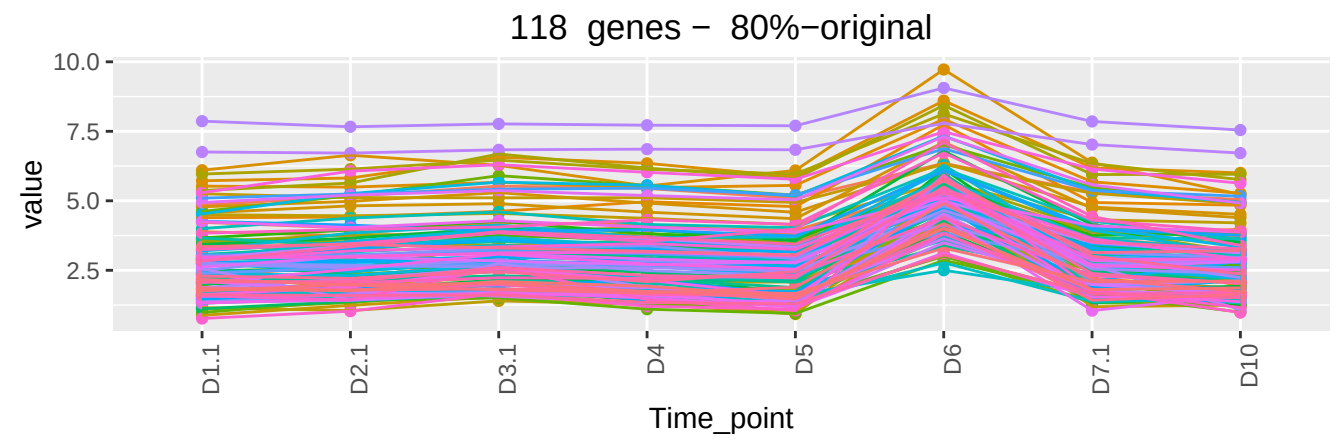
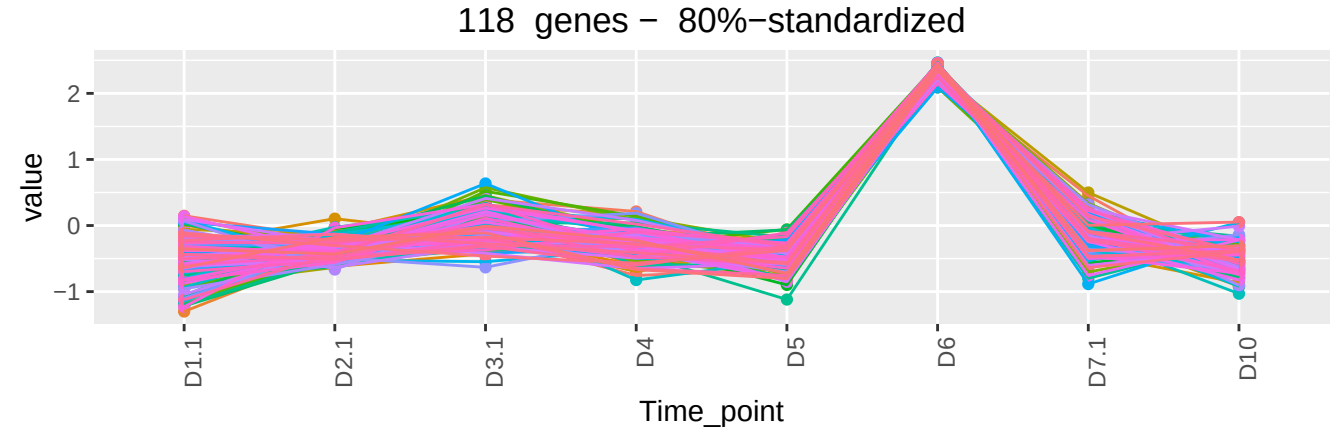
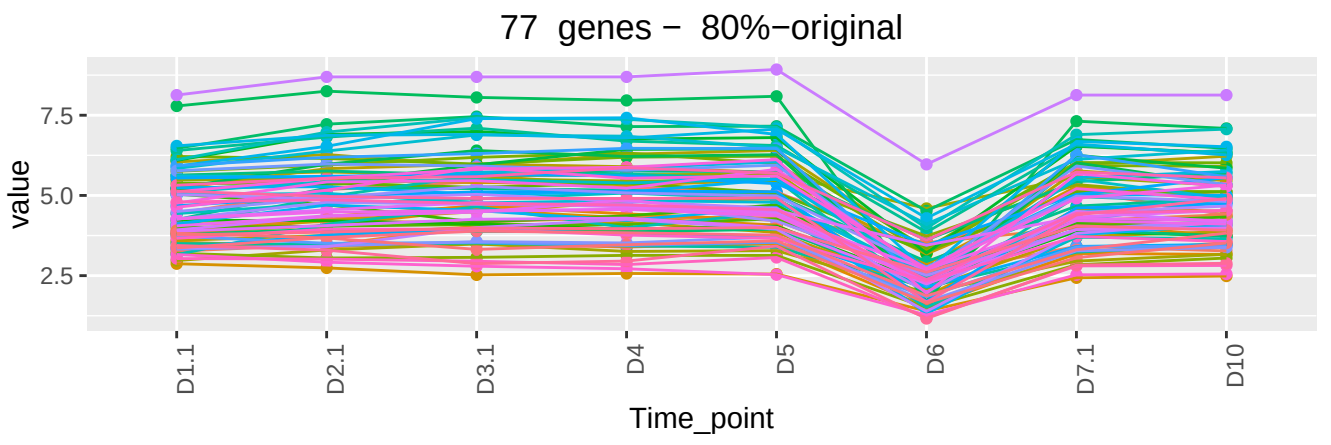

118 genes − 80%−original
10.0
7.5
5.0
2.5
value
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


118 genes − 80%−standardized
2
1
0
−1
value
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


77 genes − 80%−original
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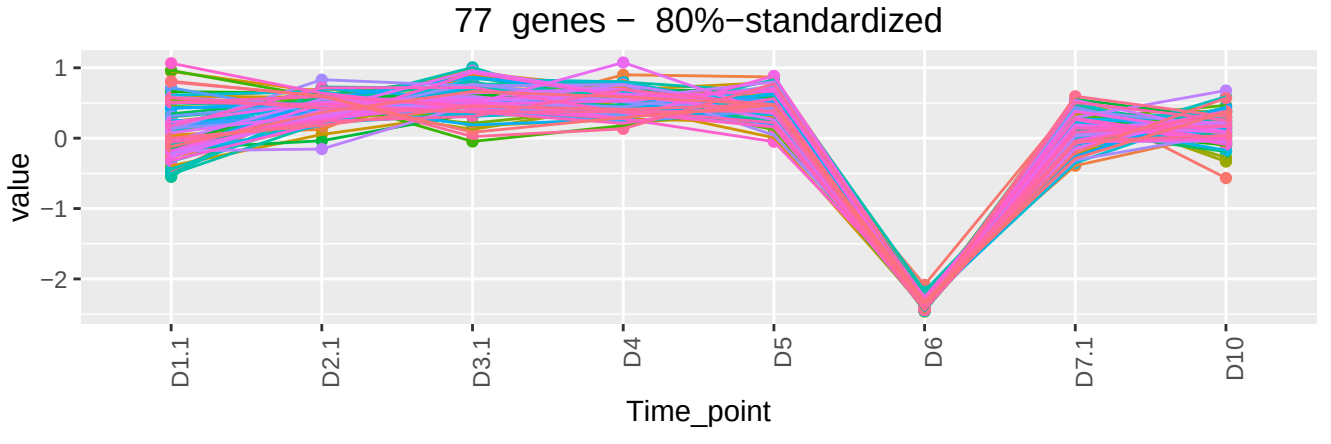
5.0
2.5
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


77 genes − 80%−standardized
value
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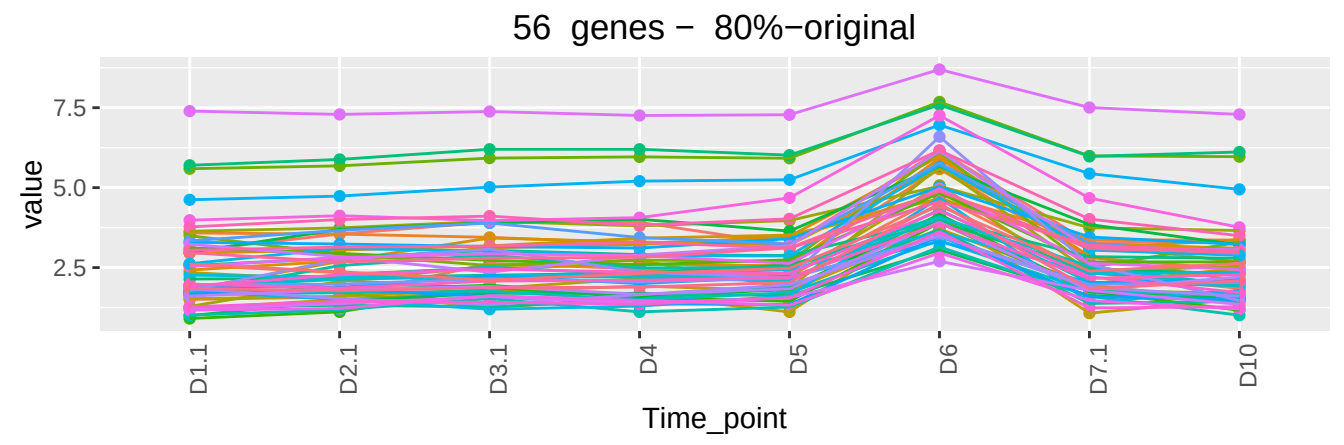
0
−1
−2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


56 genes − 80%−original
value
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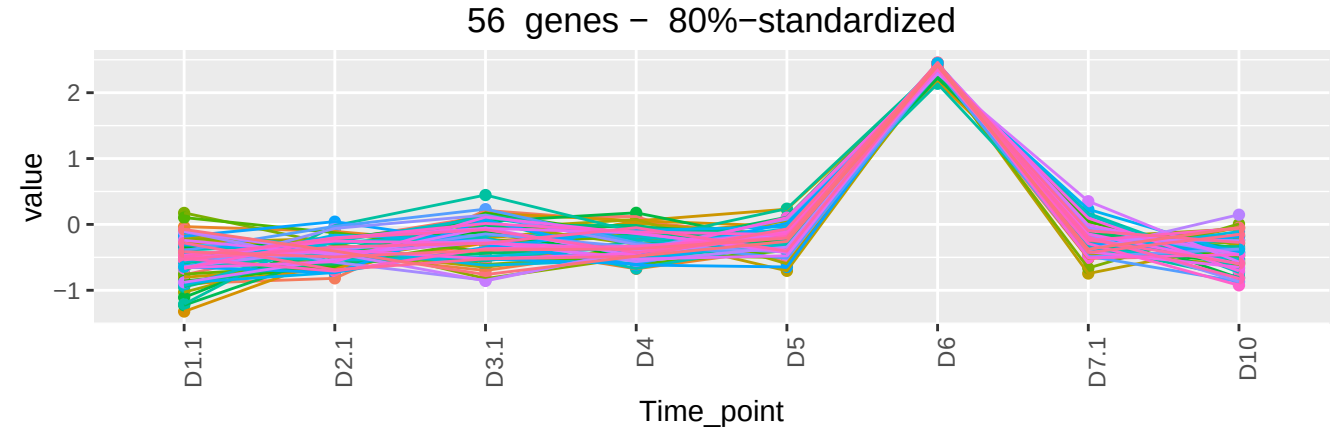
2.5
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


56 genes − 80%−standardized
value
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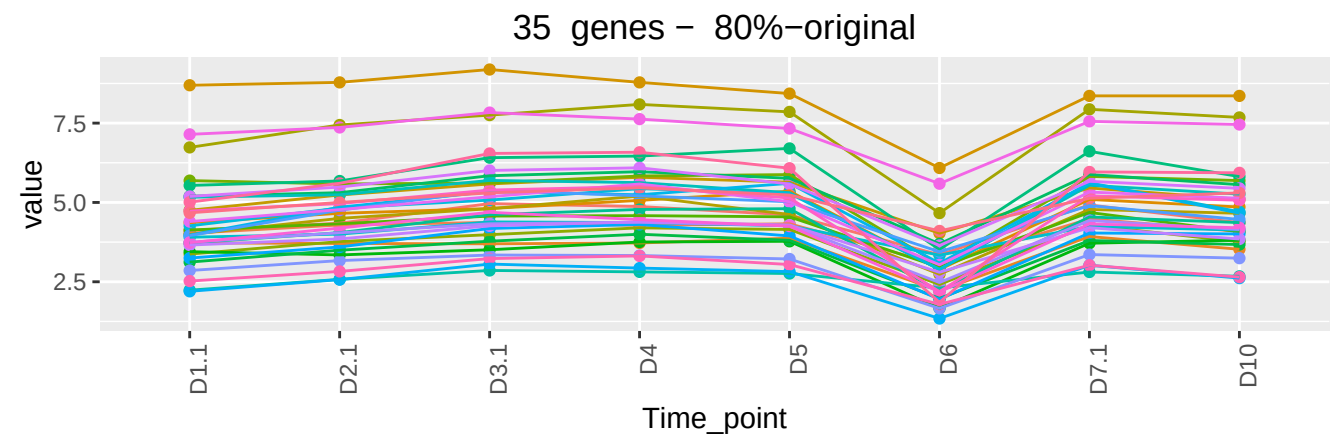
−1
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


35 genes − 80%−original
value
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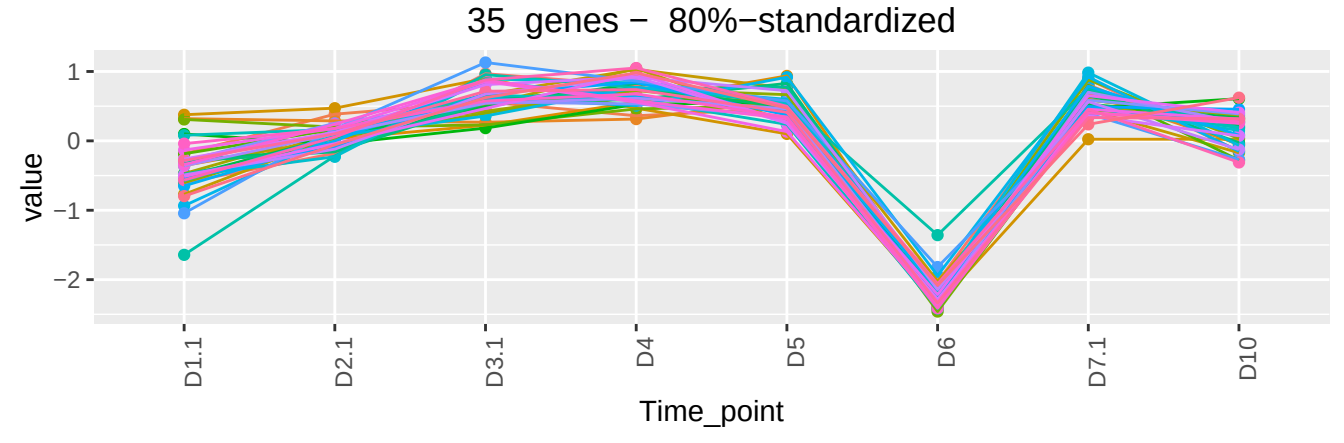
2.5
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


35 genes − 80%−standardized
value
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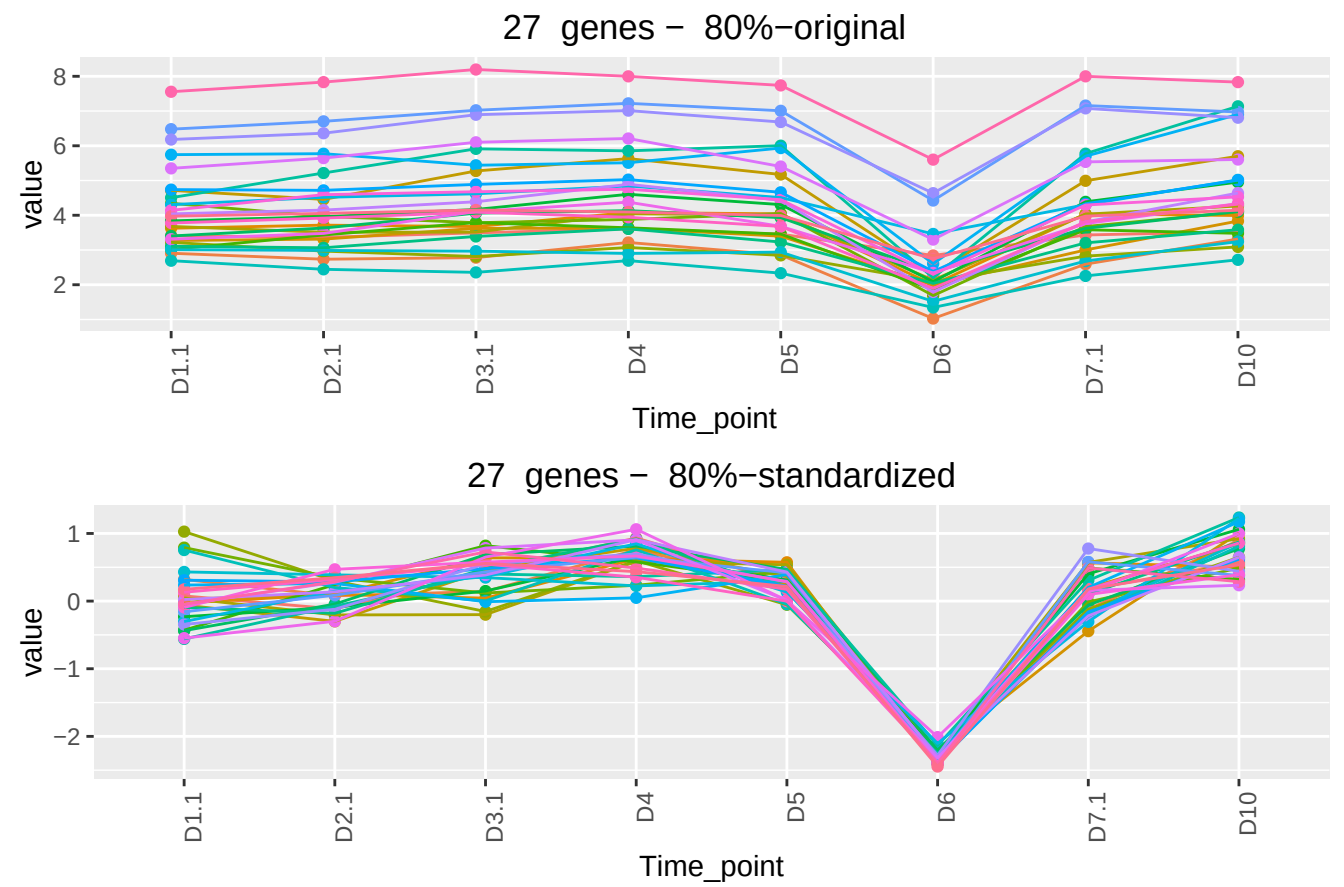
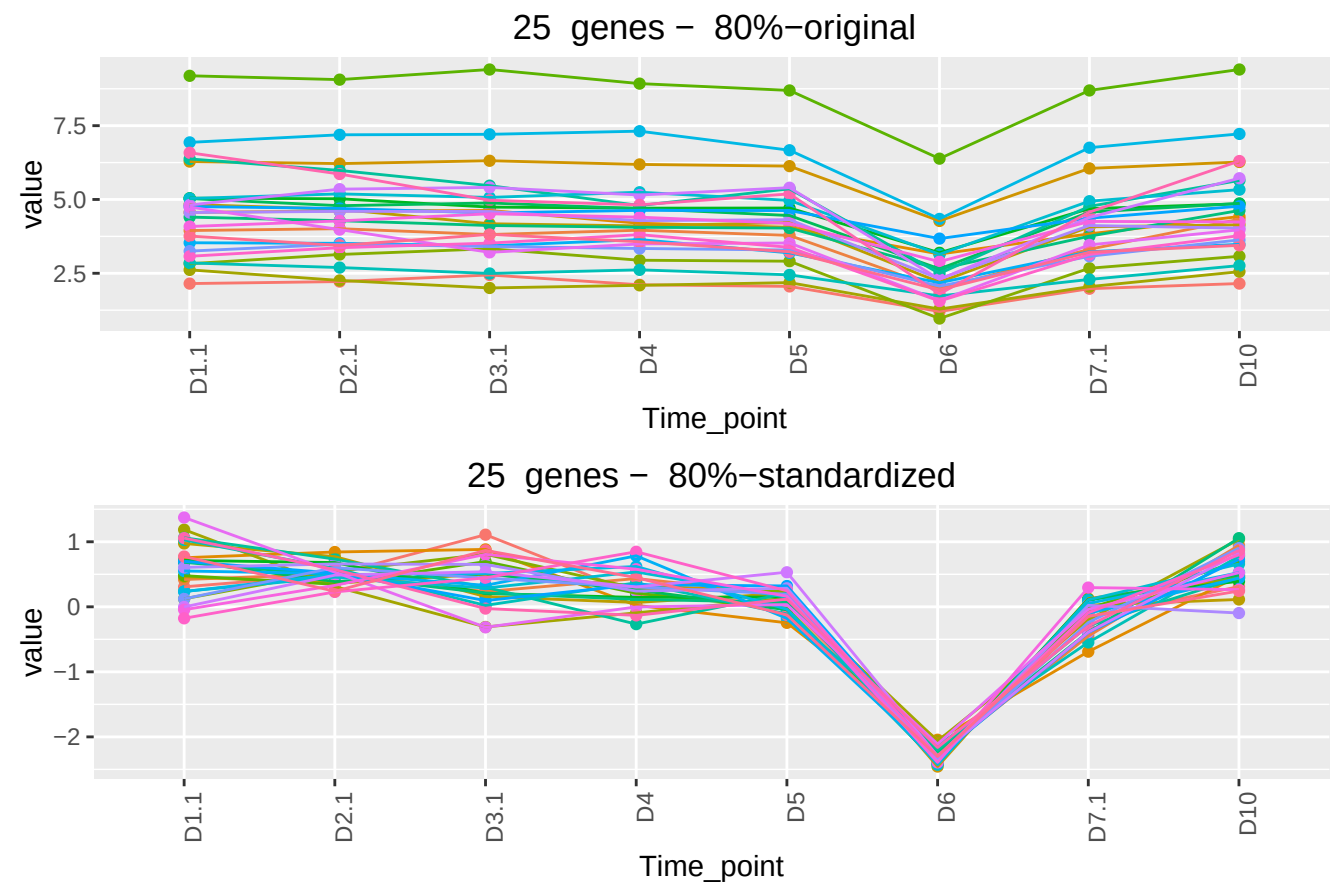
−1
−2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


