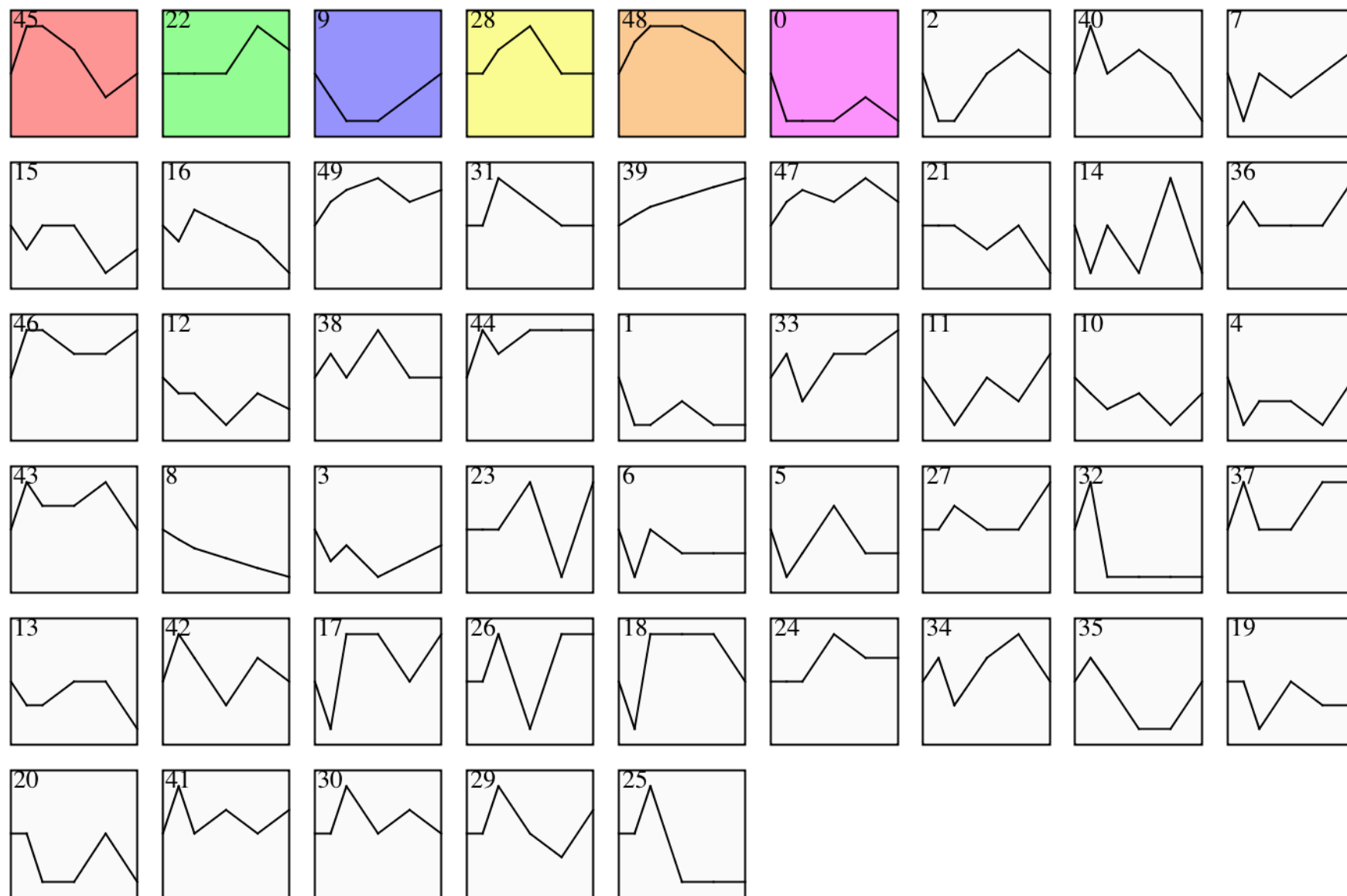


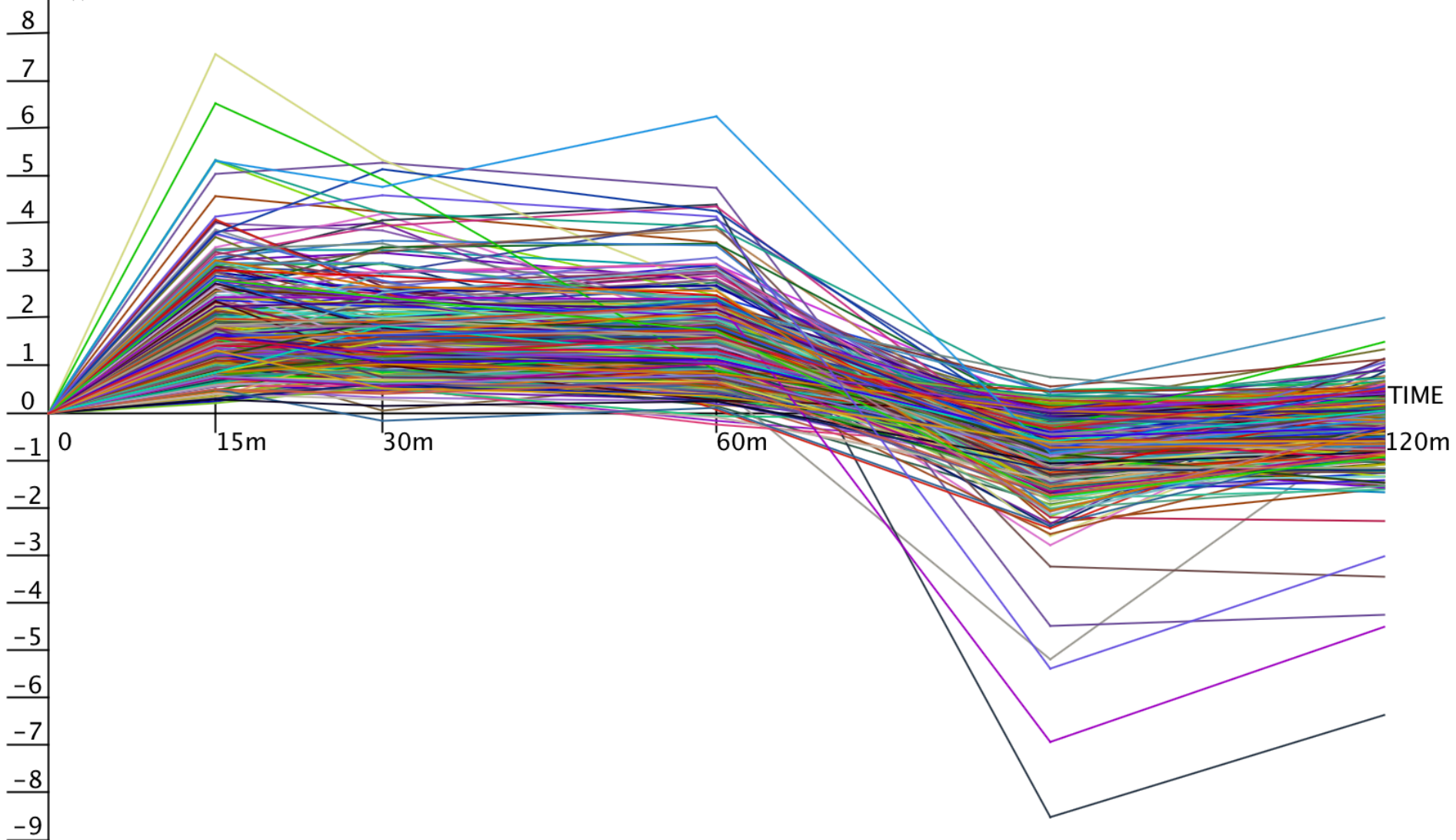
ANOVA	WT
p < 0.05	2600 (42%)
p < 0.01	1727 (28%)
p < 0.001	1015 (16.4%)
p < 0.0001	574 (9.3%)
B & H p < 0.05	302 (4.88%)
Bonferroni p < 0.05	1936 (31.3%)

Clusters ordered based on number of genes and profiles ordered by significance (default)



