

H3K27me3 – DownstreamIntergenic

Average proportion of gene region

1.00
0.75
0.50
0.25
0.00

AT1G10920
AT1G12210
AT1G12220
AT1G12280
AT1G12290
AT1G15890
AT1G17600
AT1G17610
AT1G17615
AT1G27170
AT1G27180
AT1G31540
AT1G50180
AT1G51480
AT1G52660
AT1G53350
AT1G56510
AT1G56520
AT1G56540
AT1G57650
AT1G58390
AT1G58400
AT1G58410
AT1G58602
AT1G58807
AT1G58848
AT1G59124
AT1G59218
AT1G59620
AT1G59780
AT1G61180
AT1G61190
AT1G61300
AT1G61310
AT1G62630
AT1G63350
AT1G63360
AT1G63730
AT1G63740
AT1G63750
AT1G63860
AT1G63870
AT1G63880
AT1G64070
AT1G65850
AT1G66090
AT1G69550
AT1G72840
AT1G72850
AT1G72860
AT1G72870
AT1G72890
AT1G72900
AT1G72910
AT1G72920
AT1G72930
AT1G72940
AT1G72950
AT1G16870
AT2G17050
AT2G17060
AT3G04210
AT3G04220
AT3G07040
AT3G14460
AT3G14470
AT3G15700
AT3G25510
AT3G50950
AT3G51560
AT3G51570
AT4G10780
AT4G11170
AT4G12010
AT4G12020
AT4G14370
AT4G16860
AT4G16890
AT4G16900
AT4G16920
AT4G16930
AT4G16940
AT4G16950
AT4G16960
AT4G16990
AT4G19500
AT4G19510
AT4G19520
AT4G19530
AT4G23510
AT4G26090
AT4G27190
AT4G27220
AT4G33300
AT4G36140
AT4G36150
AT5G04720
AT5G05400
AT5G11250
AT5G17680
AT5G17880
AT5G17890
AT5G17970
AT5G18350
AT5G18360
AT5G18370
AT5G22690
AT5G40060
AT5G40090
AT5G40100
AT5G40910
AT5G41540
AT5G41550
AT5G41740
AT5G41750
AT5G43470
AT5G43730
AT5G43740
AT5G44510
AT5G44870
AT5G45000
AT5G45050
AT5G45060
AT5G45200
AT5G45210
AT5G45220
AT5G45230
AT5G45240
AT5G45250
AT5G45260
AT5G45440
AT5G45490
AT5G46260
AT5G46270
AT5G46450
AT5G46470
AT5G46490
AT5G46510
AT5G46520
AT5G47250
AT5G47260
AT5G47280
AT5G48620
AT5G48770
AT5G48780
AT5G49140
AT5G51630
AT5G58120
AT5G63020
AT5G66900
AT5G66910

Expression

- Intermediate Expression
- Low Expression
- No Expression