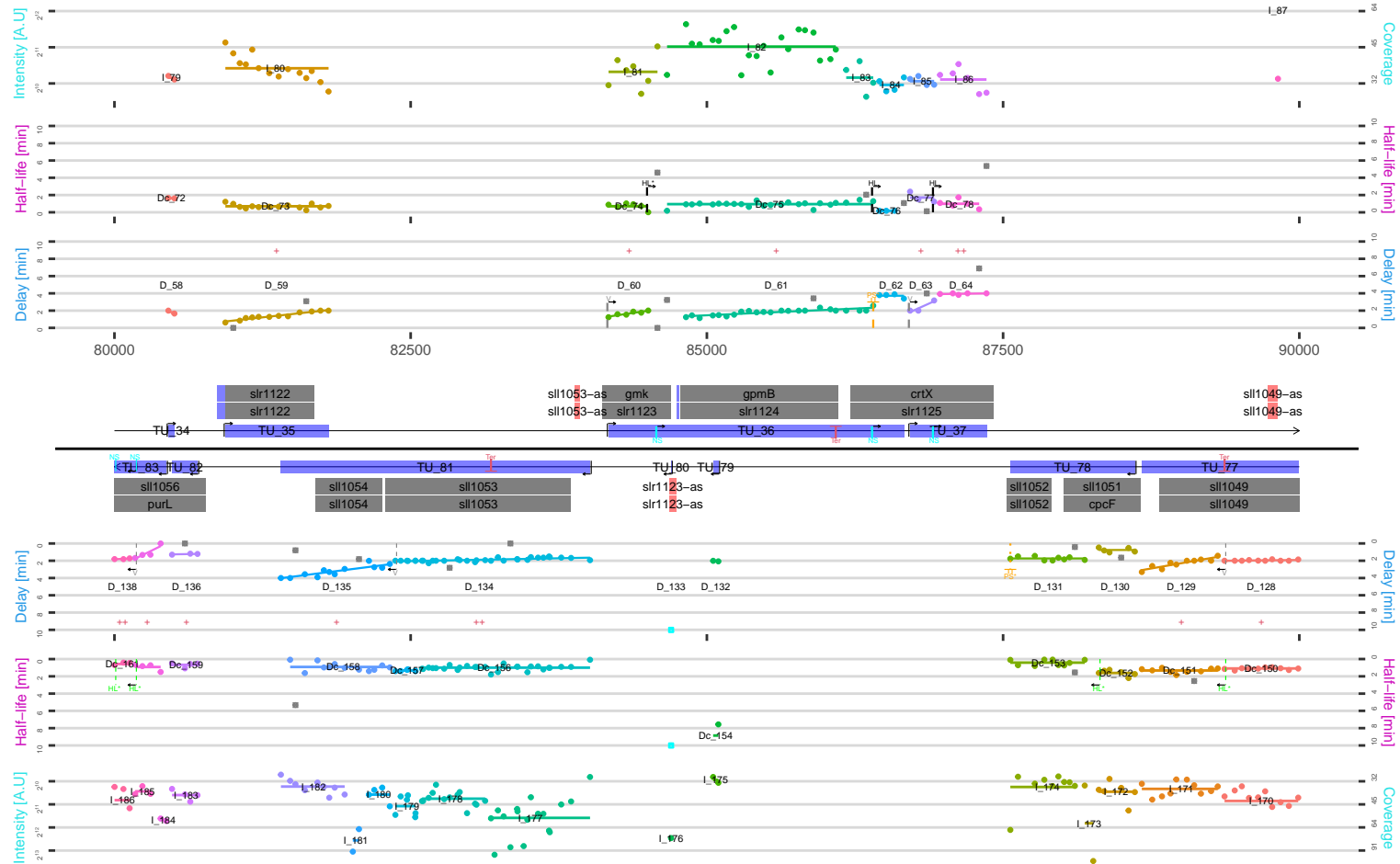
Term: termination (1), NS: new start (0), PS: pausing site (0), iTSS_L: internal starting site (0)

ID: 530–559; Term: termination (1), NS: new start (0), PS: pausing site (1), iTSS_L: internal starting site (0)