27 genes − 80%−original
value
8
6
4
2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point
27 genes − 80%−standardized
value
1
0
−1
−2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point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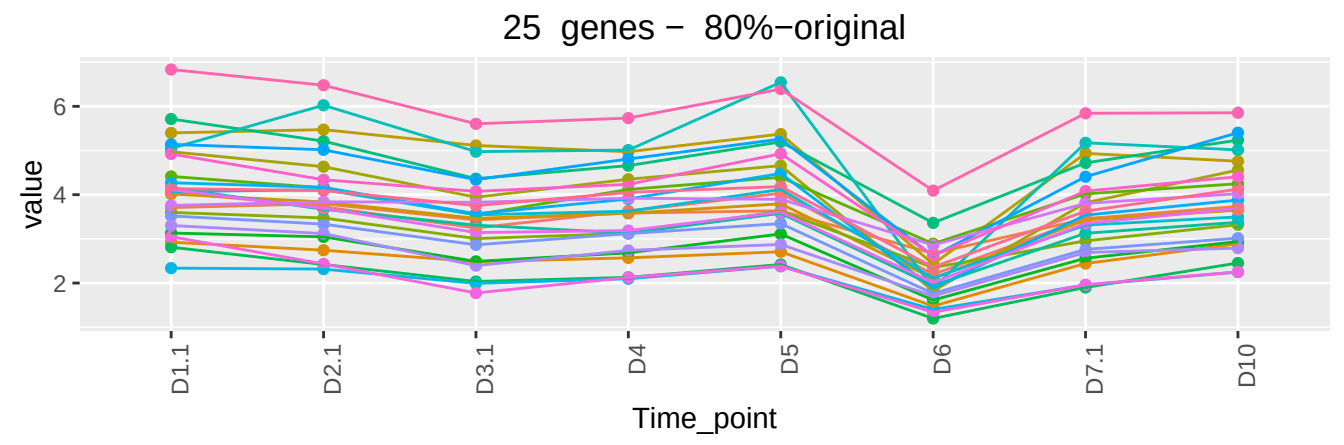
value
7.5
5.0
2.5
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point
25 genes − 80%−standardized
value
1
0
−1
−2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


25 genes − 80%−original
value
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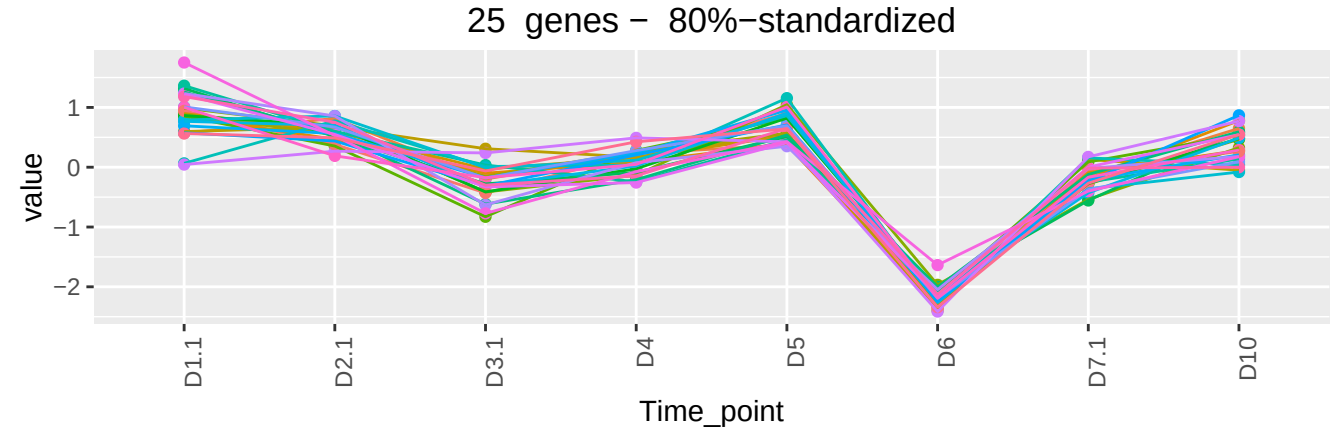
4
2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


25 genes − 80%−standardized
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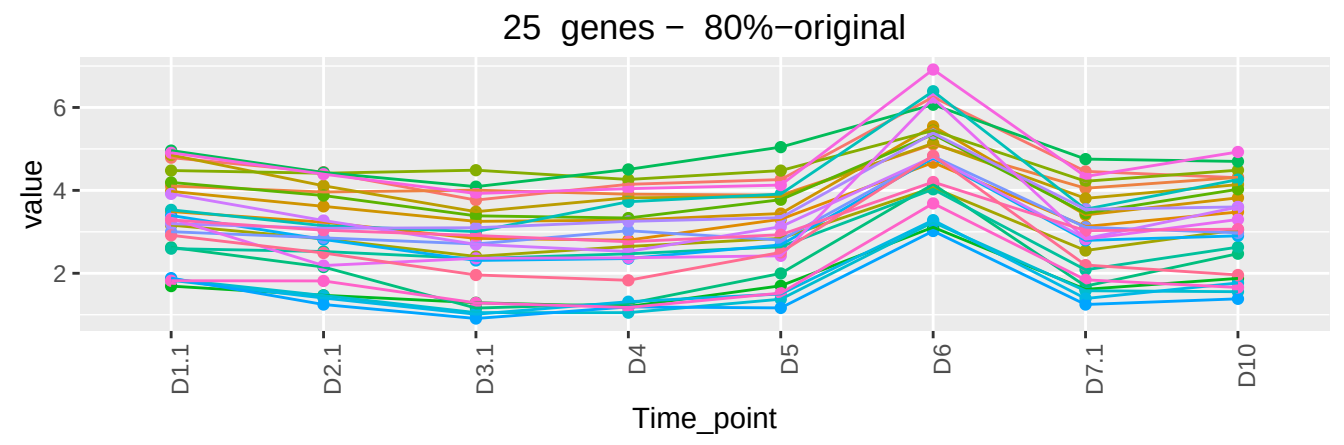
−2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


25 genes − 80%−original
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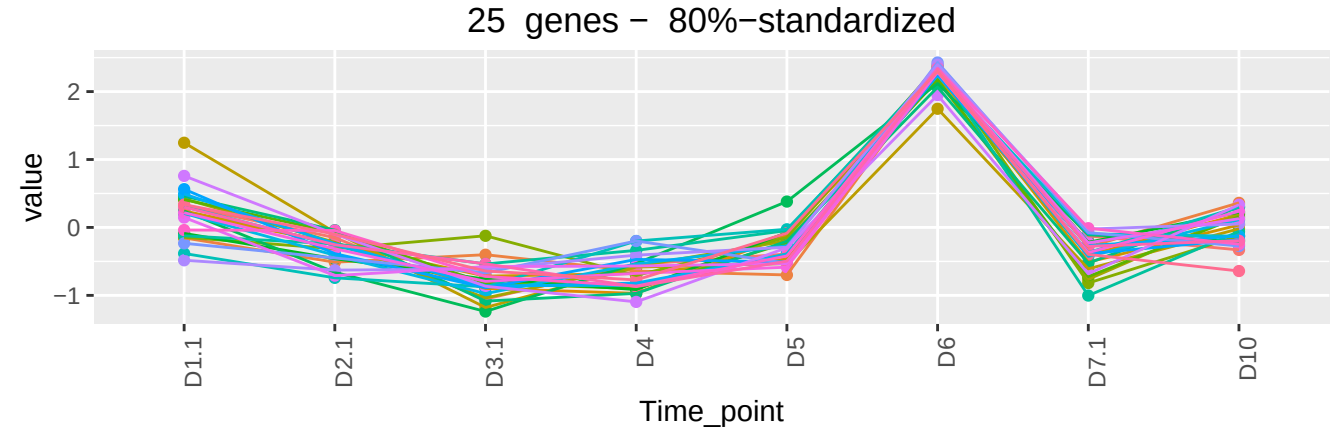
D3.1
D4
D5
D6
D7.1
D10
Time_point


25 genes − 80%−standardized
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−1
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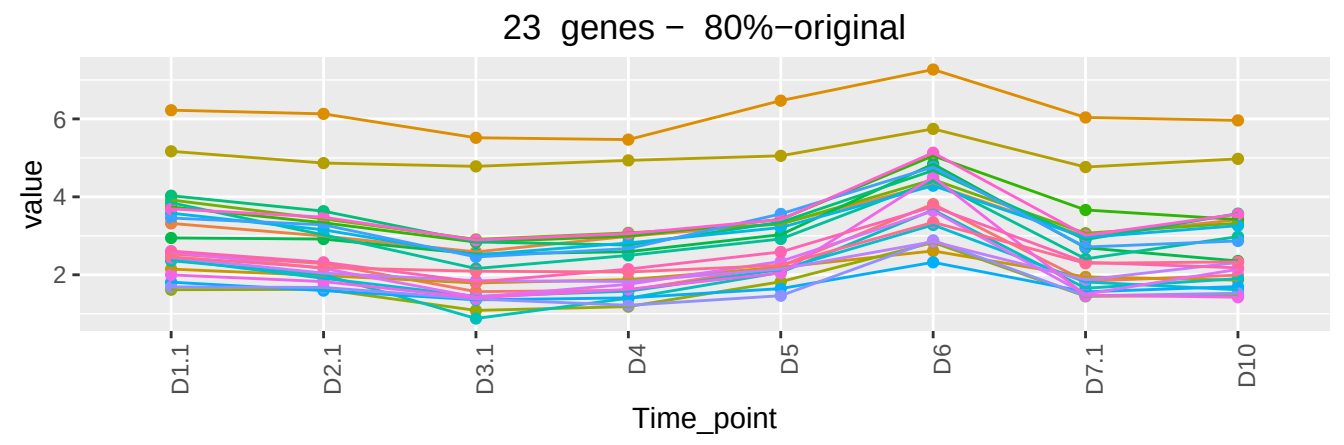
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


23 genes − 80%−original
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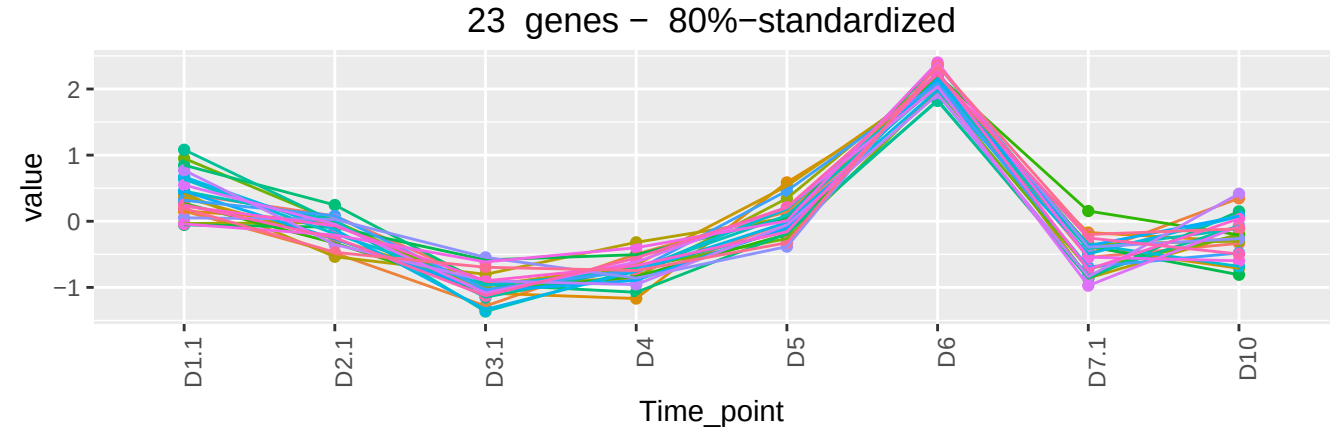
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


23 genes − 80%−standardized
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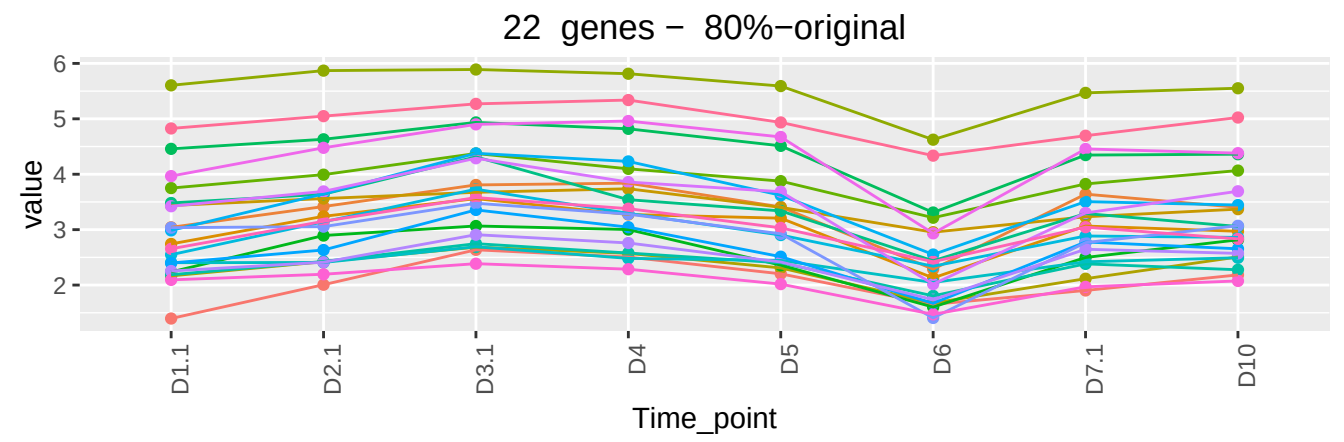
−1
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


22 genes − 80%−original
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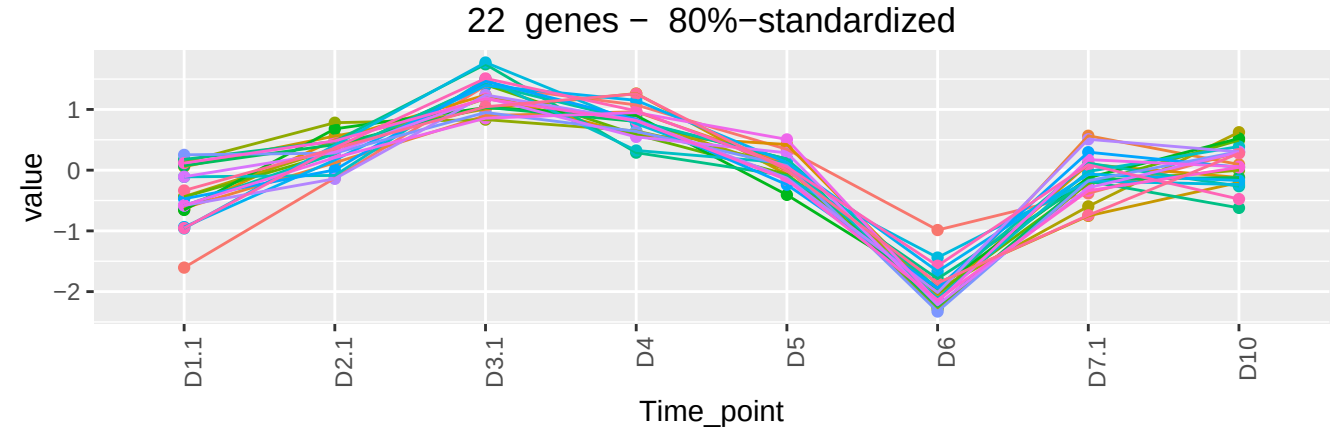
2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


22 genes − 80%−standardized
value
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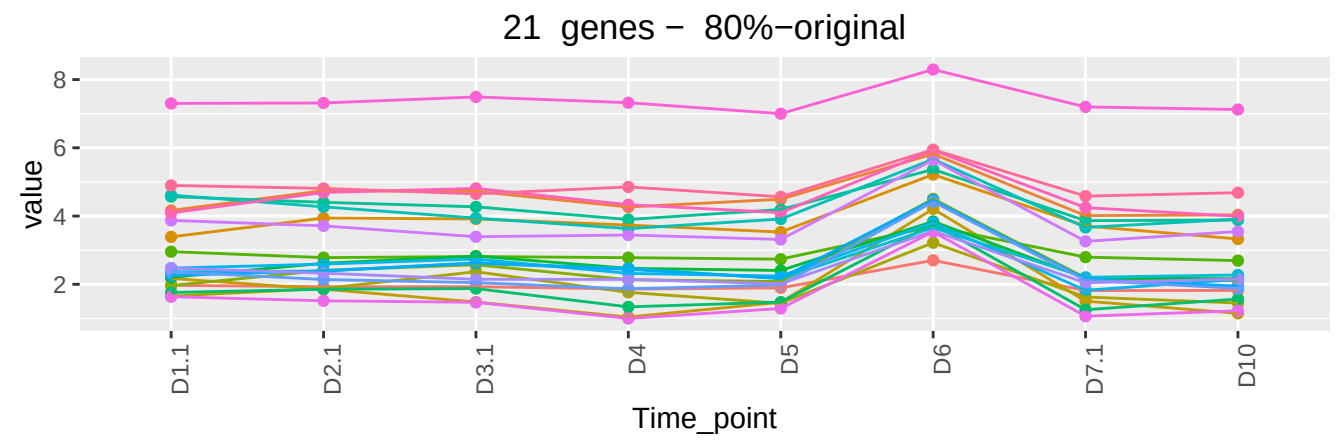
−2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


21 genes − 80%−original
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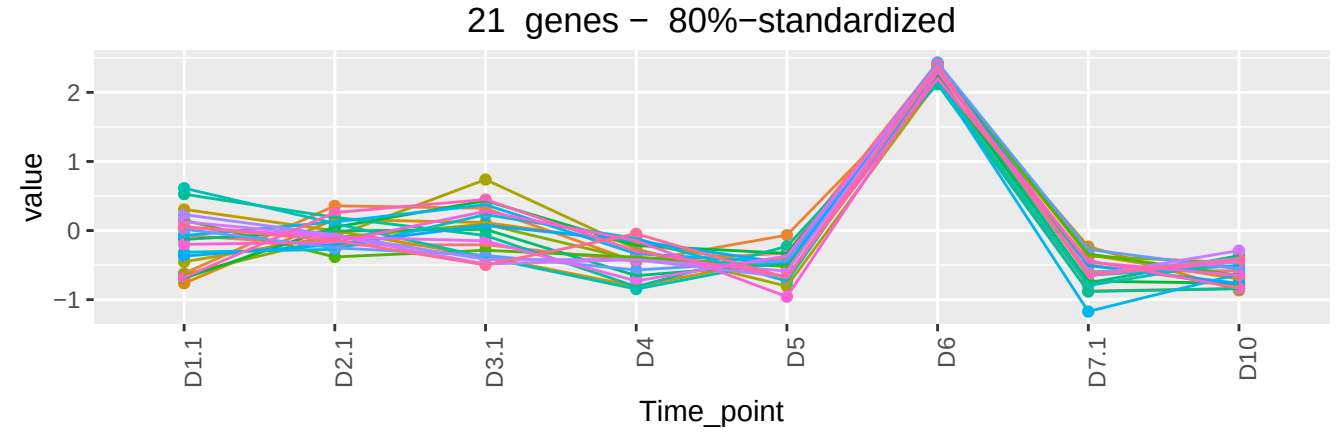
4
2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point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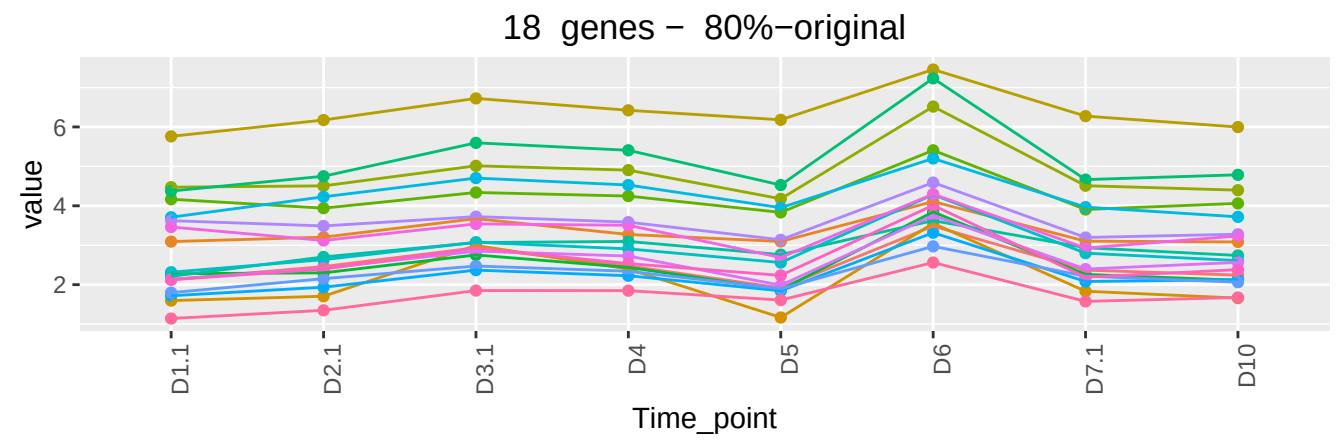
−1
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point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D1.1
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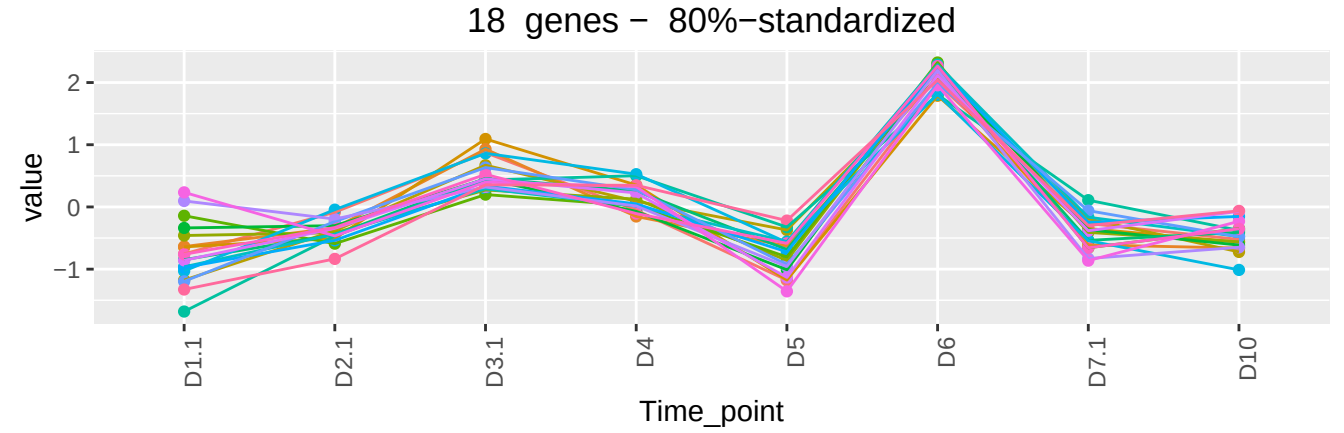
D3.1
D4
D5
D6
D7.1
D10
Time_point