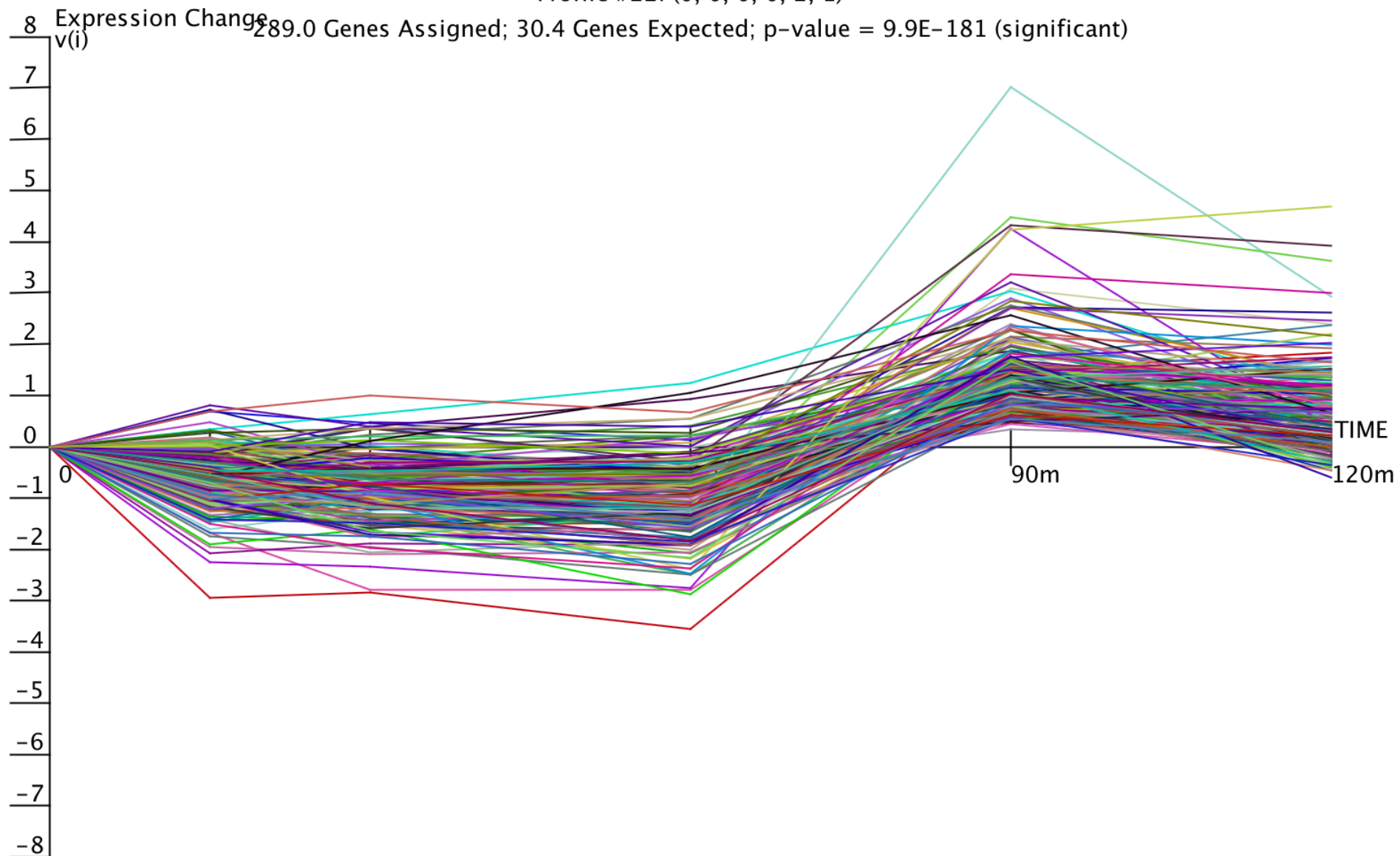
Profile #45: (0, 2, 2, 1, -1, 0)