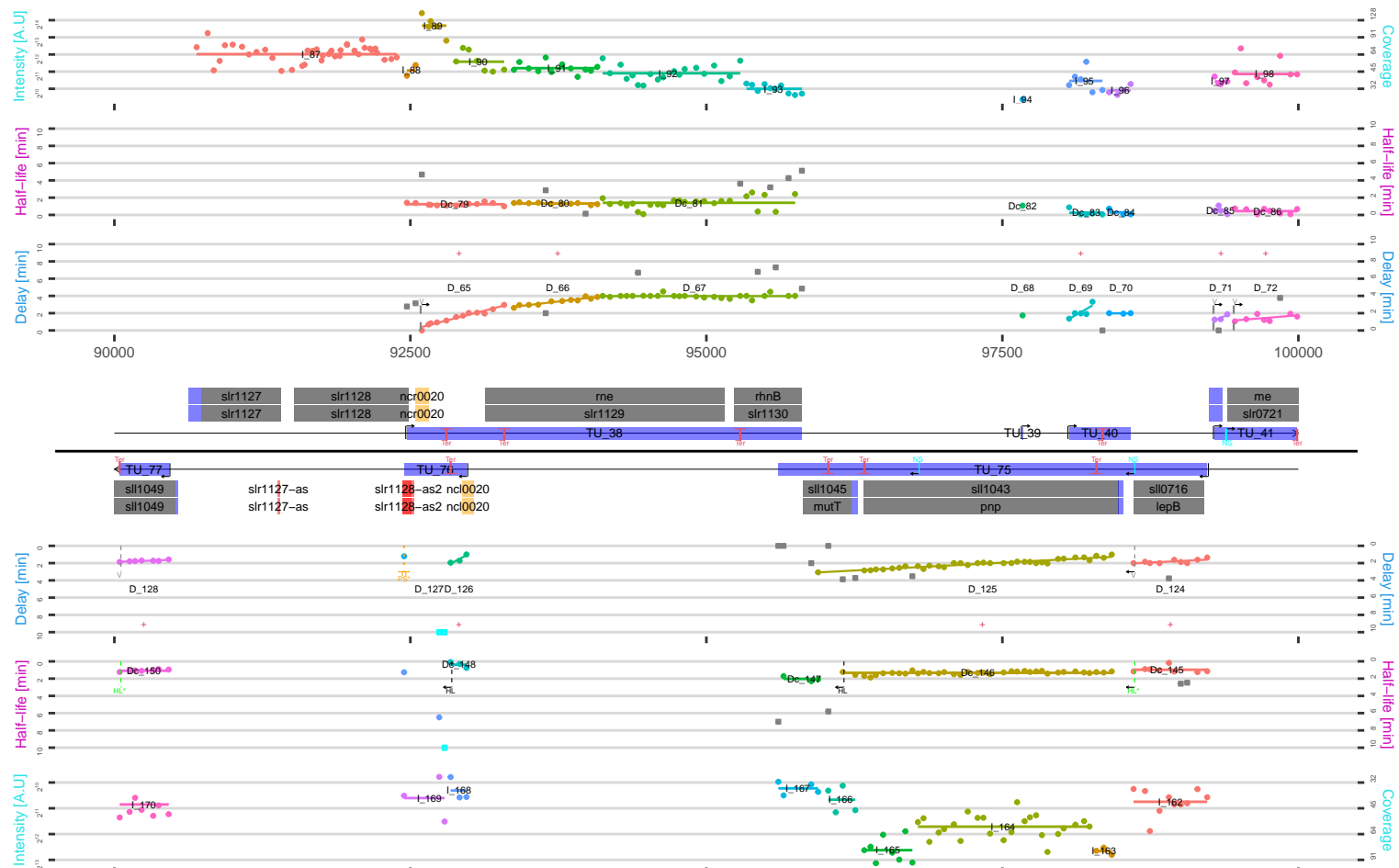


Term: termination (2), NS: new start (1), PS: pausing site (3), iTSS_L: internal starting site (0)

ID: 560-622; Term: termination (1), NS: new start (3), PS: pausing site (2), iTSS_L: internal starting site (0)



ID: 623-739; Term: termination (5), NS: new start (1), PS: pausing site (1), iTSS_L: internal starting site (3)