18 genes − 80%−standardized
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−1
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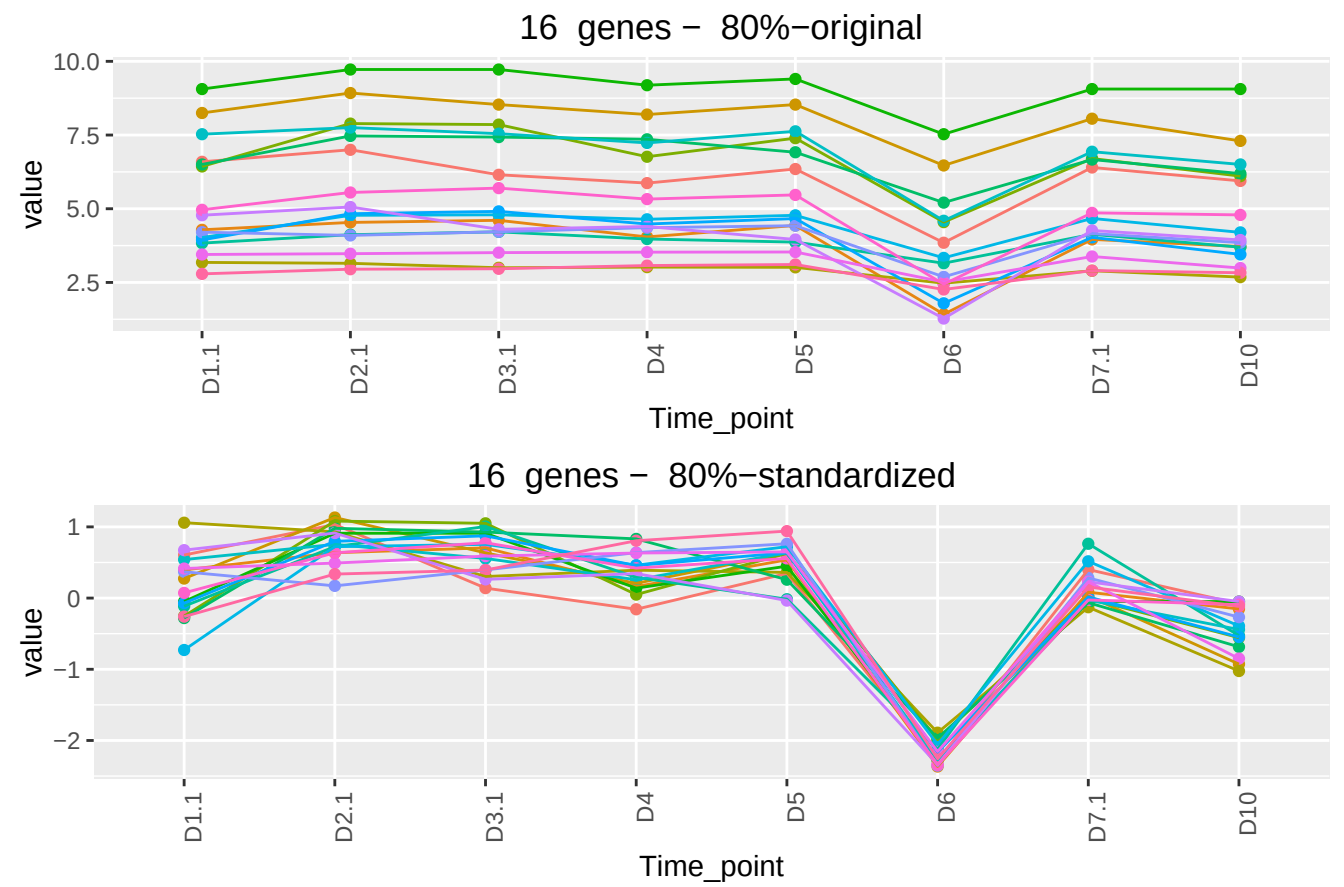
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


16 genes − 80%−original
10.0
7.5
5.0
2.5
value
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point
16 genes − 80%−standardized
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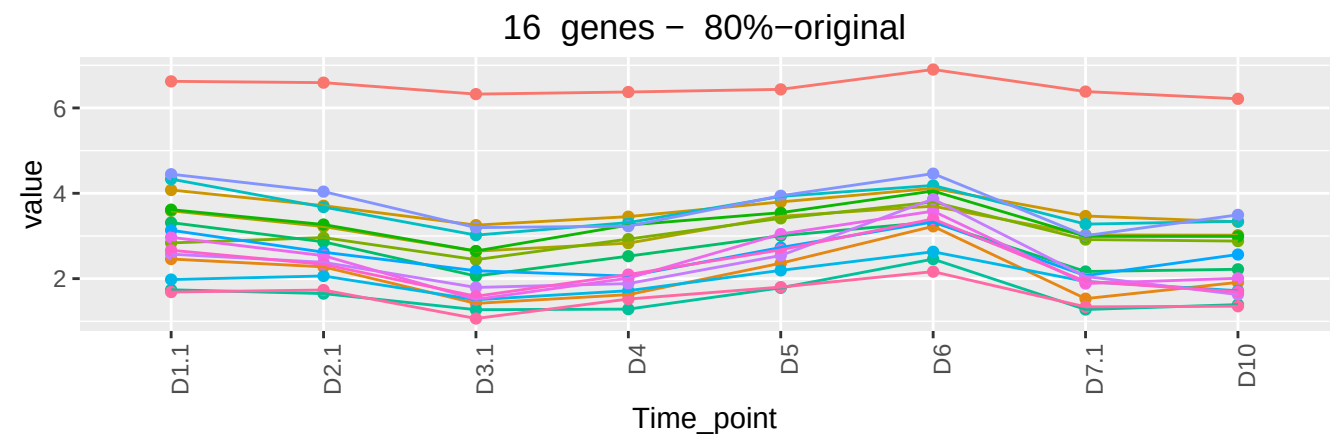
−1
−2
value
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point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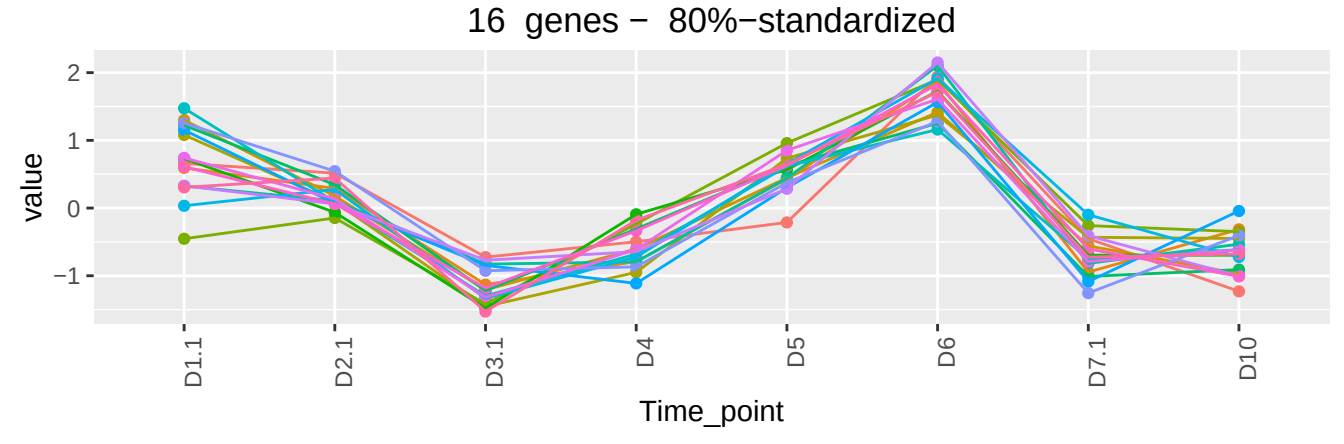
Time_point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D3.1
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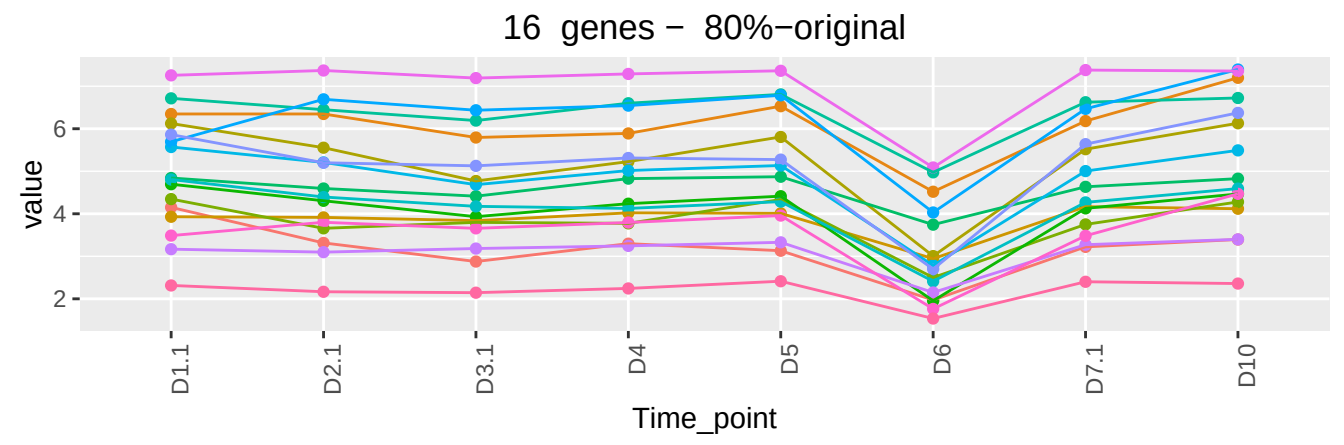
Time_point


16 genes − 80%−original
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D3.1
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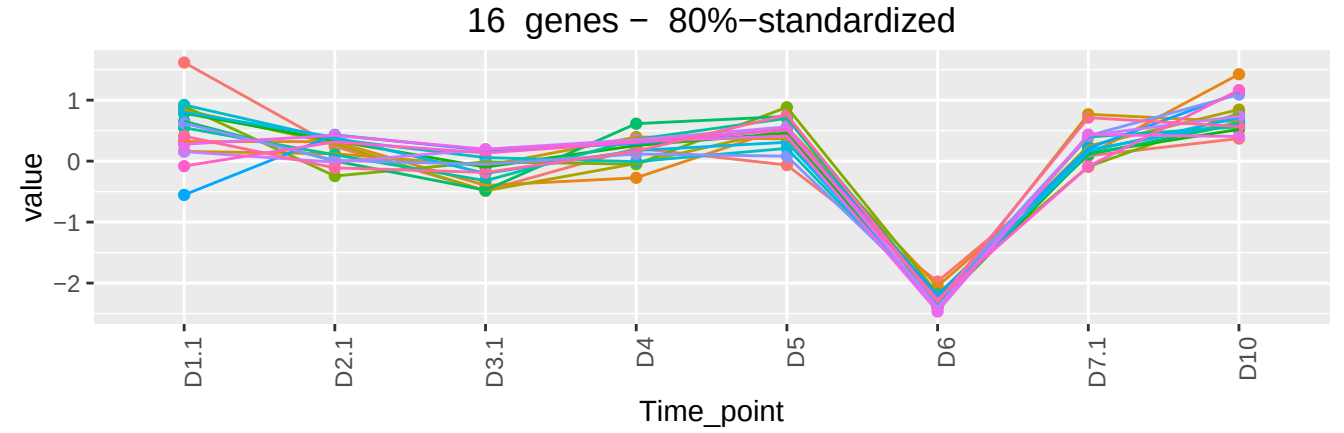
D7.1
D10
Time_point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−2
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D3.1
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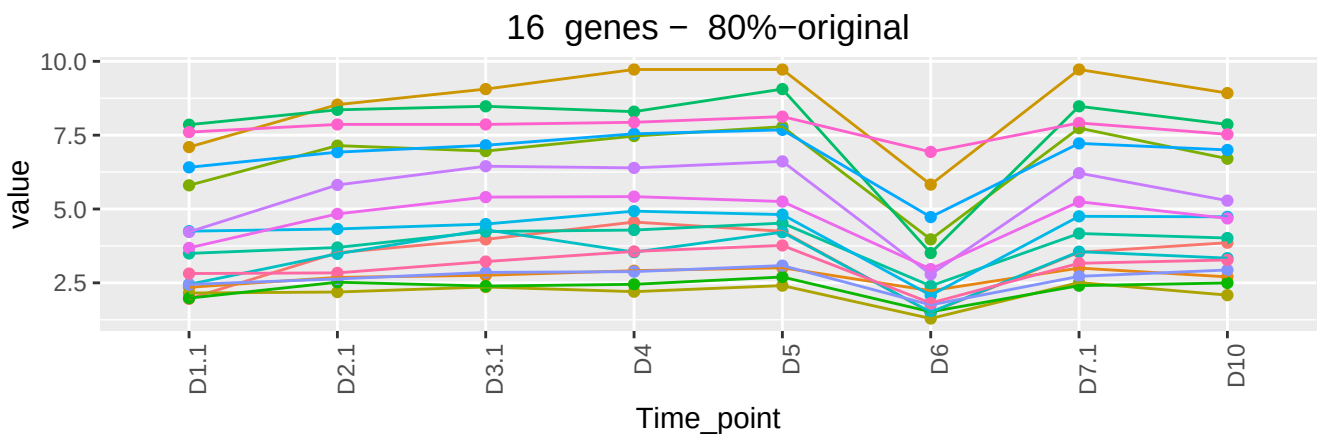
Time_point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10.0
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value
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D2.1
D3.1
D4
D5
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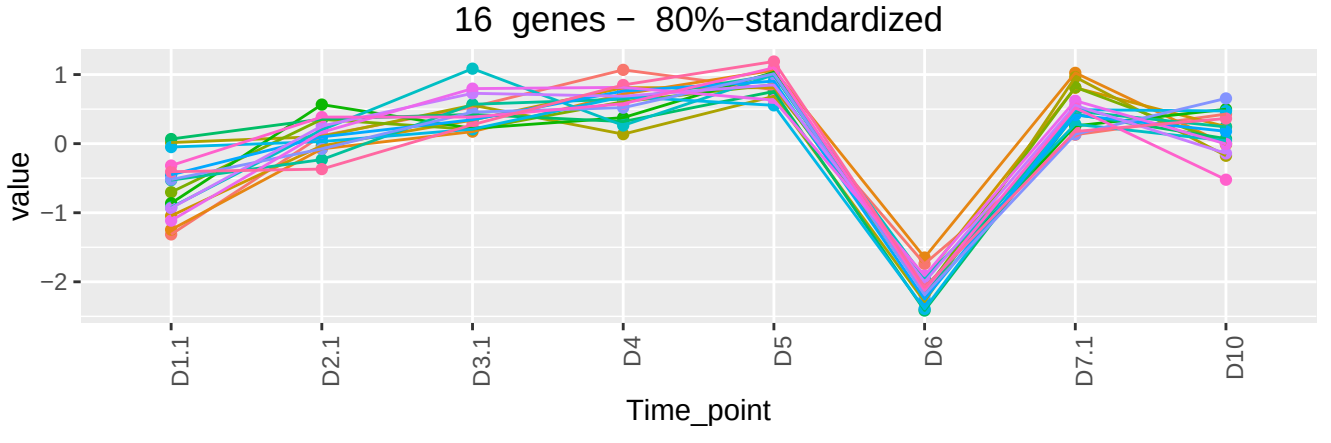
D10
Time_point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value
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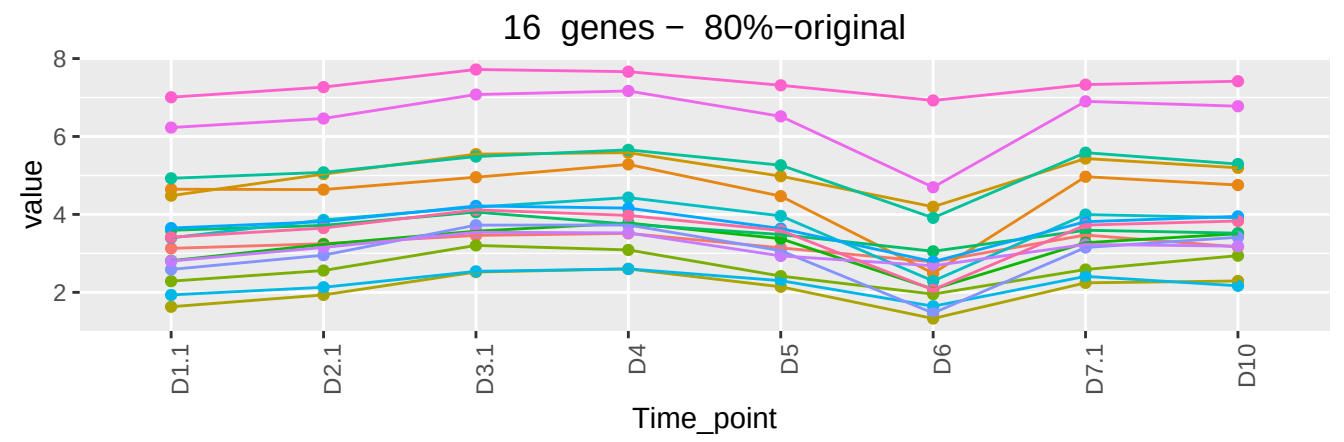
D10
Time_point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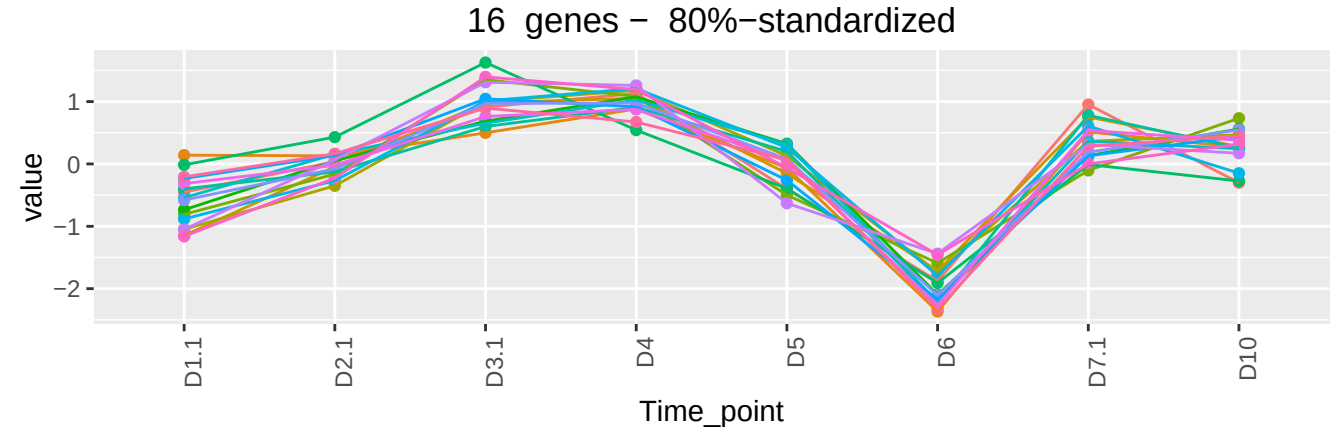
D5
D6
D7.1
D10
Time_point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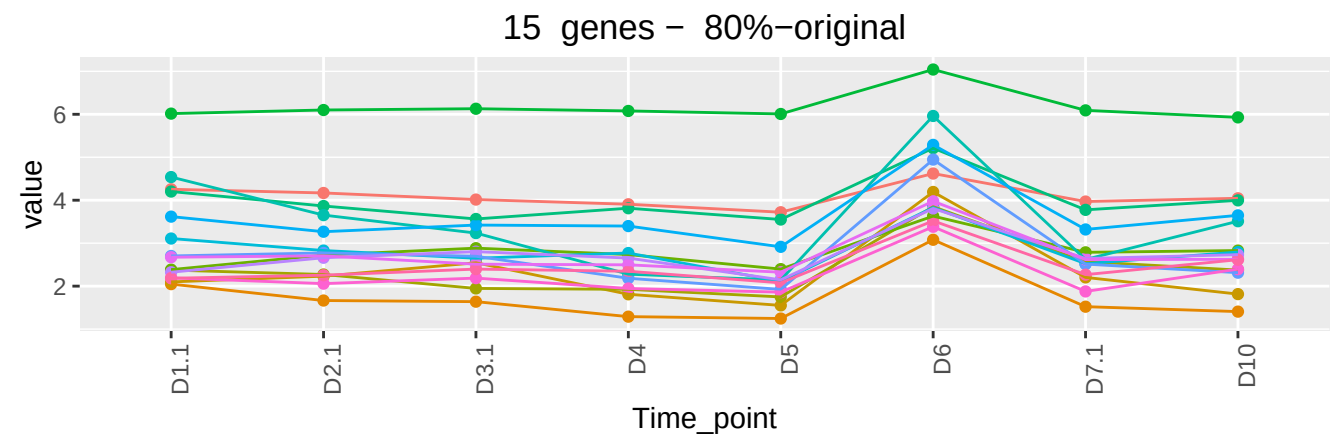
Time_point