9 Expression Change
v(i) 570.0 Genes Assigned; 51.1 Genes Expected; p-value = 0.00 (significant)



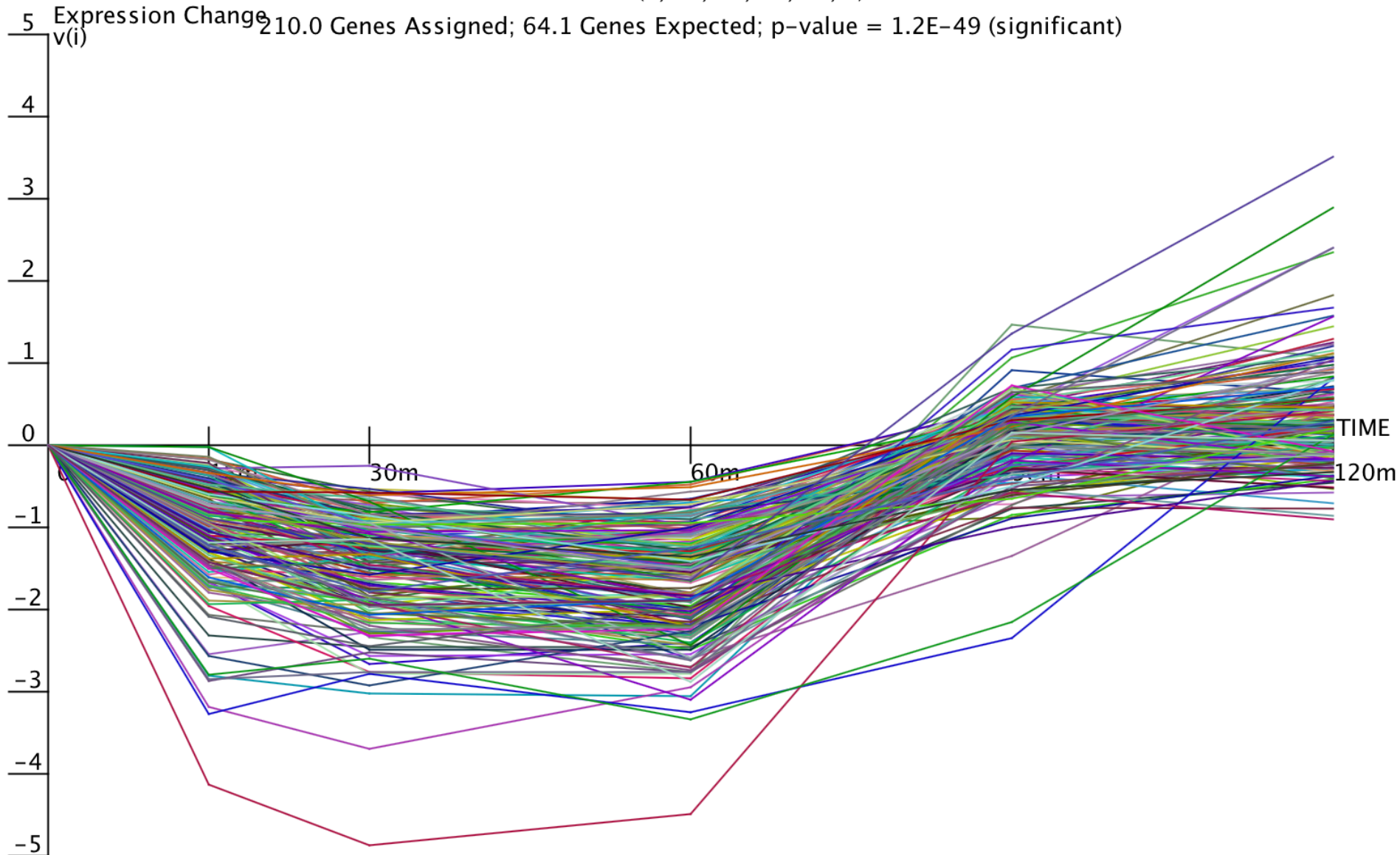
Profile #22: (0, 0, 0, 0, 2, 1)