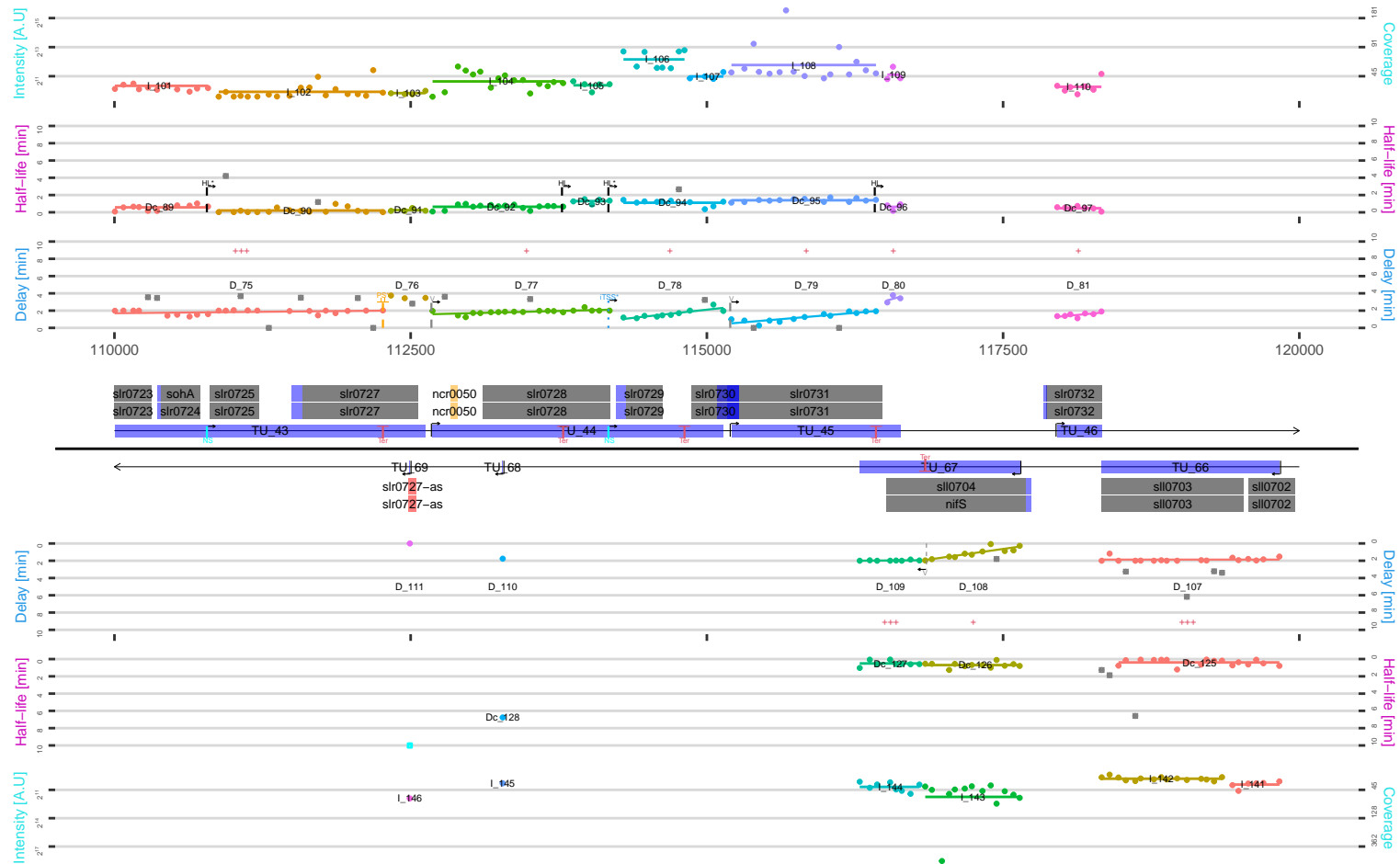


ID: 740–782; Term: termination (1), NS: new start (1), PS: pausing site (1), iTSS_L: internal starting site (1)