15 genes − 80%−original
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D1.1
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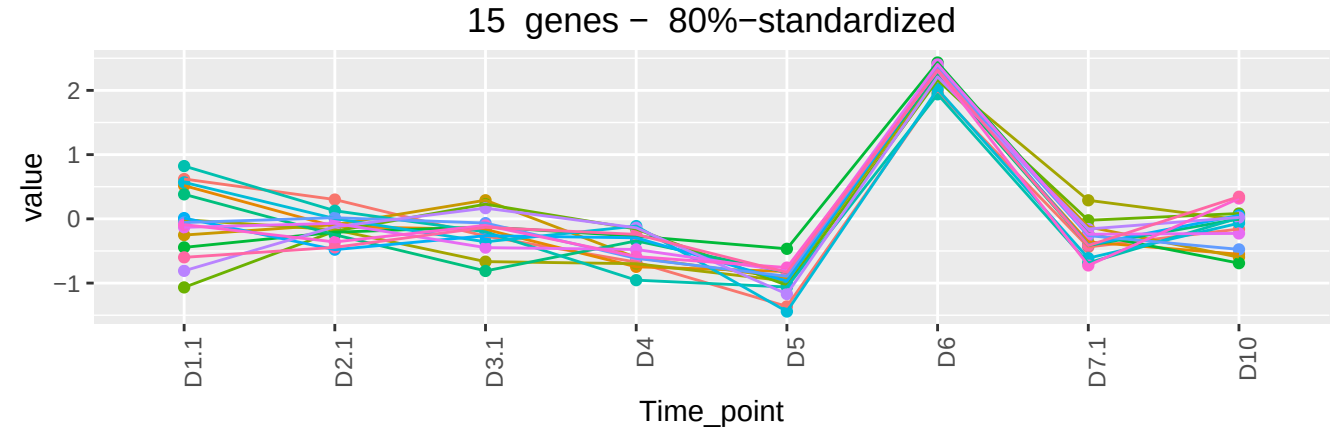
D7.1
D10
Time_point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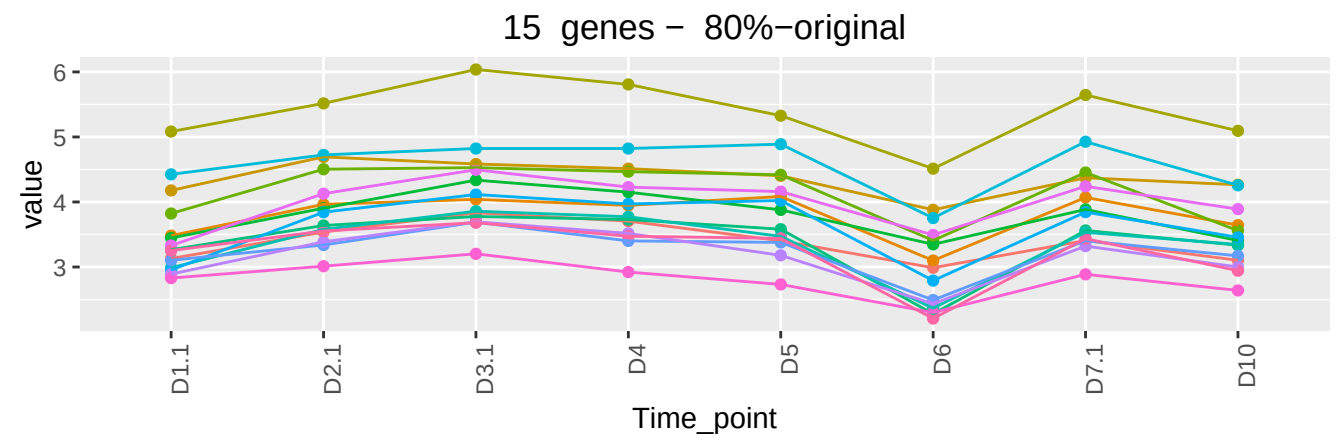
D7.1
D10
Time_point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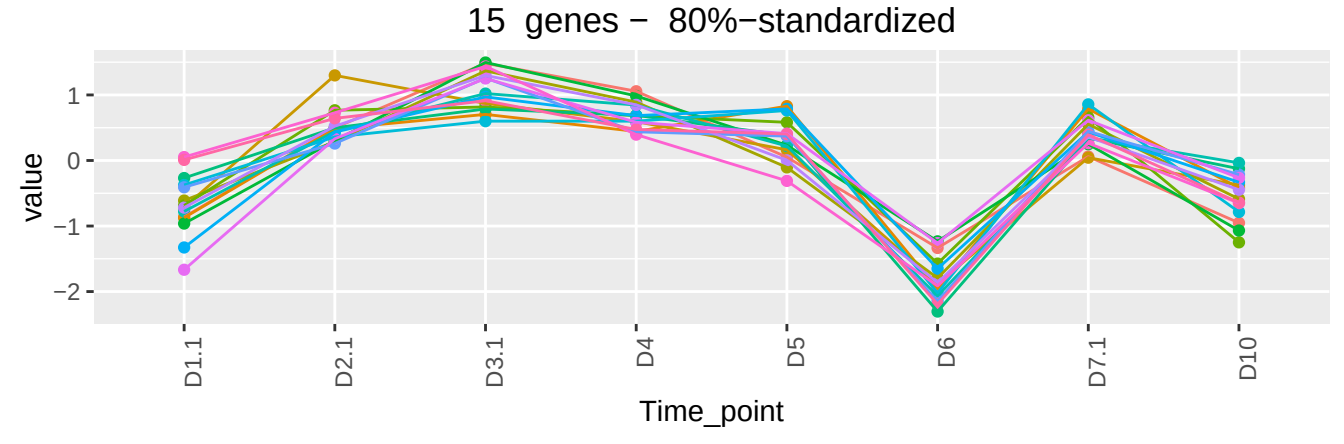
D5
D6
D7.1
D10
Time_point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−2
D1.1
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D3.1
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D5
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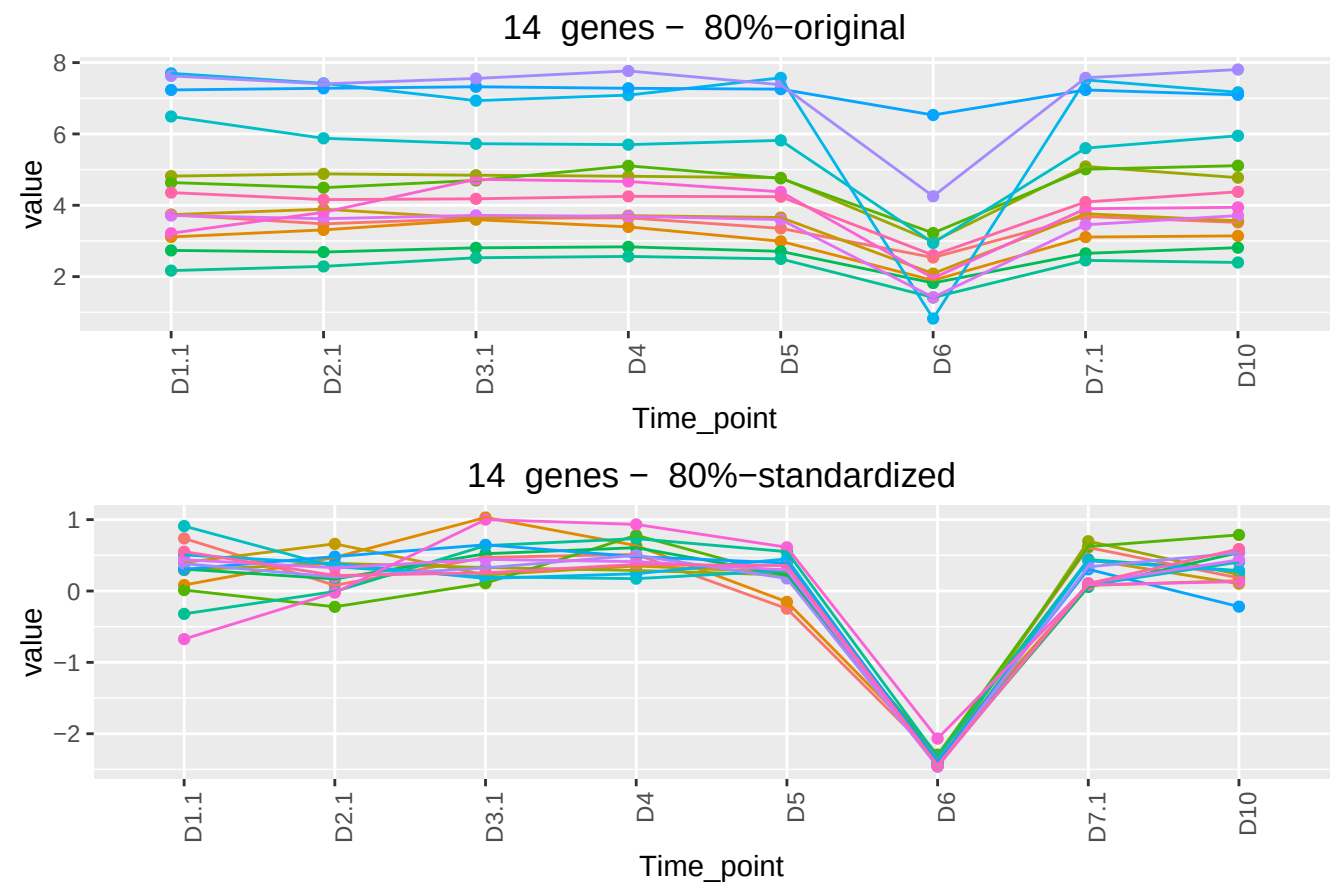
D7.1
D10
Time_point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value
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6
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2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point
14 genes − 80%−standardized
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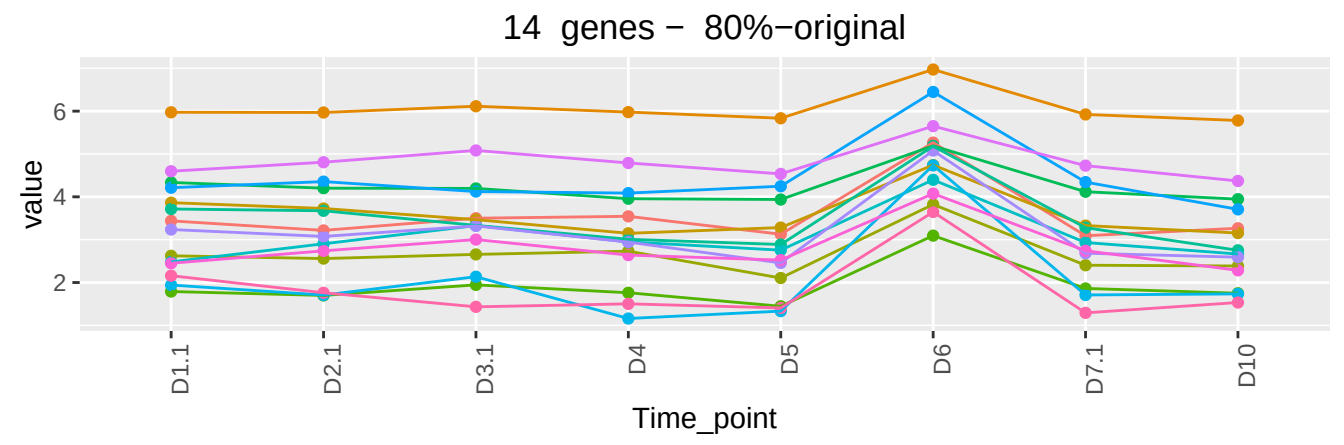
−2
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point