289.0 Genes Assigned; 30.4 Genes Expected; p-value = 9.9E-181 (significant)



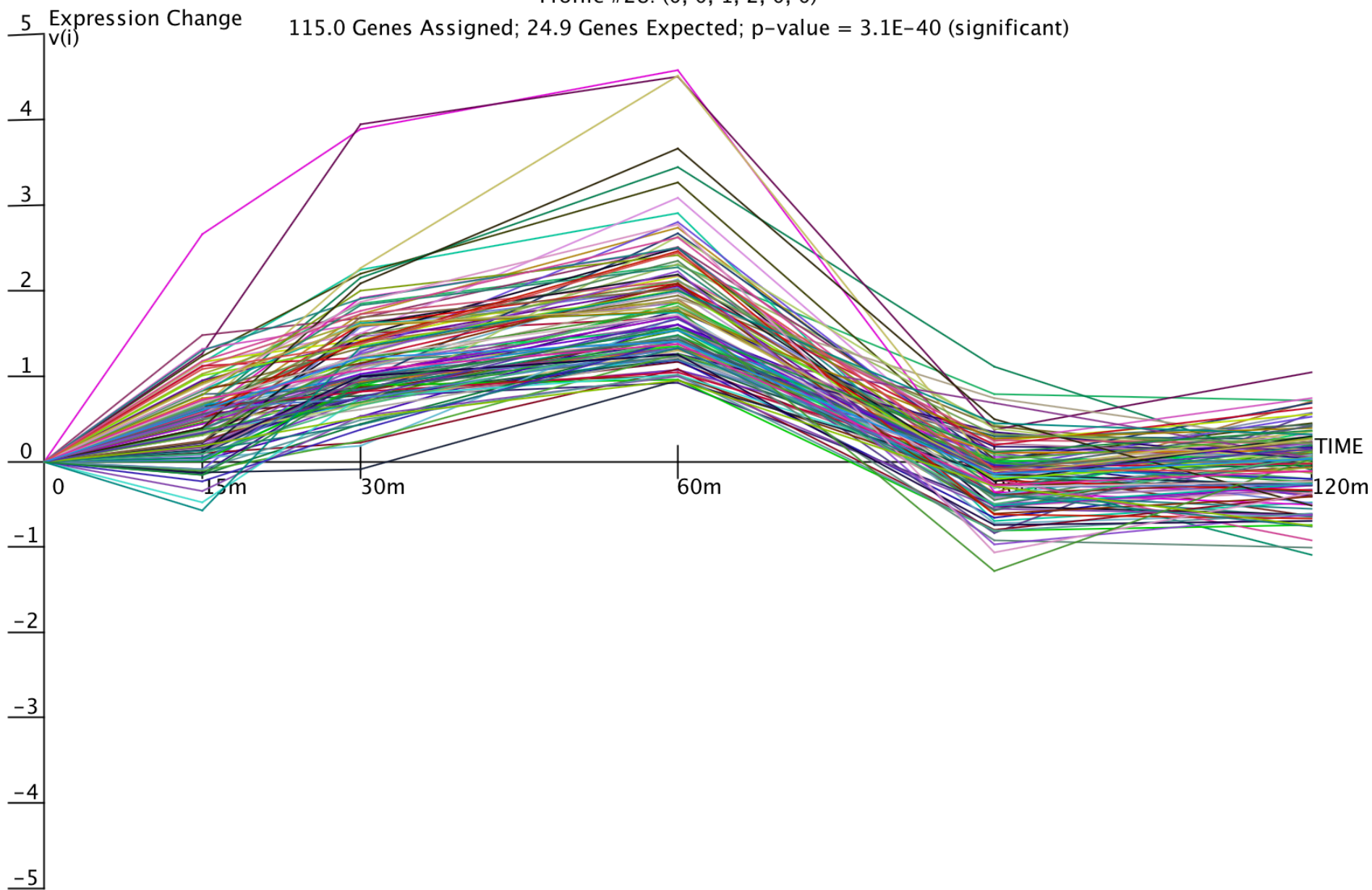
Profile #9: (0, -1, -2, -2, -1, 0)

210.0 Genes Assigned; 64.1 Genes Expected; p-value = 1.2E-49 (significant)



Profile #28: (0, 0, 1, 2, 0, 0)

115.0 Genes Assigned; 24.9 Genes Expected; p-value = 3.1E-40 (significant)



Profile #48: (0, 2, 3, 3, 2, 0)

85.0 Genes Assigned; 38.9 Genes Expected; p-value = 6.6E-11 (significant)