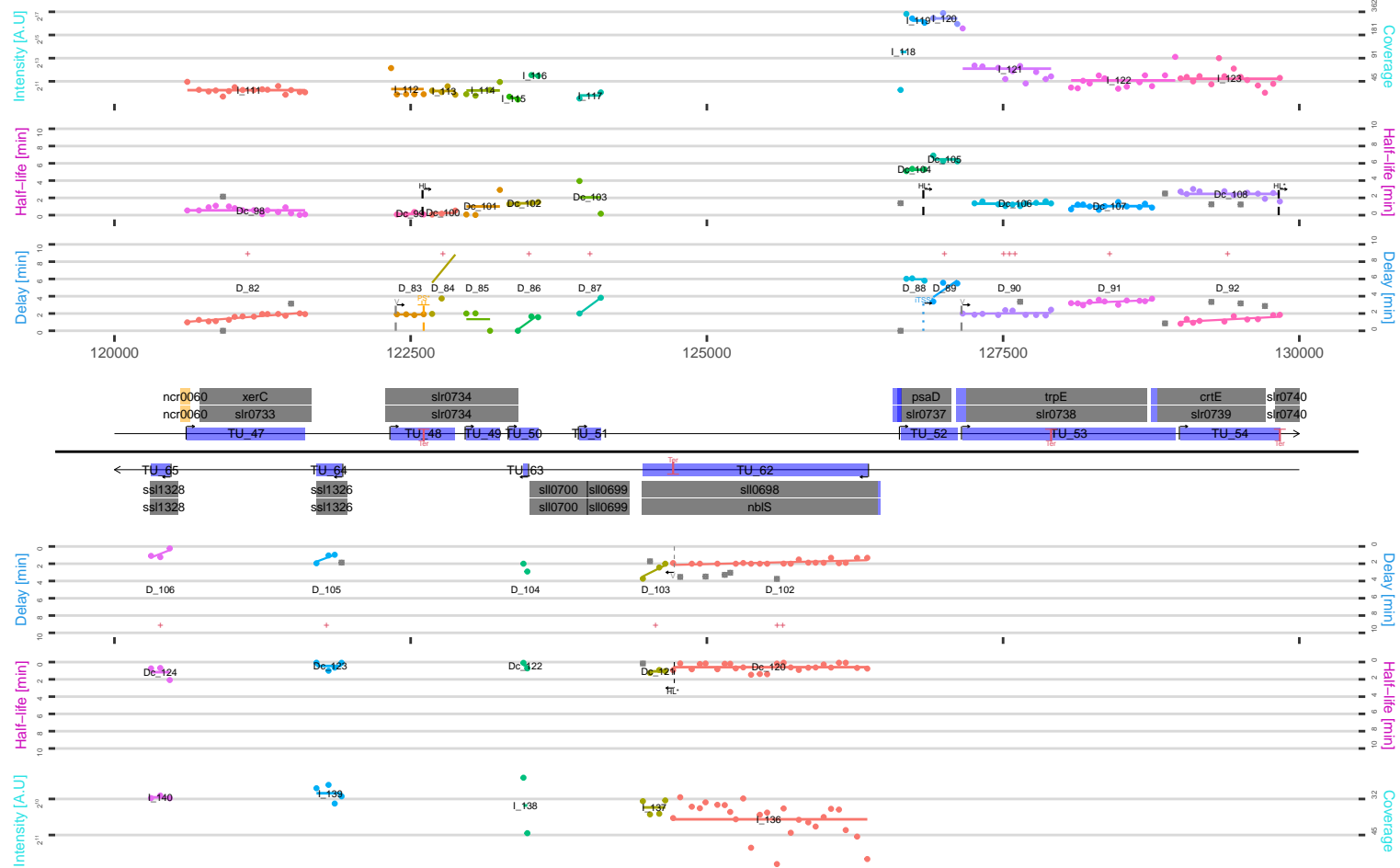
Term: termination (2), NS: new start (4), PS: pausing site (5), iTSS_L: internal starting site (2)

ID: 783–875; Term: termination (4), NS: new start (2), PS: pausing site (2), iTSS_L: internal starting site (1)