14 genes − 80%−original
value
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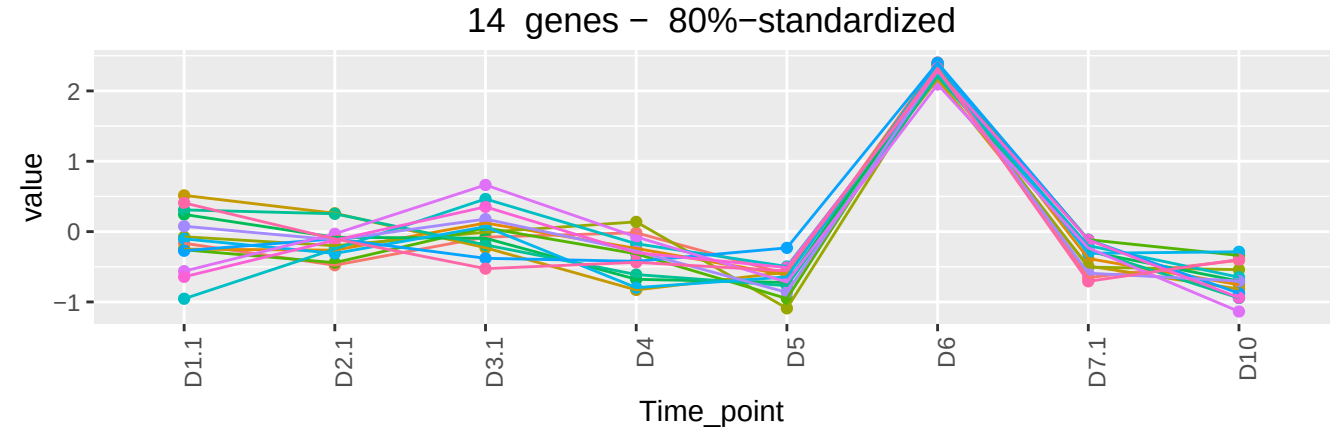
D1.1
D2.1
D3.1
D4
D5
D6
D7.1
D10
Time_point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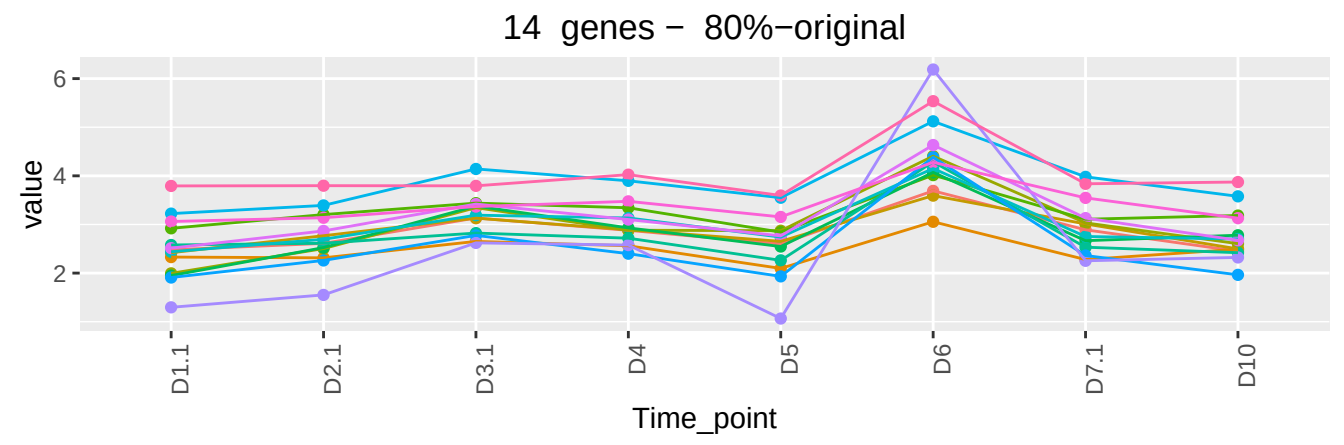
Time_point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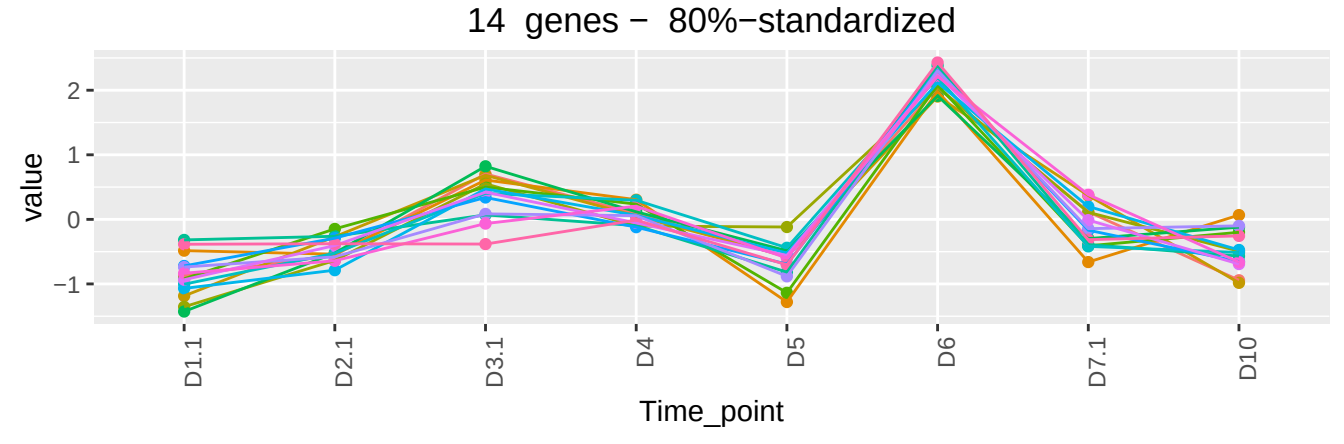
Time_point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D6
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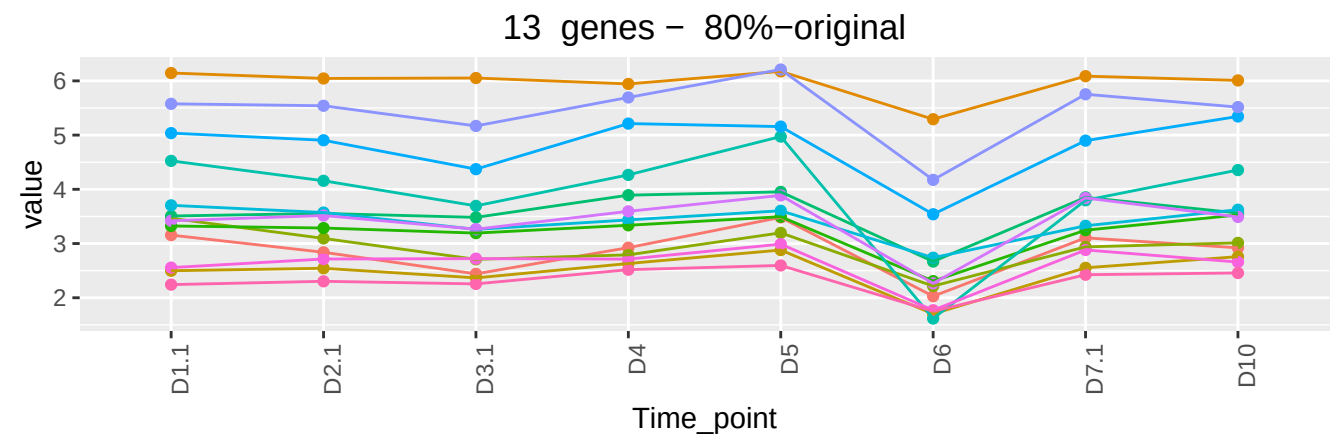
Time_point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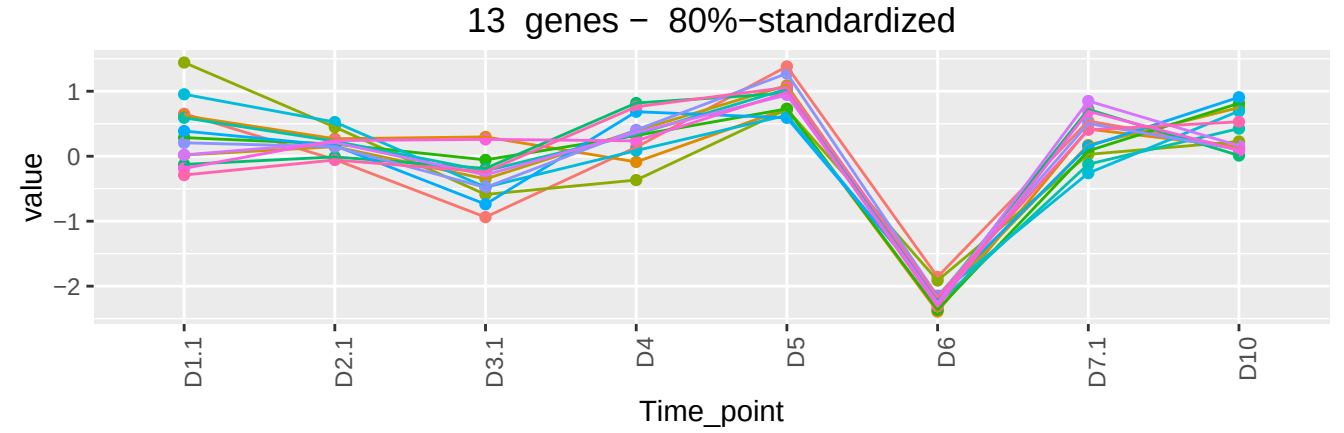
Time_point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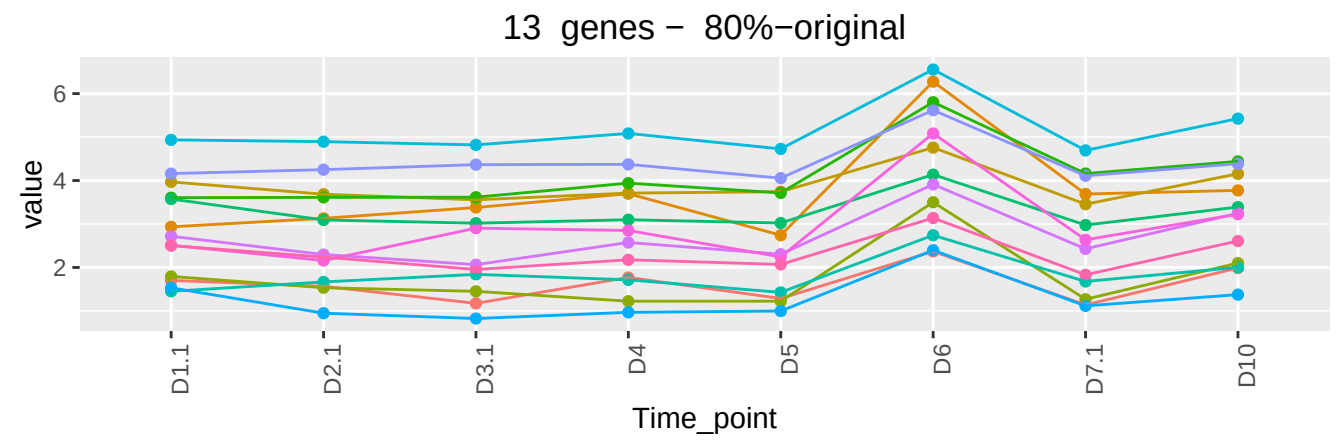
Time_point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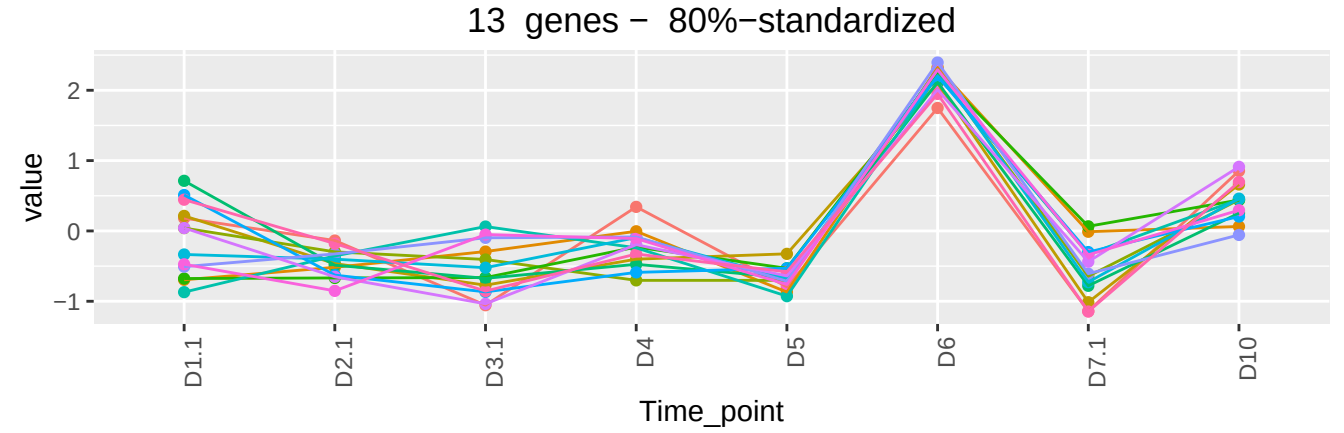
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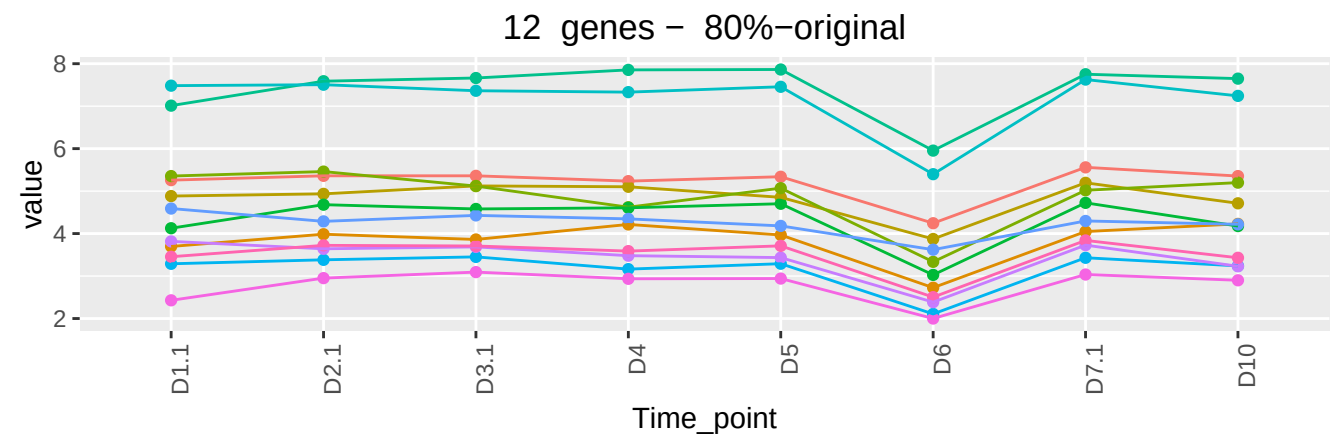
12 genes − 80%−original
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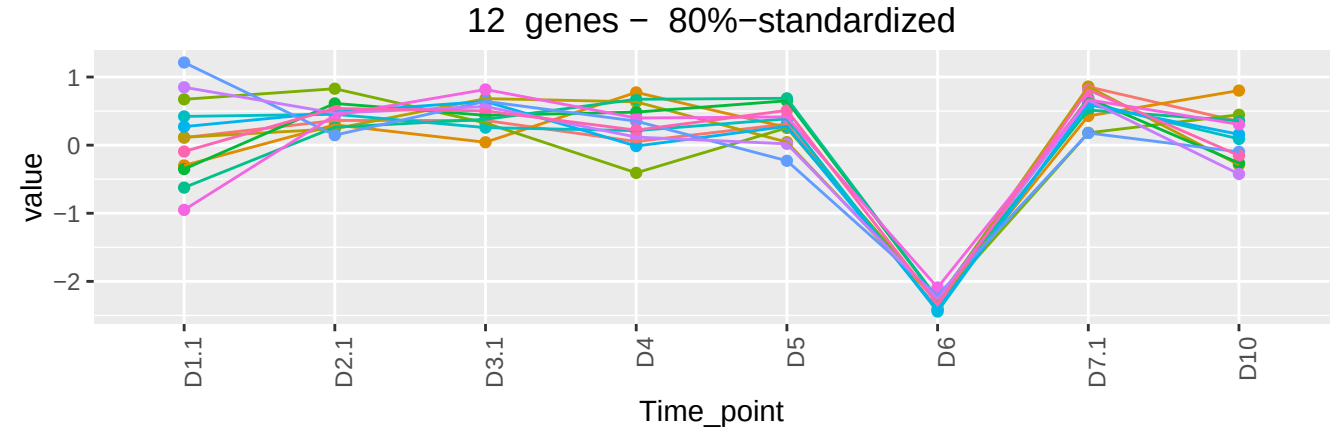
Time_point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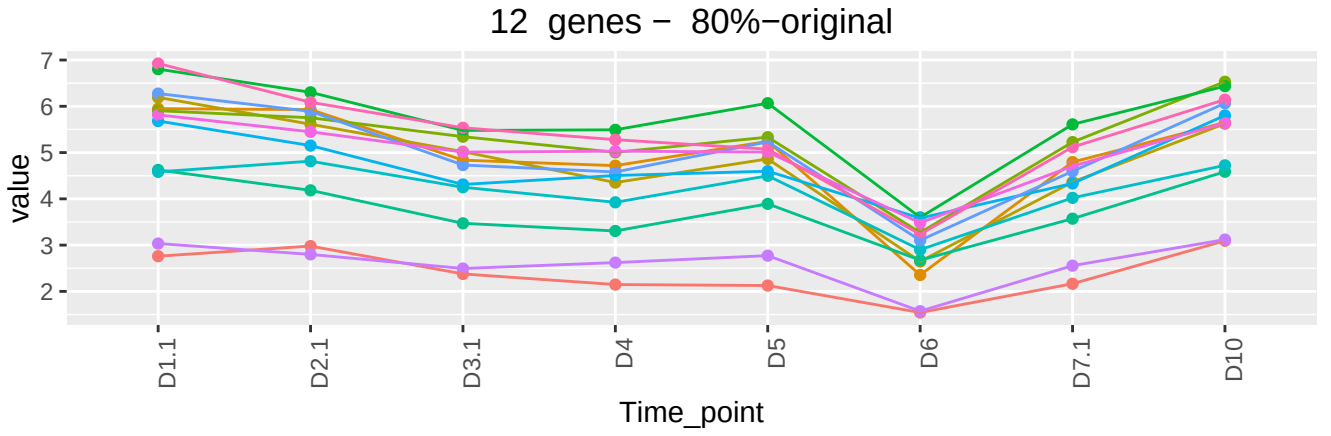
Time_point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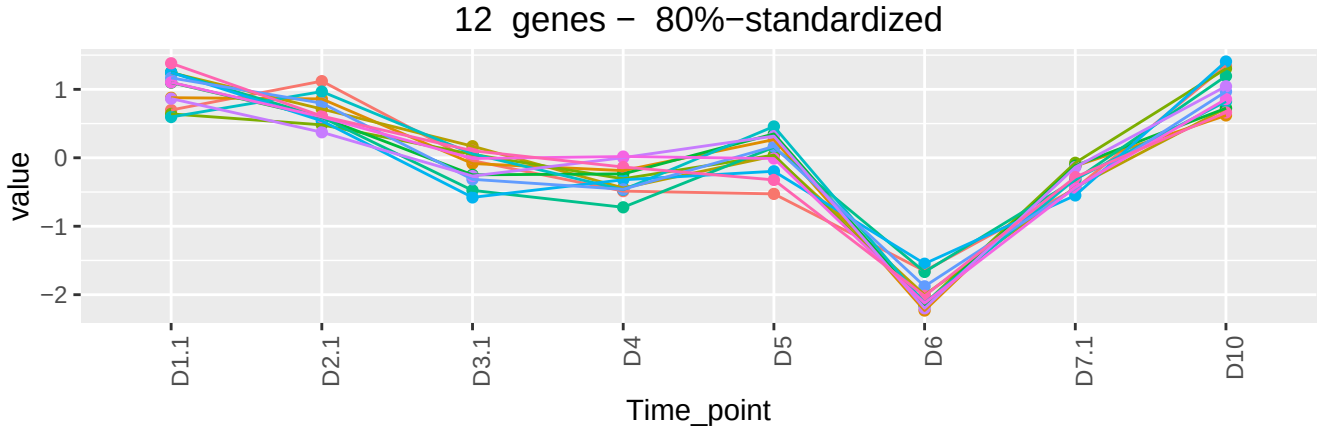
Time_point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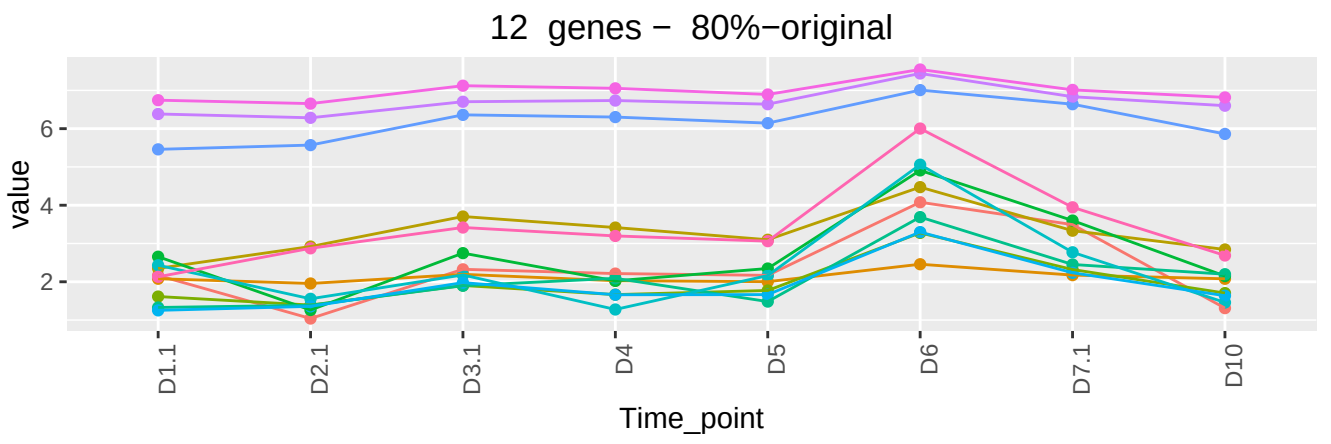
Time_point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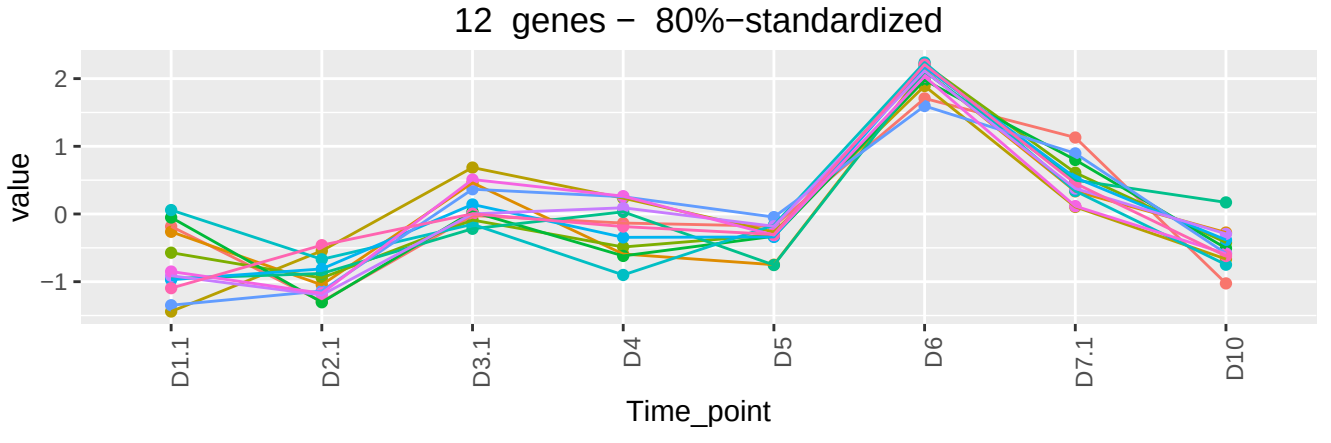
Time_point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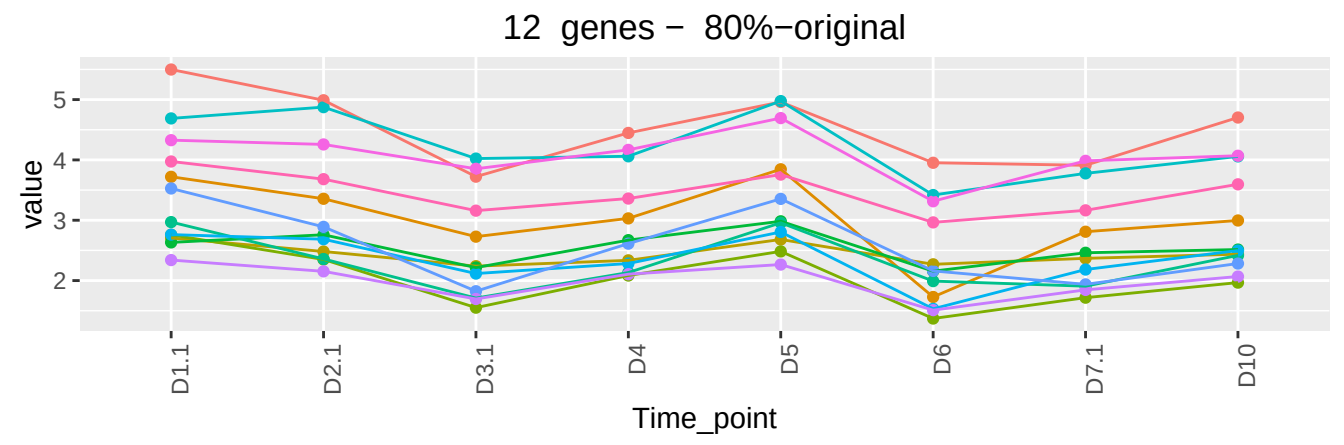
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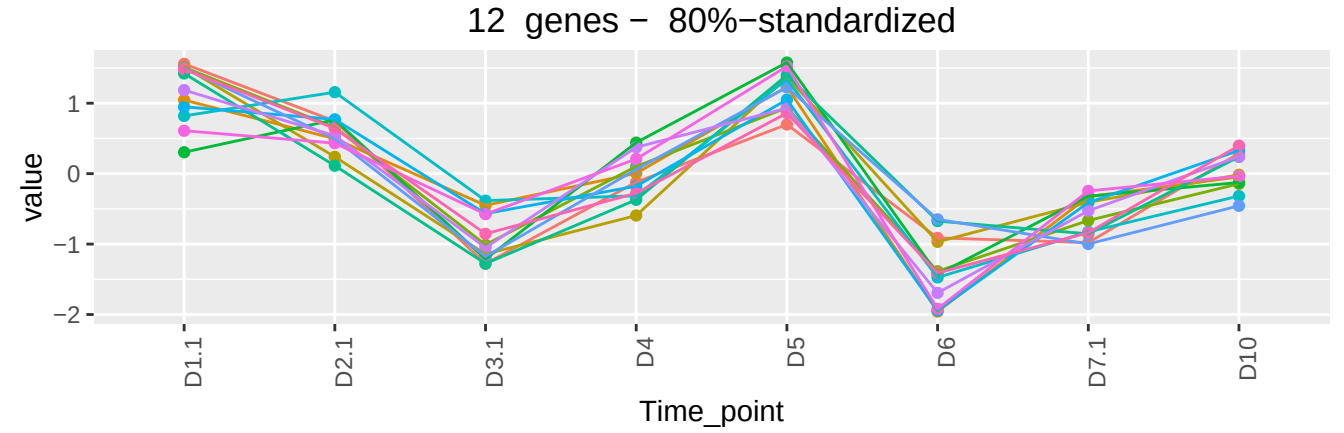
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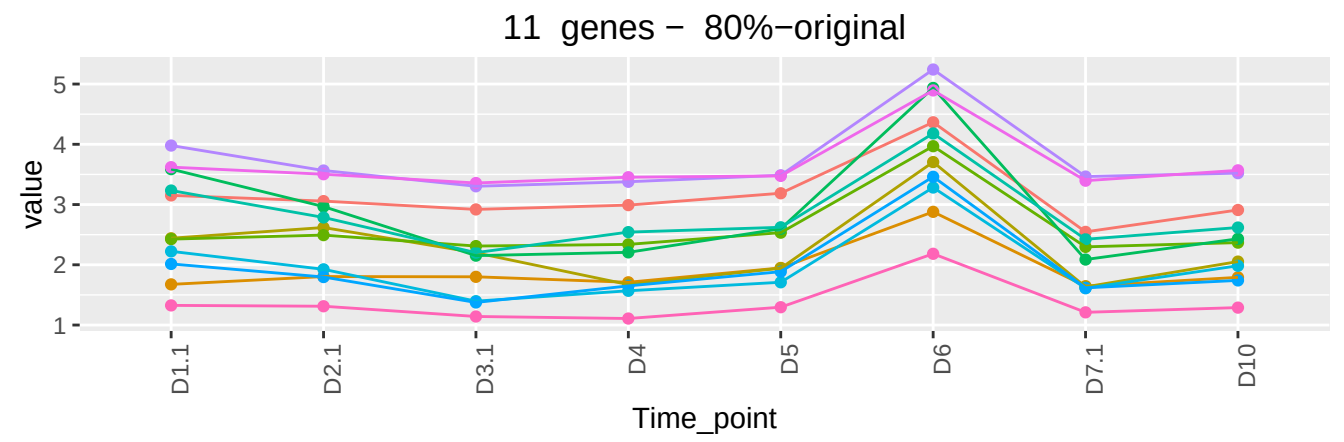
Time_point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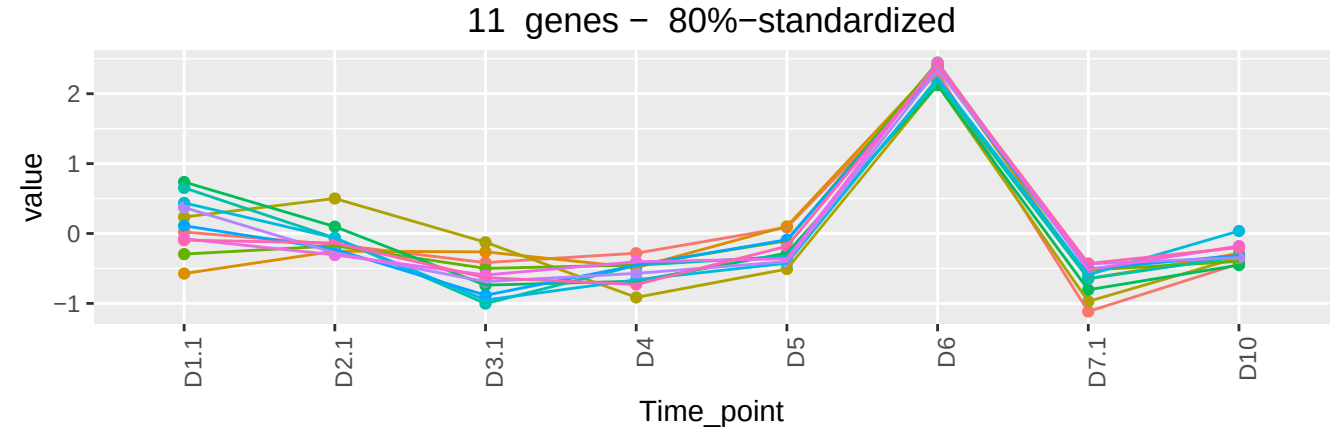
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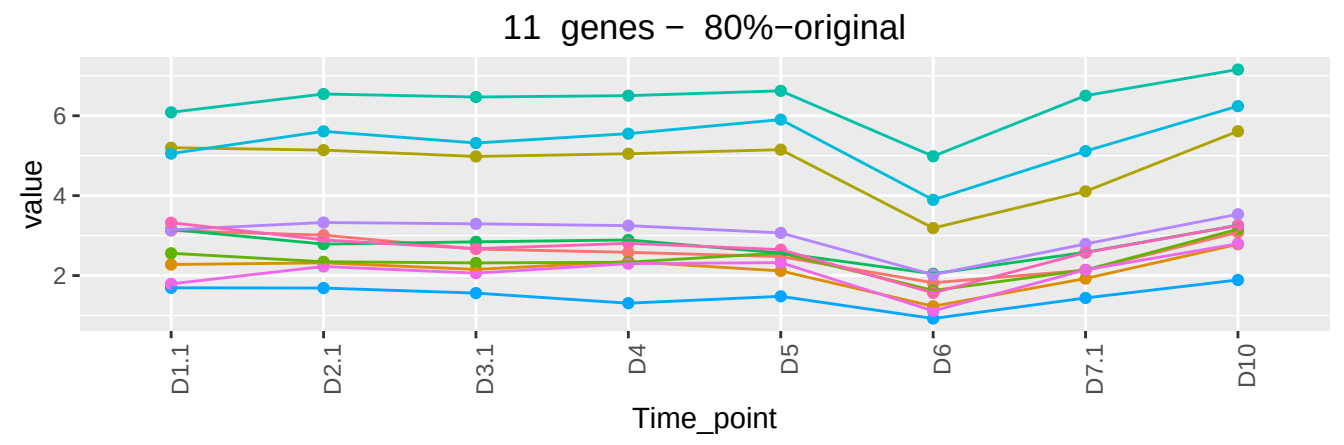
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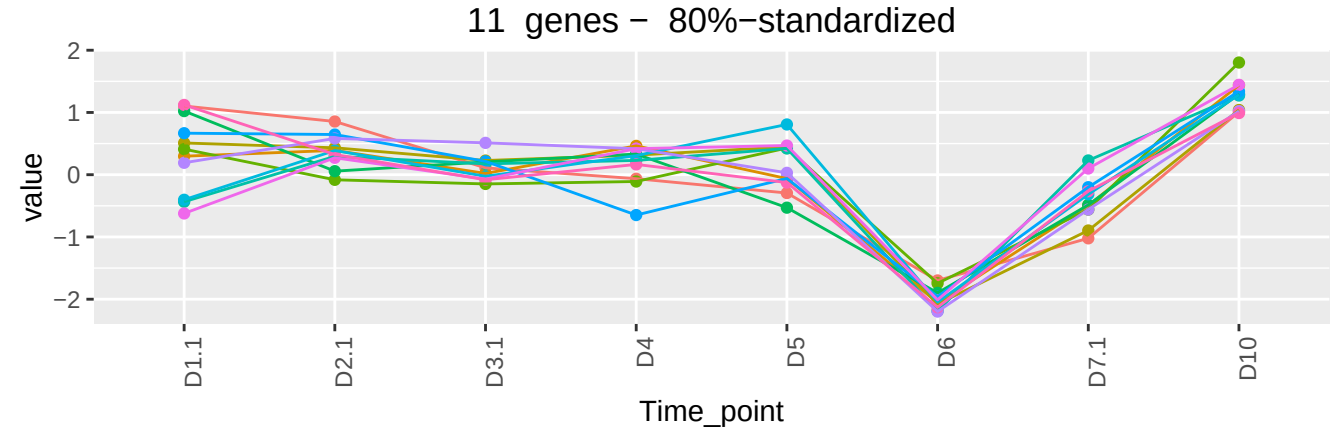
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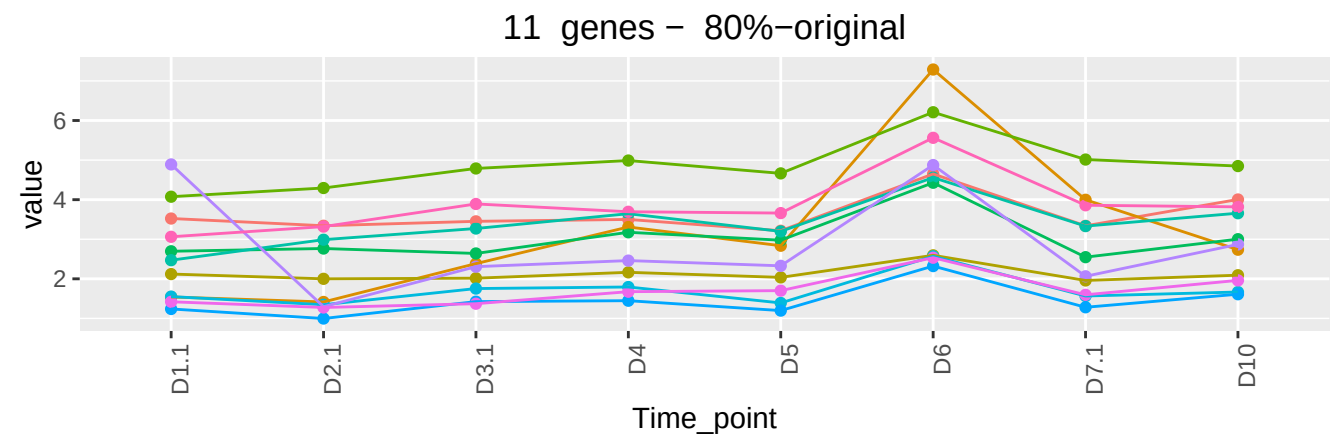
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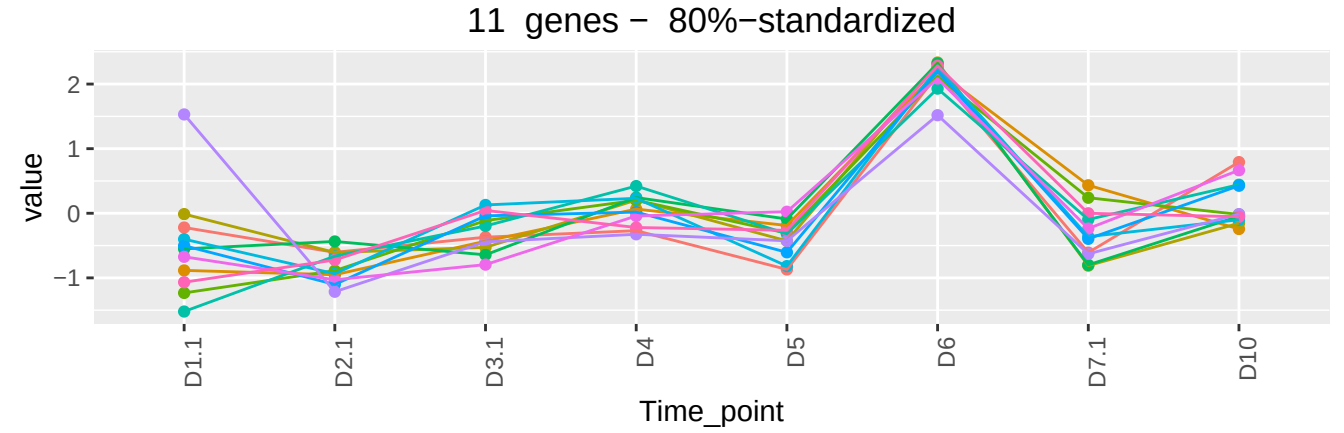
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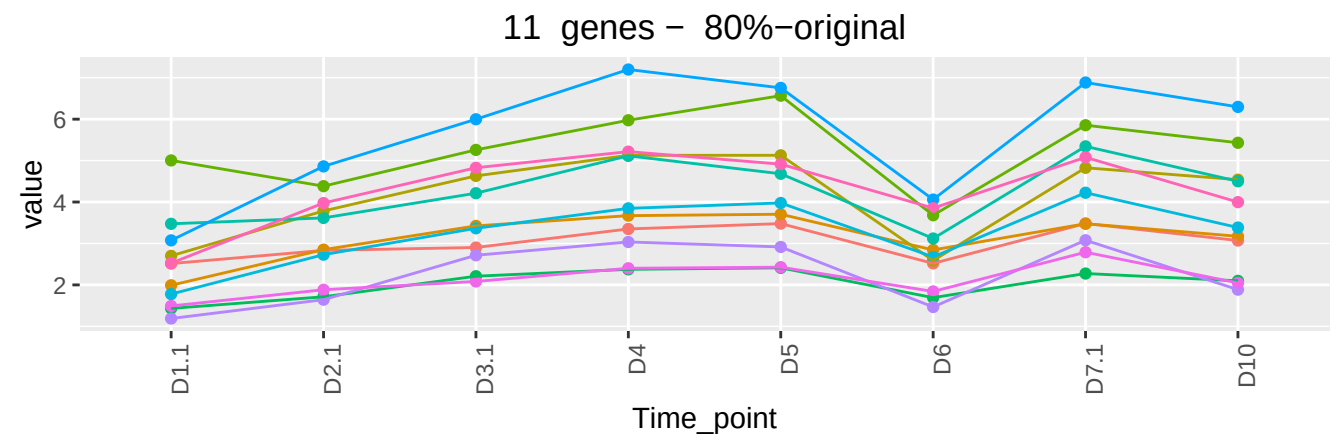
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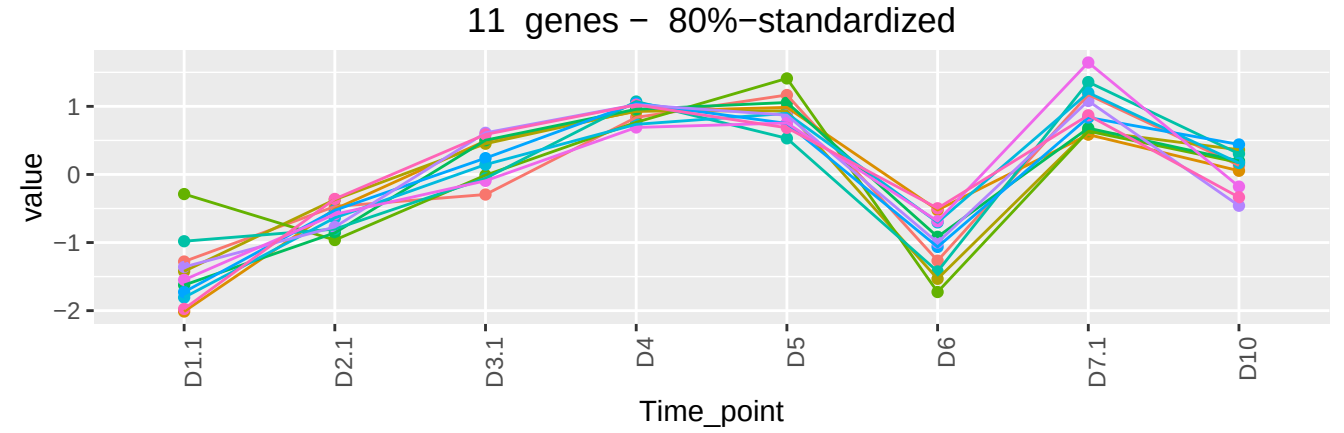
Time_point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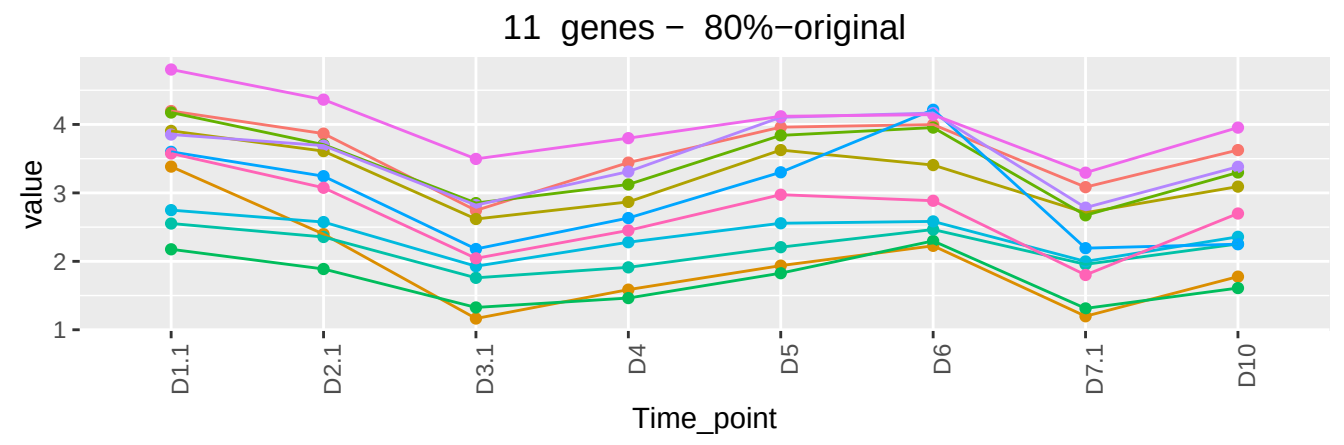
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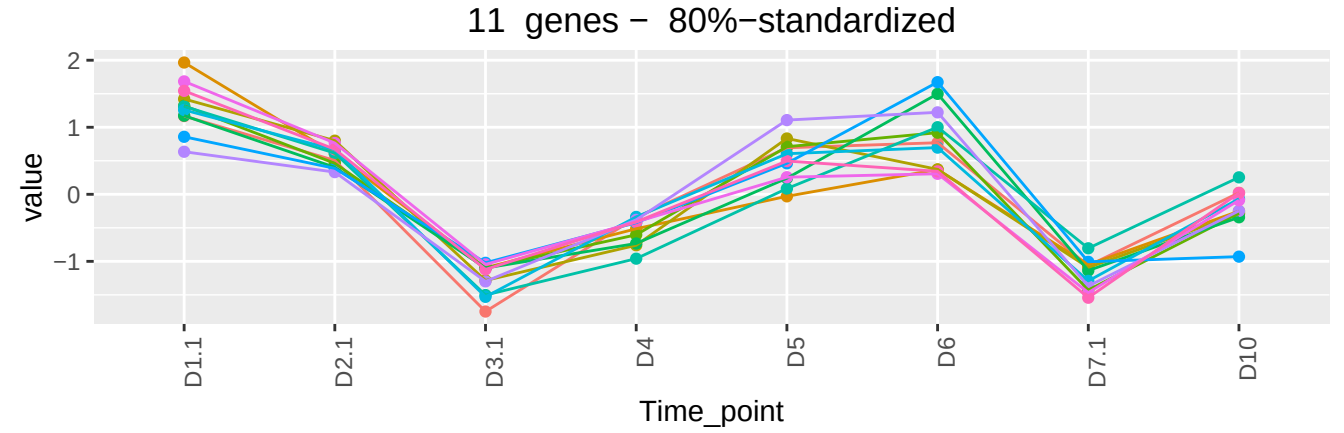
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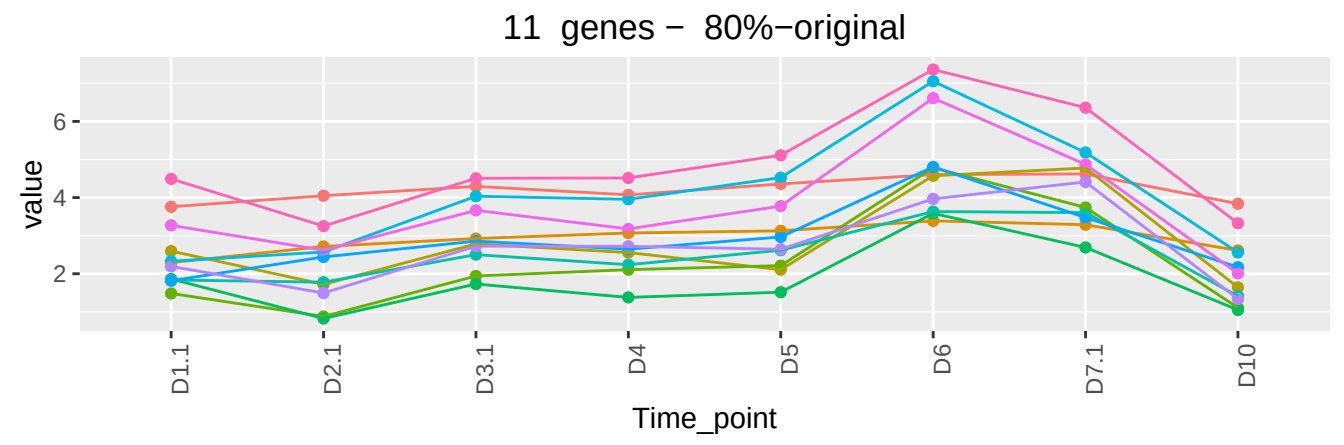
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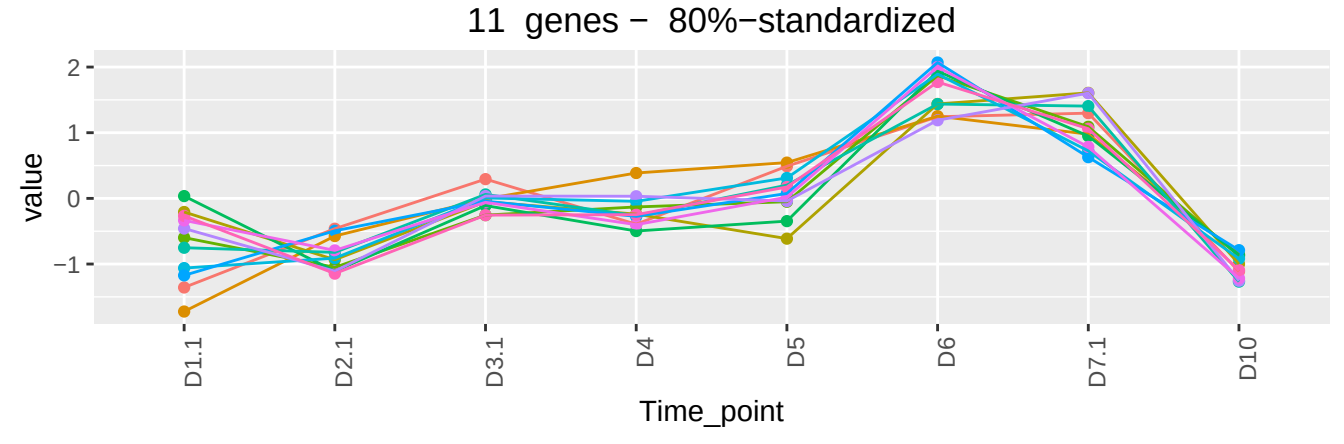
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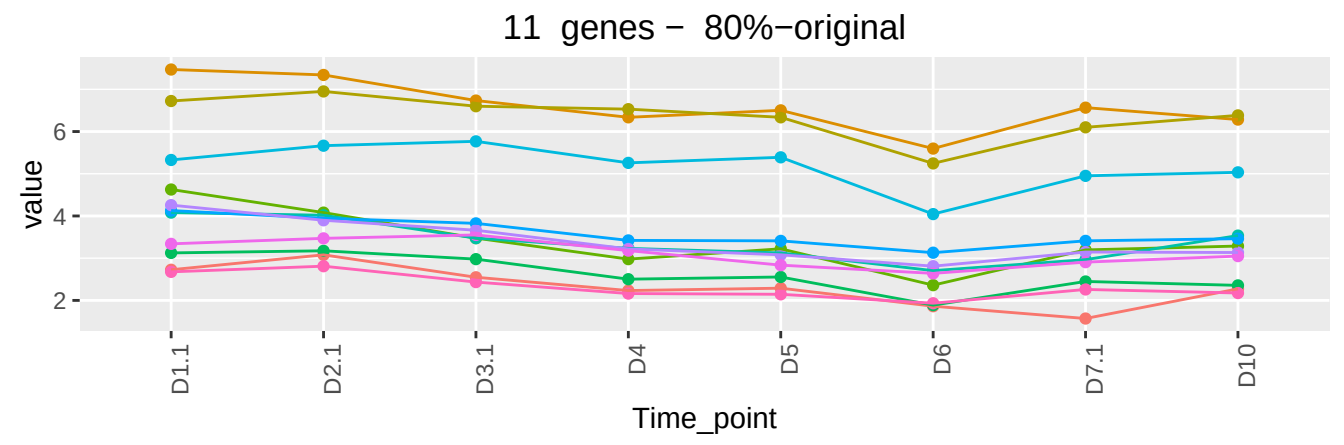
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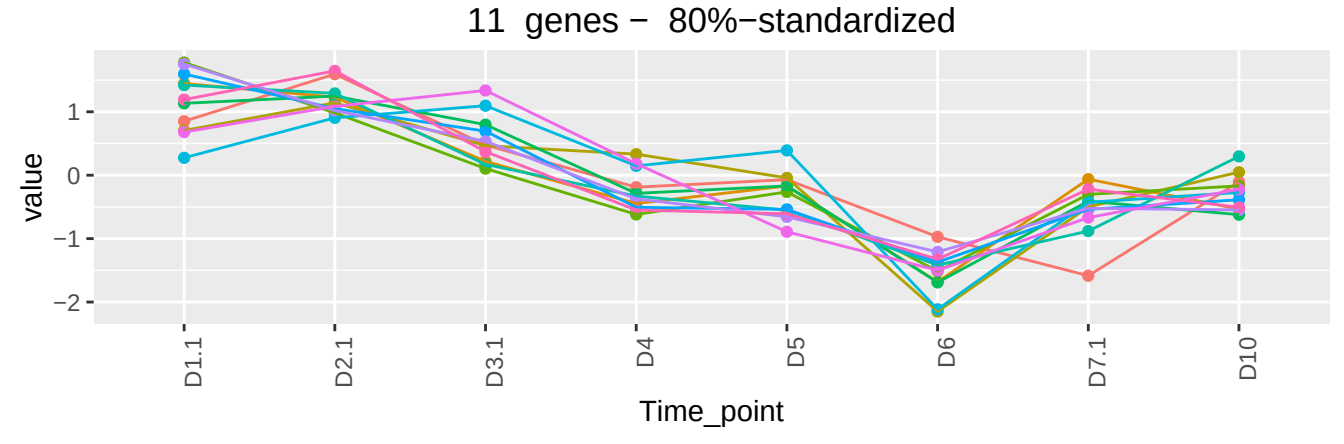
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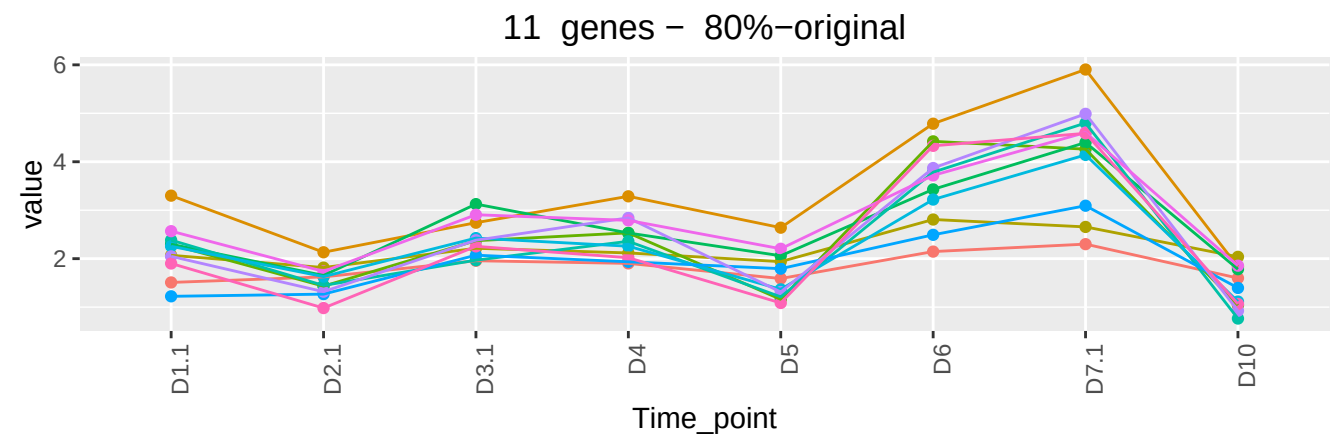