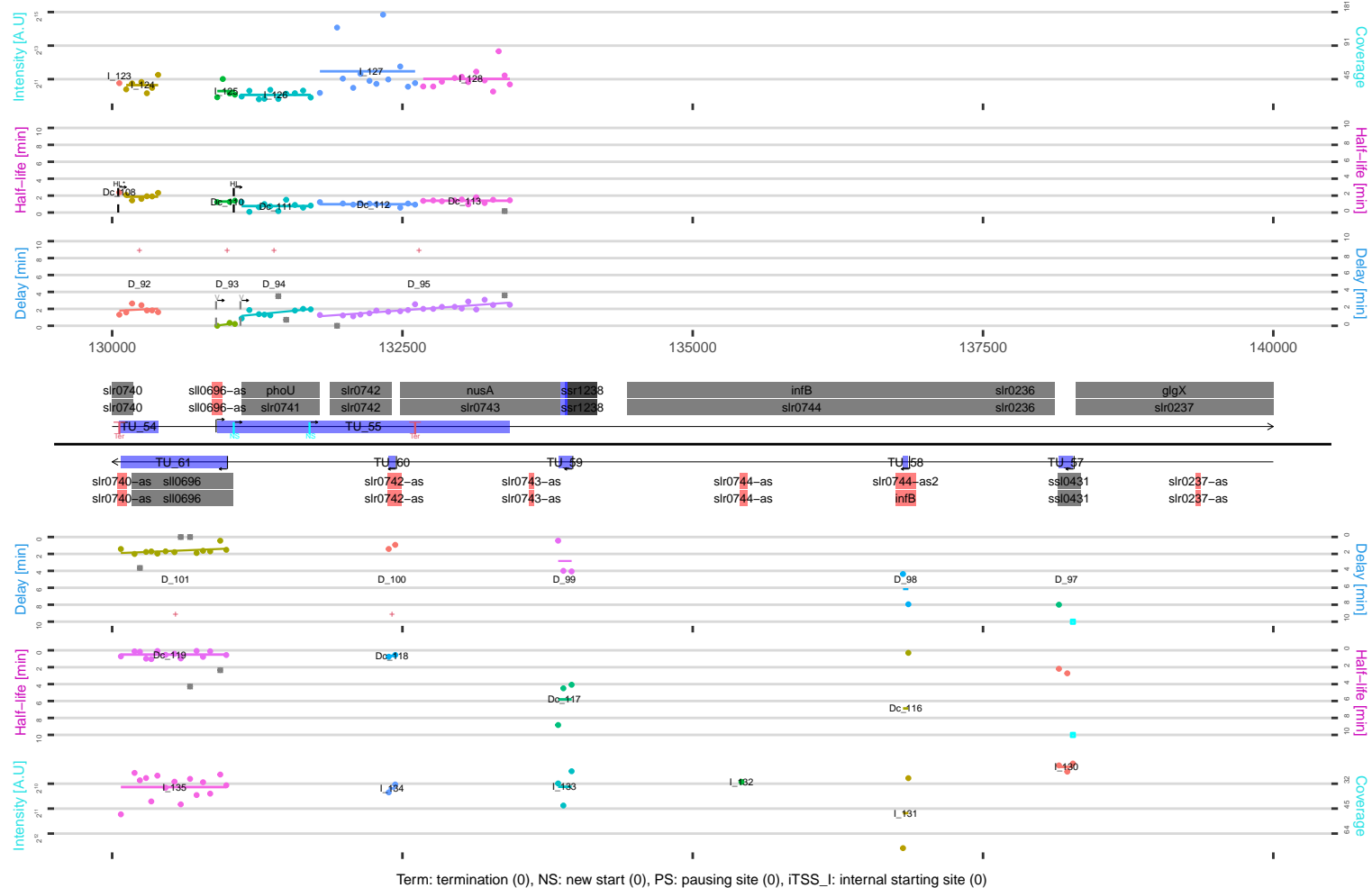
Term: termination (1), NS: new start (0), PS: pausing site (1), iTSS_L: internal starting site (0)

ID: 876-955; Term: termination (3), NS: new start (0), PS: pausing site (0), iTSS_L: internal starting site (1)



Term: termination (1), NS: new start (0), PS: pausing site (0), iTSS_L: internal starting site (1)

ID: 956-1000; Term: termination (2), NS: new start (2), PS: pausing site (1), iTSS_L: internal starting site (1)



ID: 1004–1001; FC*: significant t–test of two consecutive segments; Term: termination, NS: new start, PS: pausing site, iTSS_L: internal starting site, TI: transcription interference.