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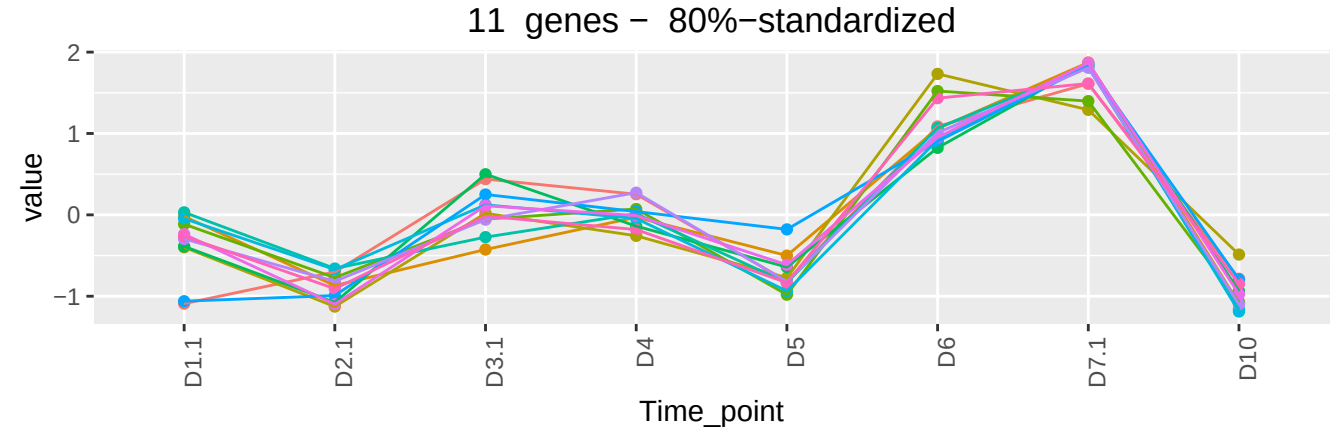
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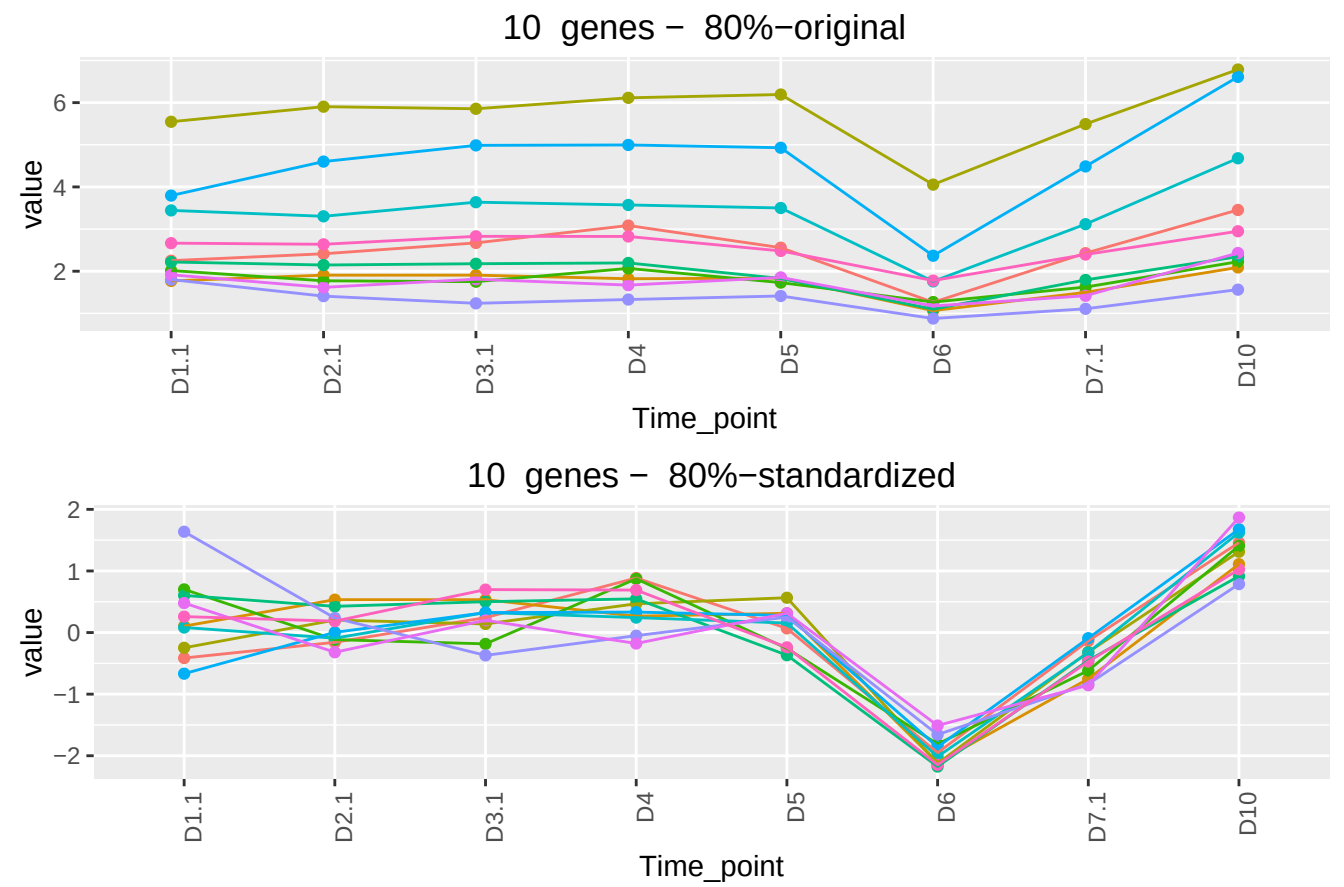
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D6
D7.1
D10
Time_point
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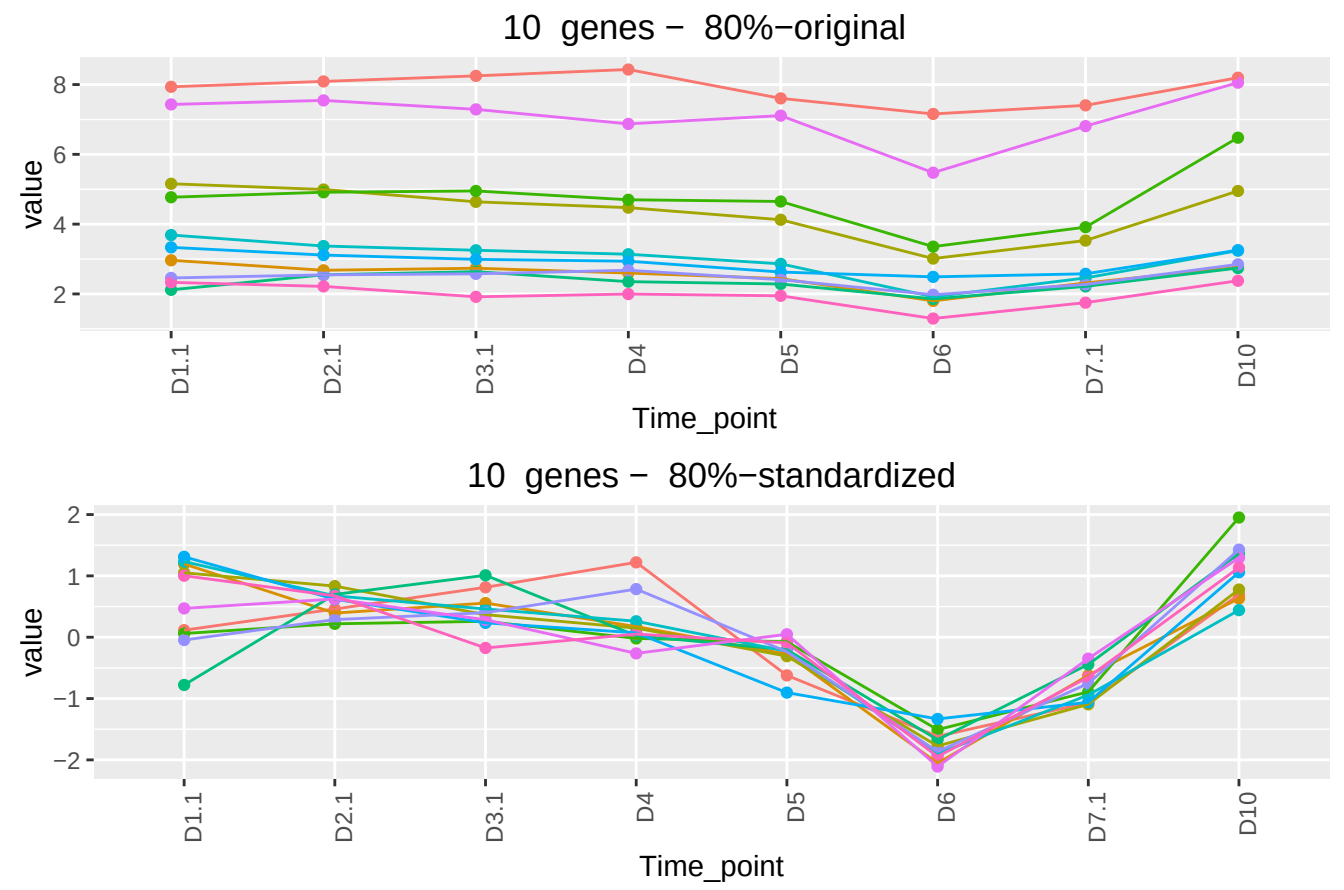
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Time_point
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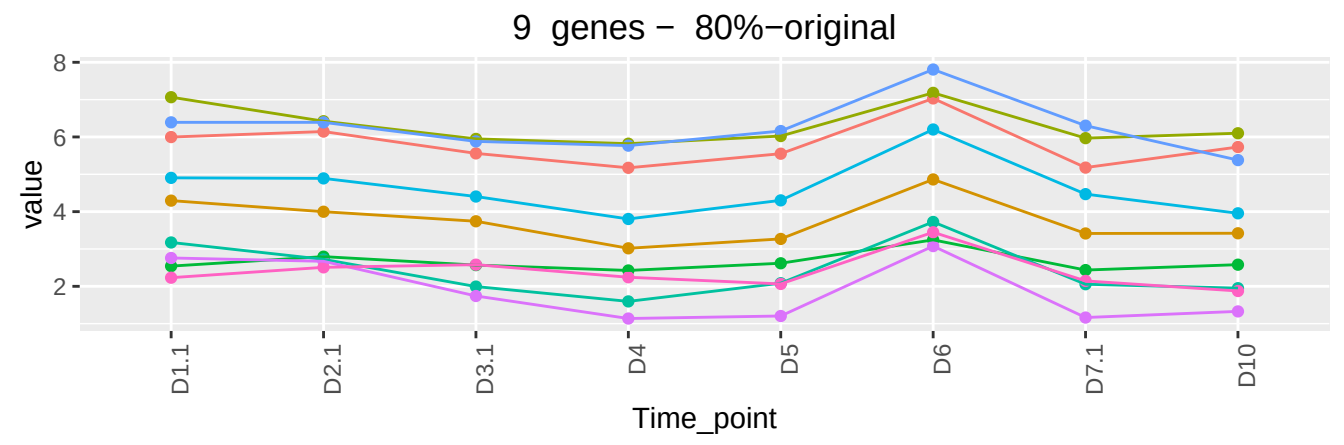
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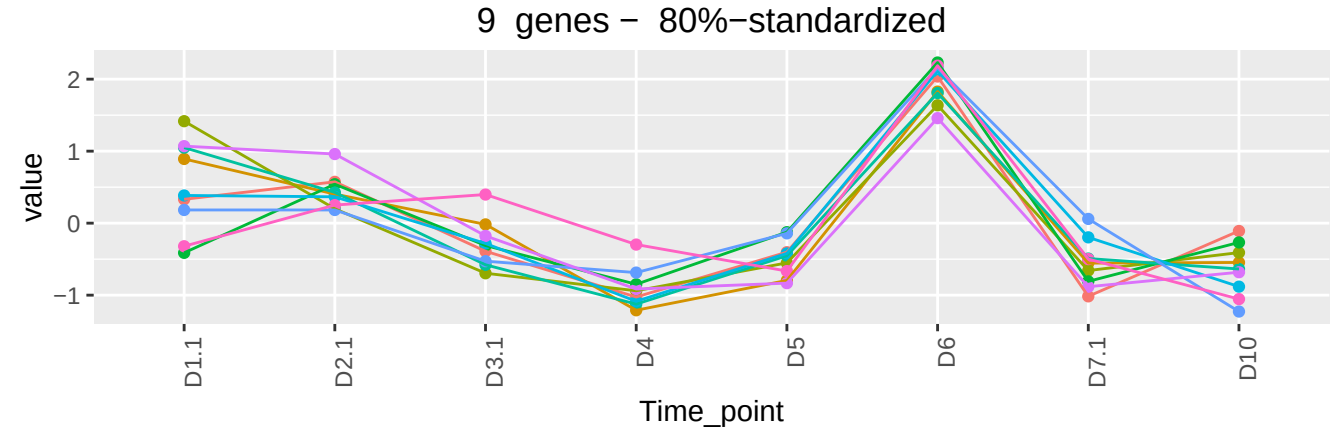
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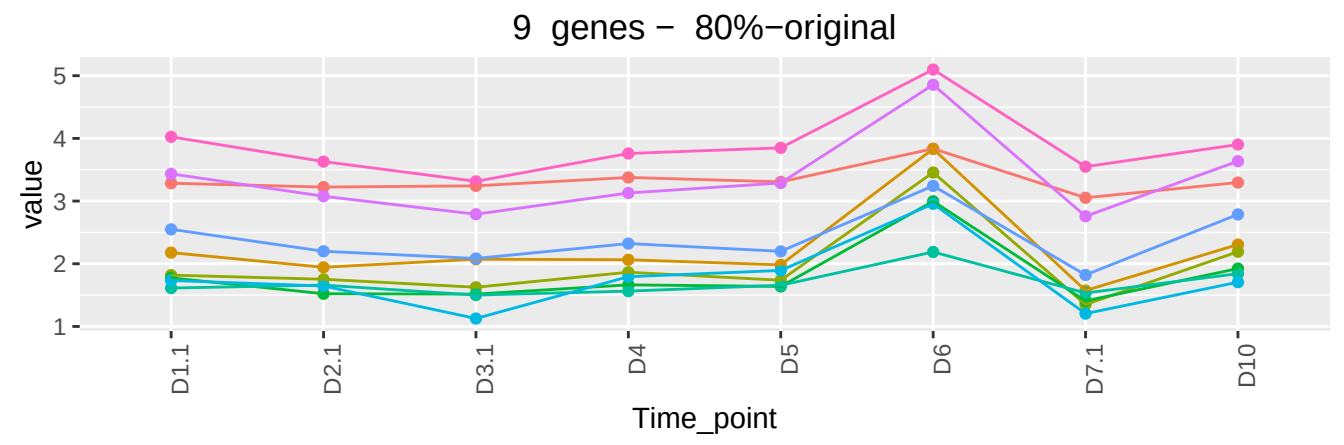
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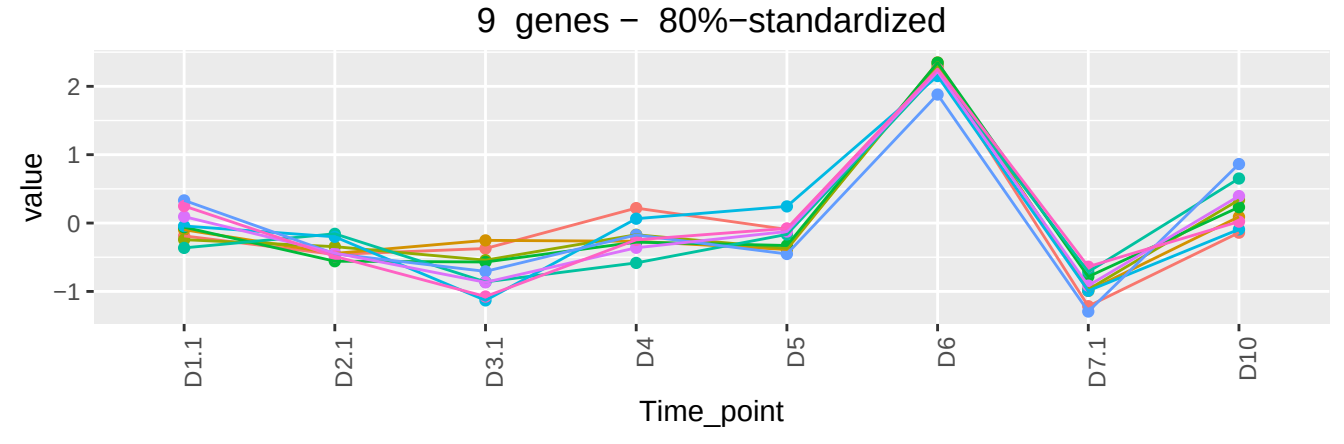
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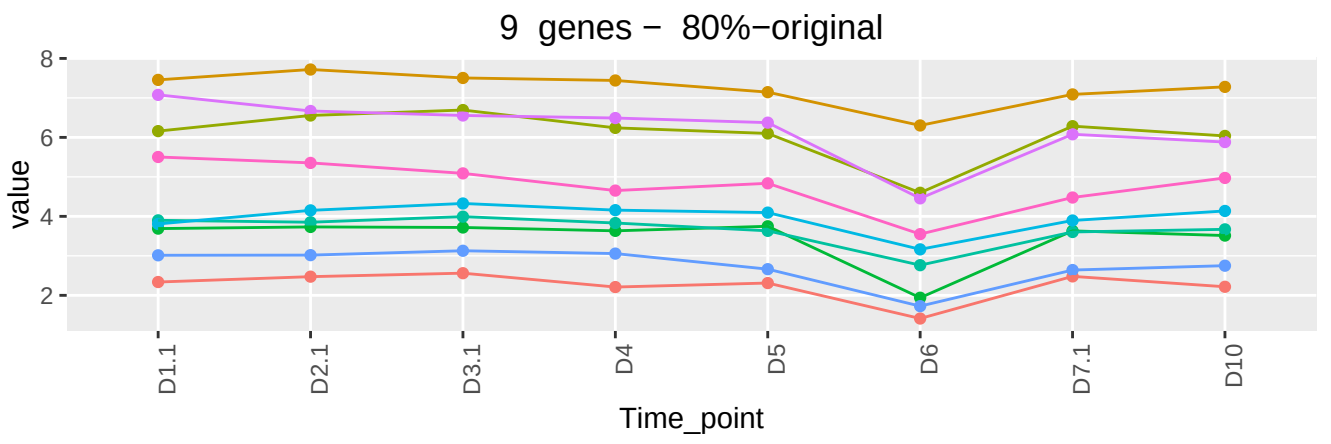
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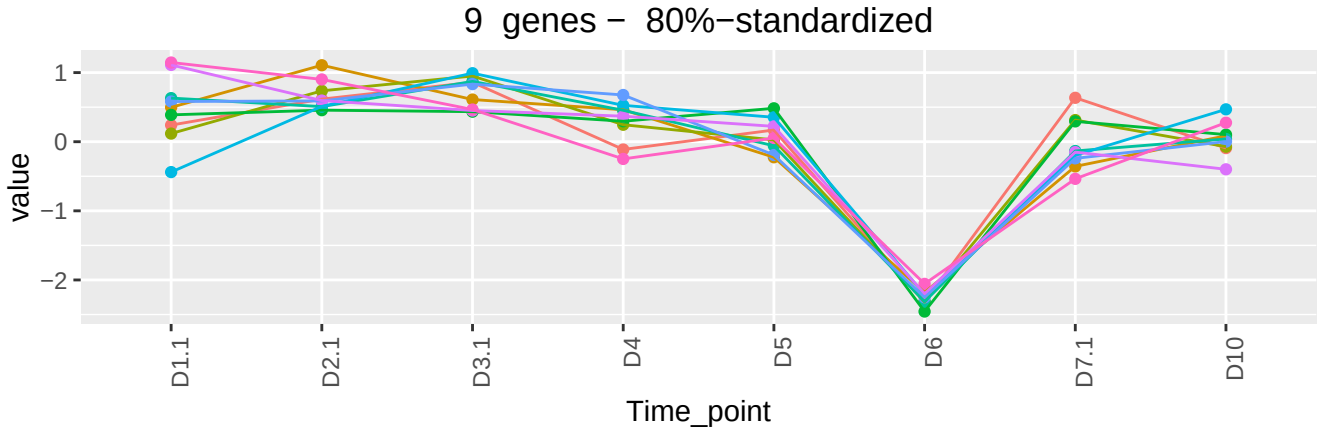
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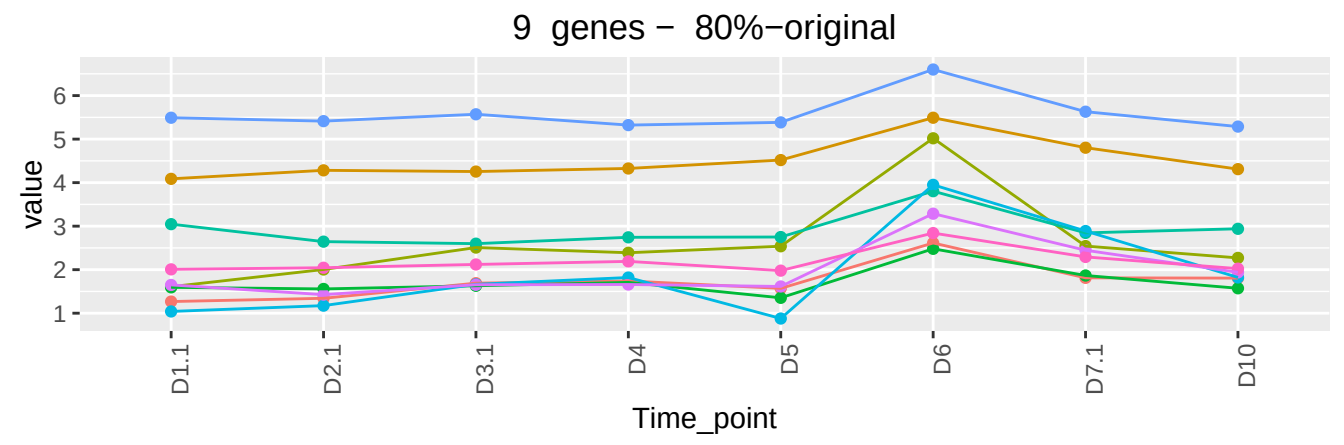
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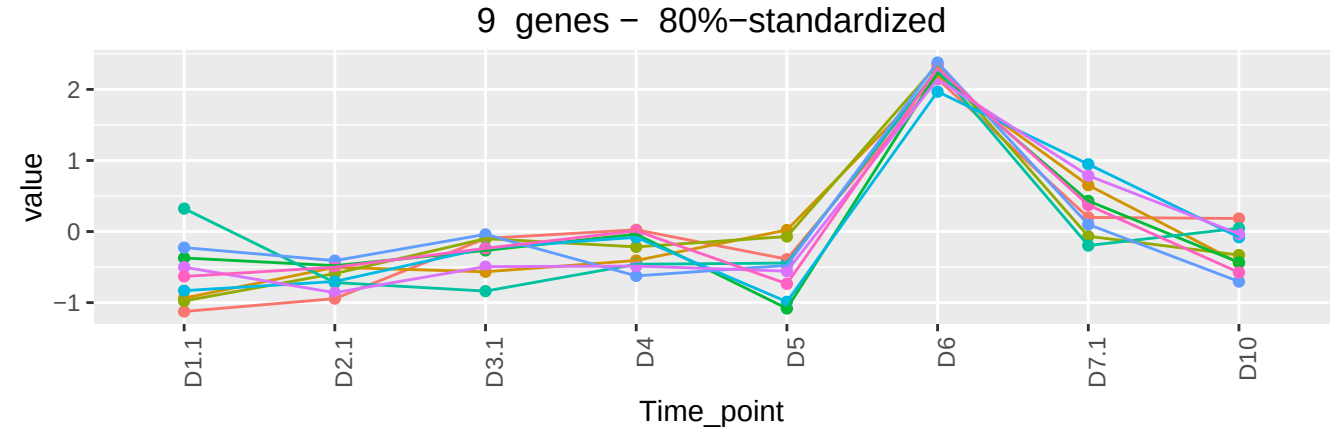
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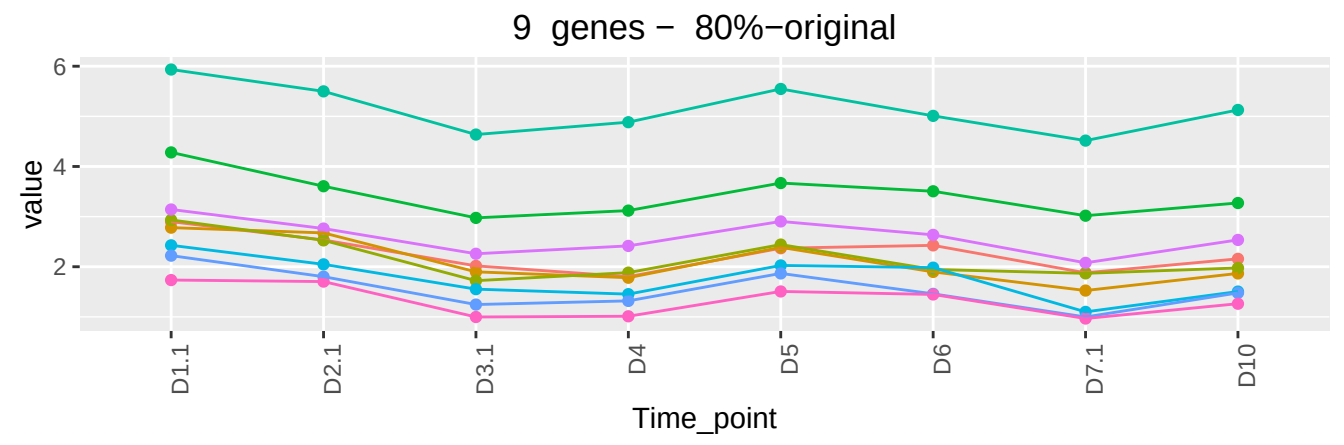
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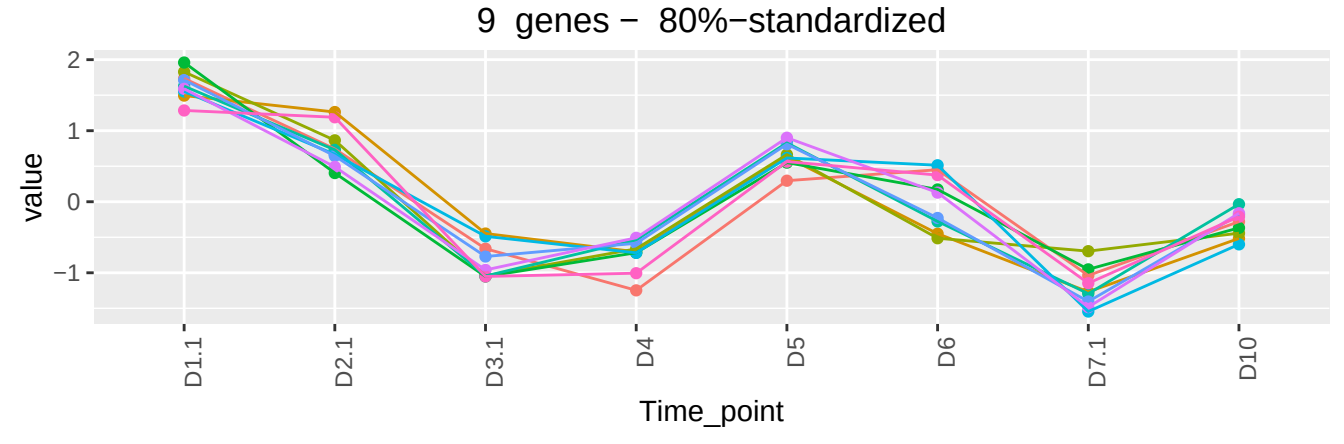
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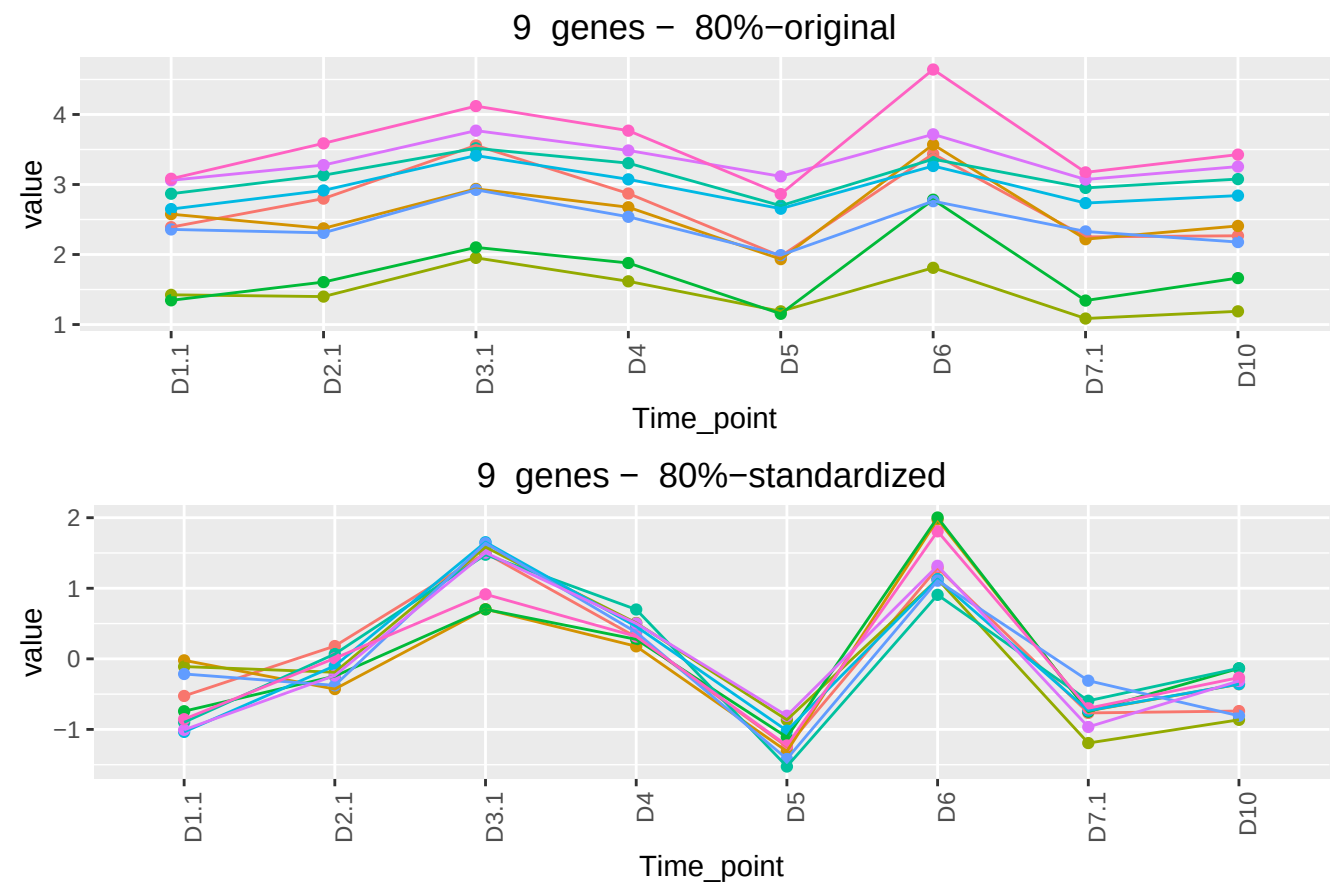
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Time_point

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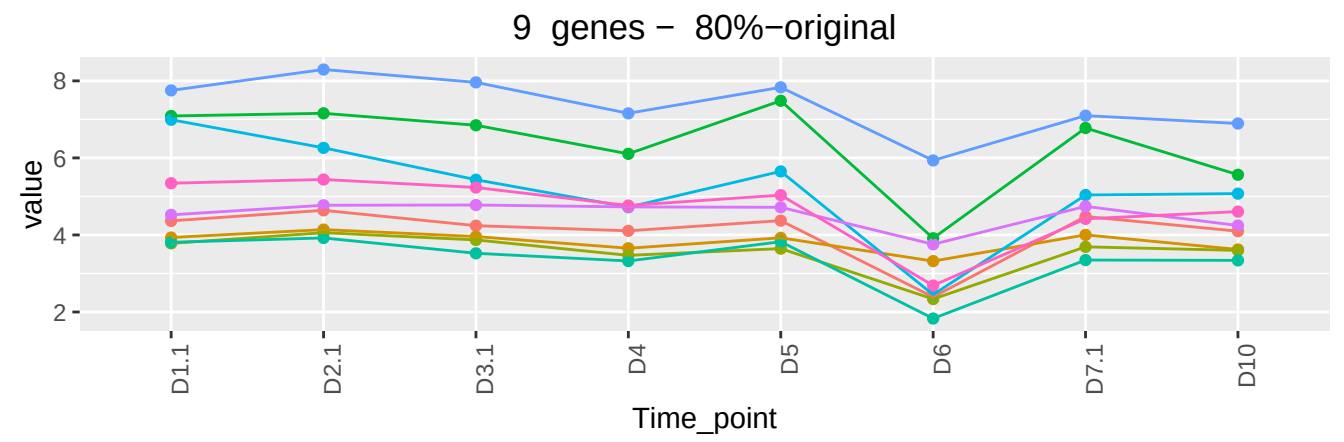
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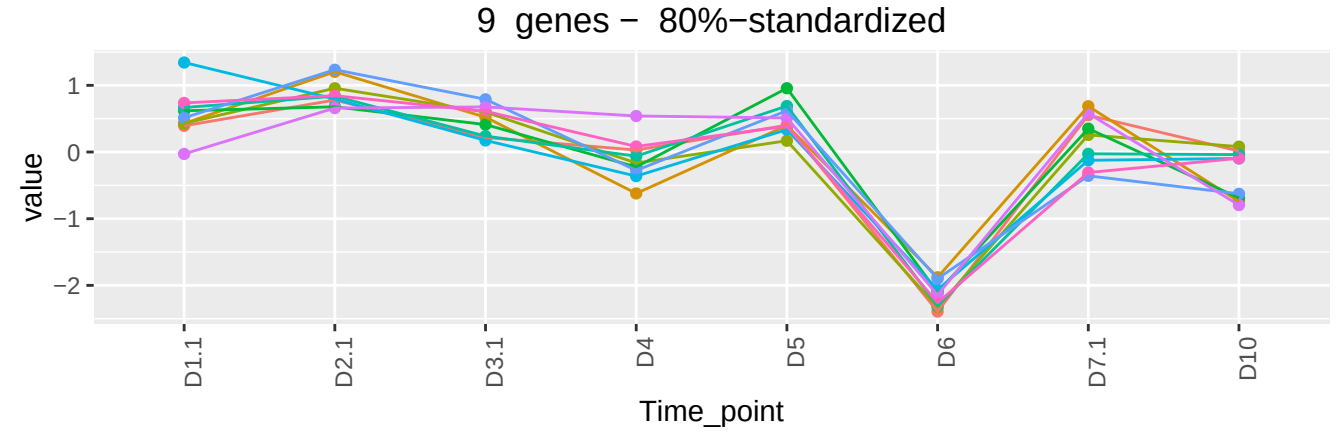
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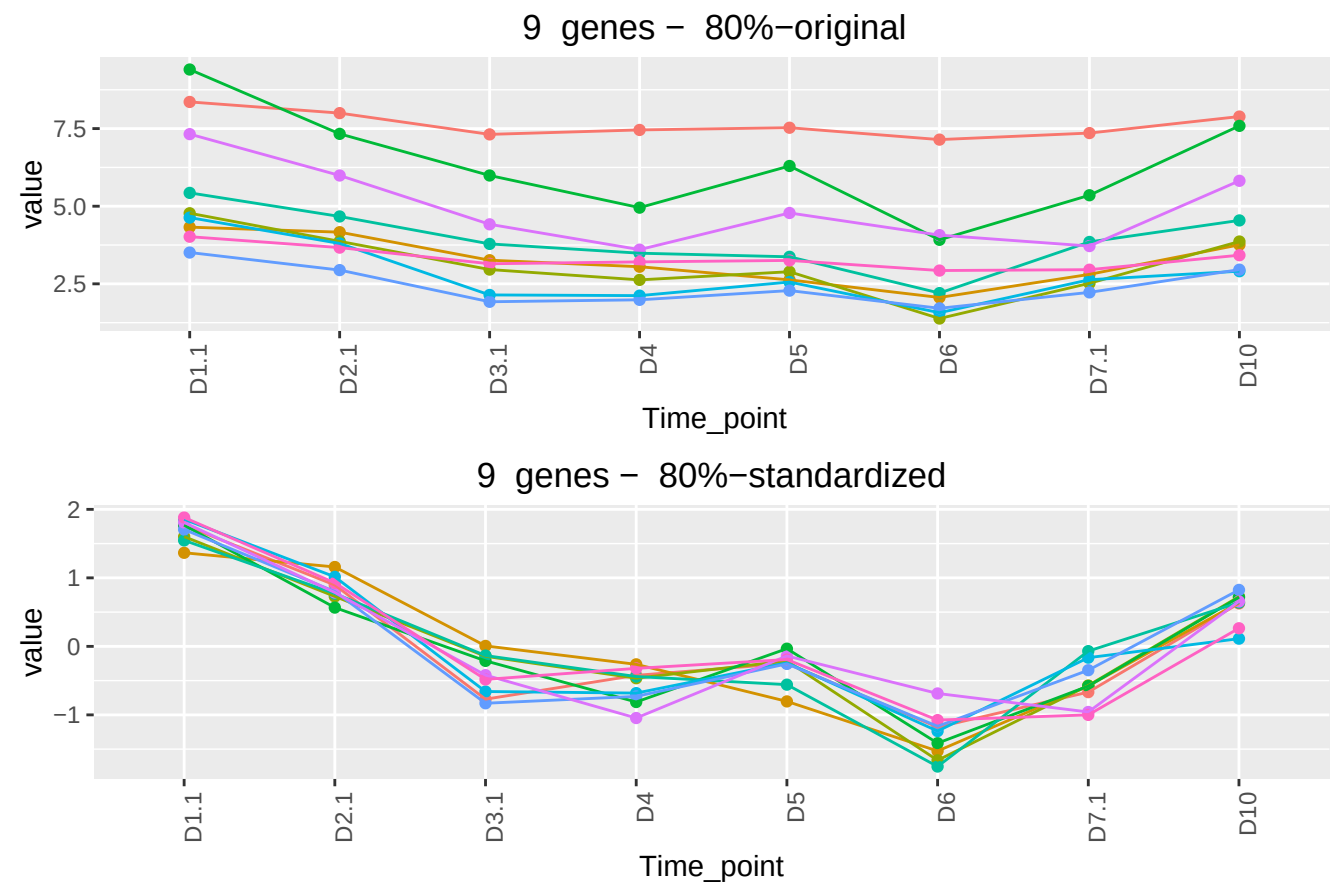
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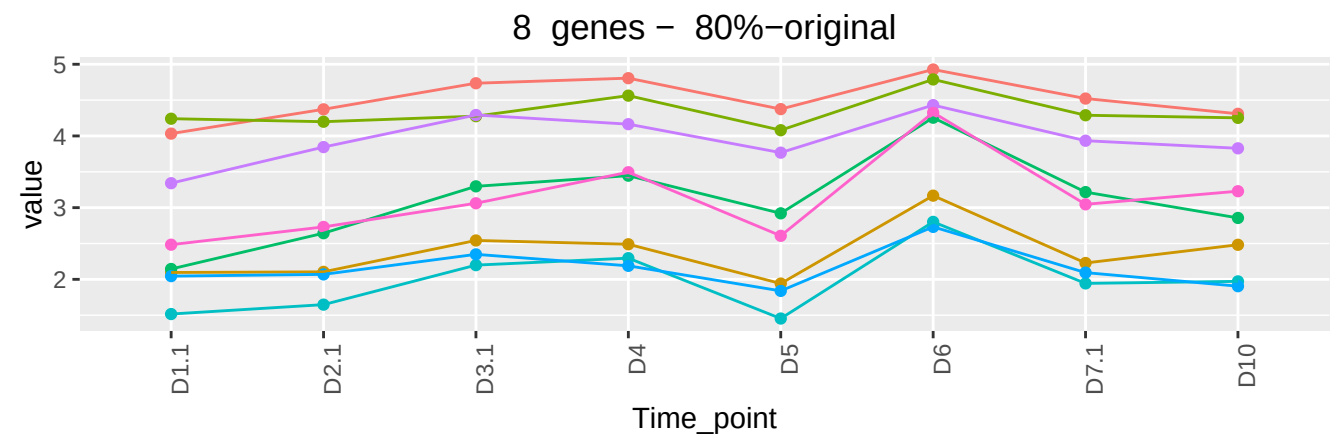
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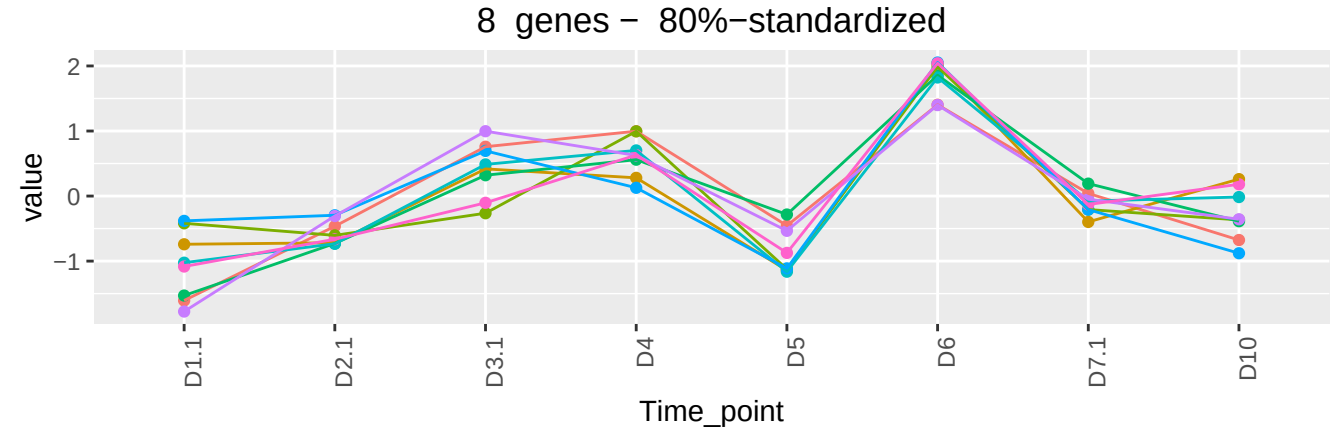
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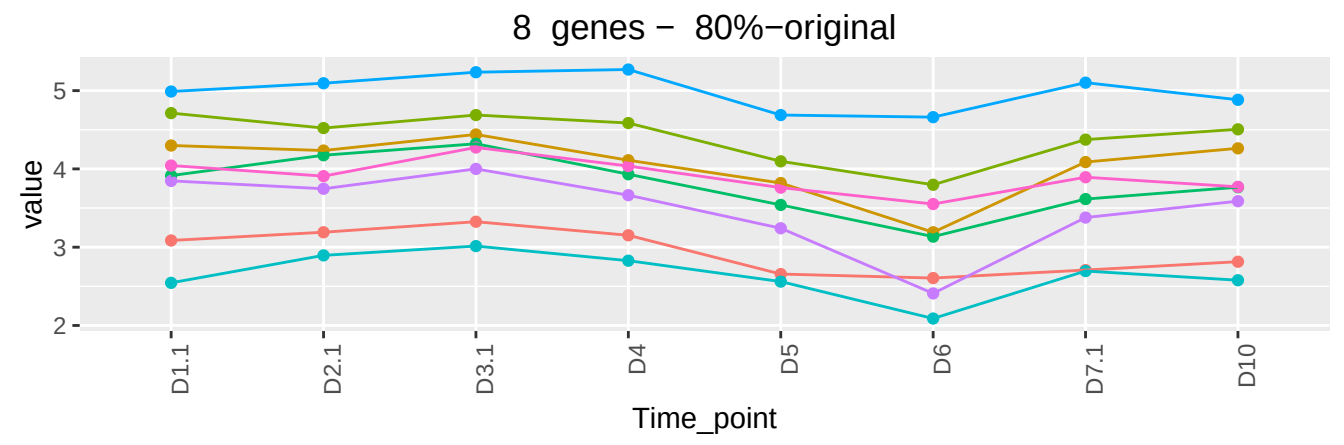
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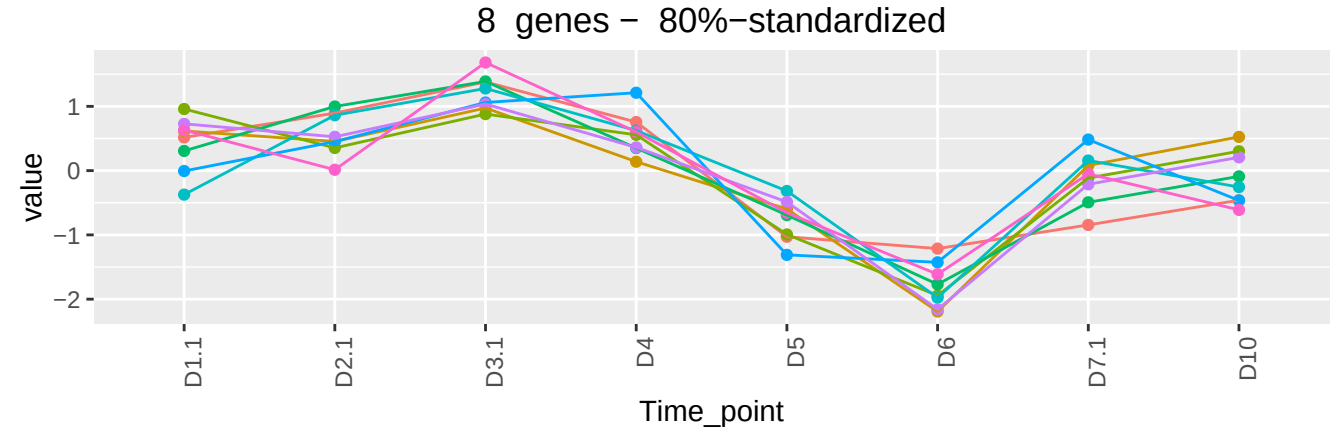
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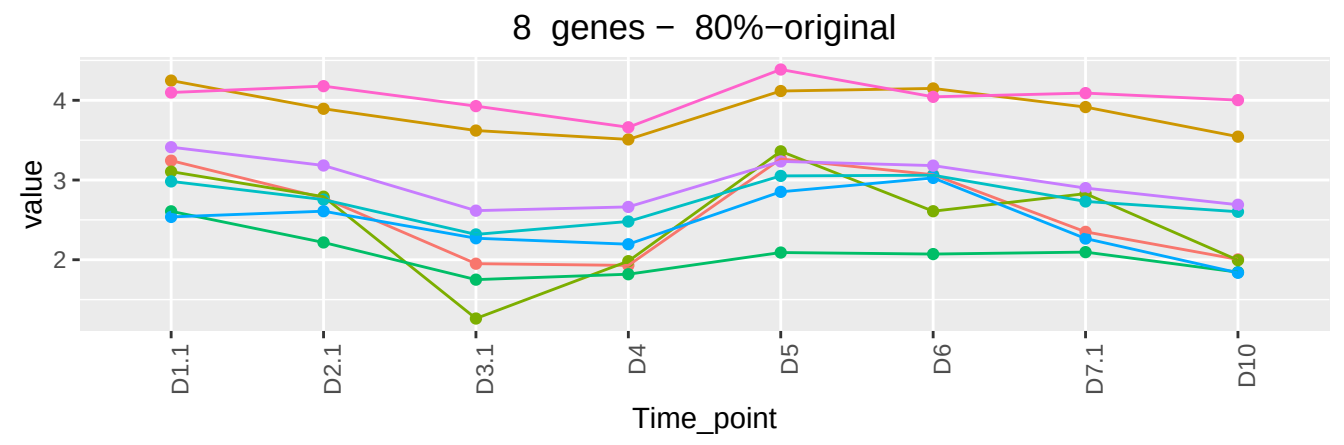
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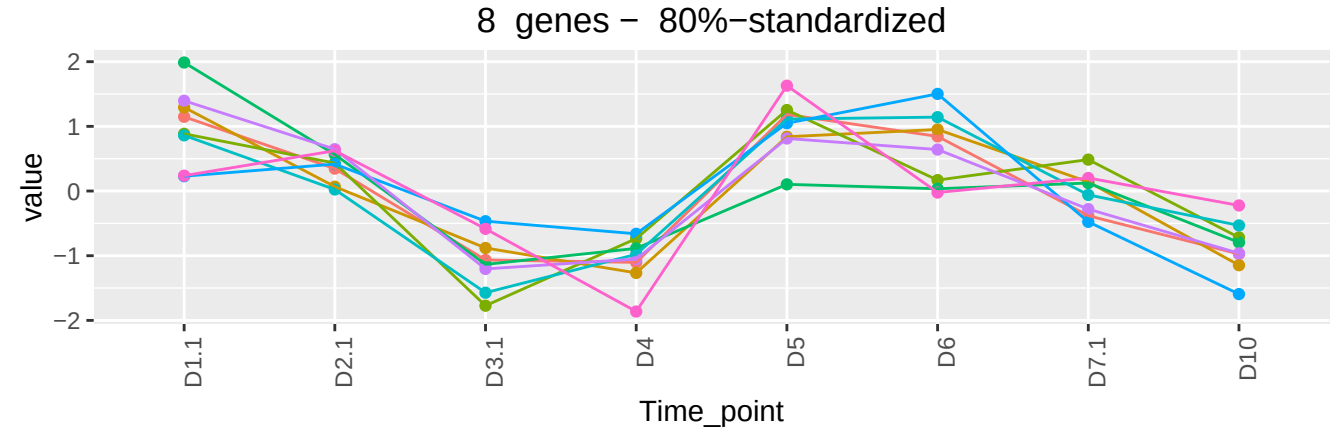
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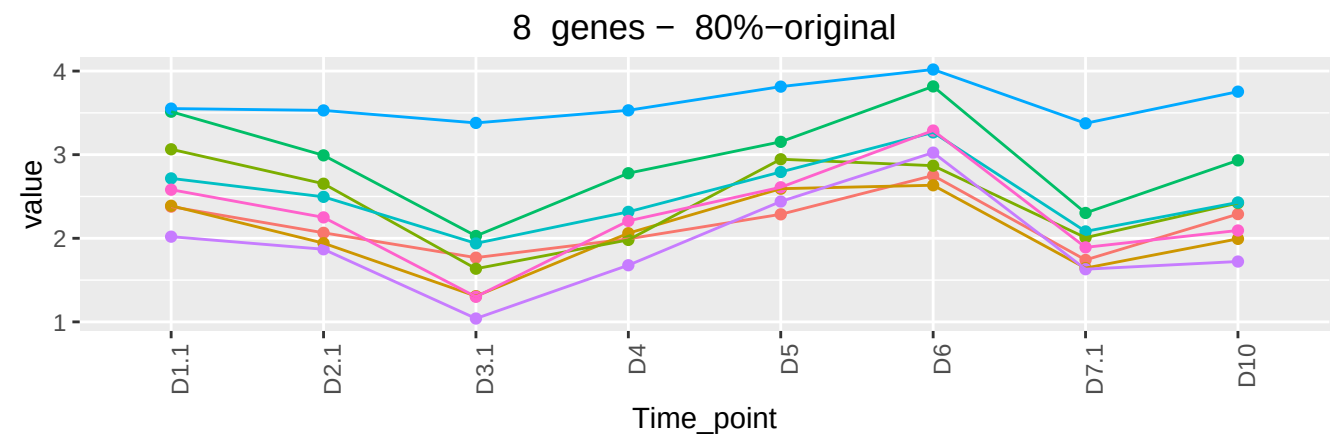
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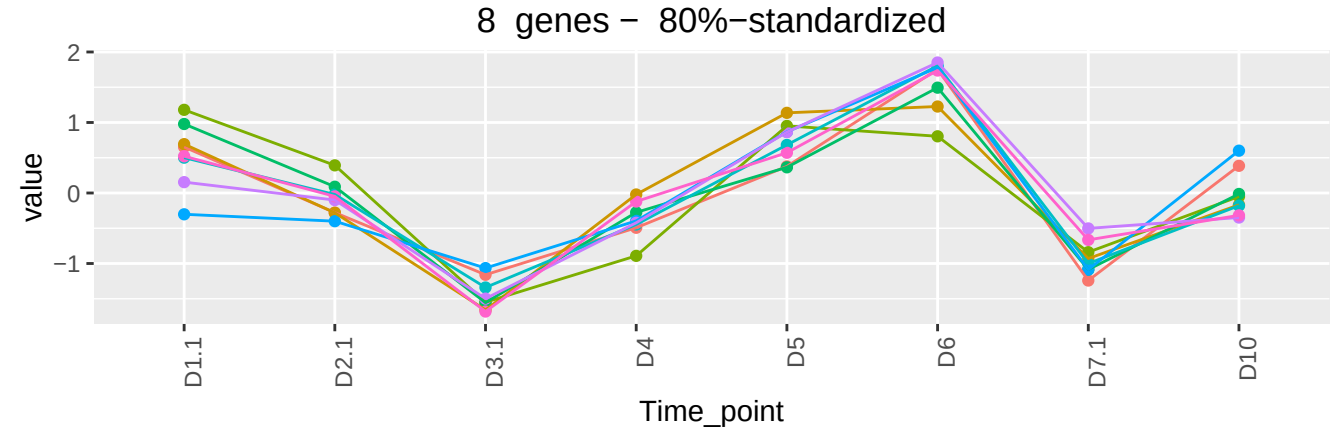
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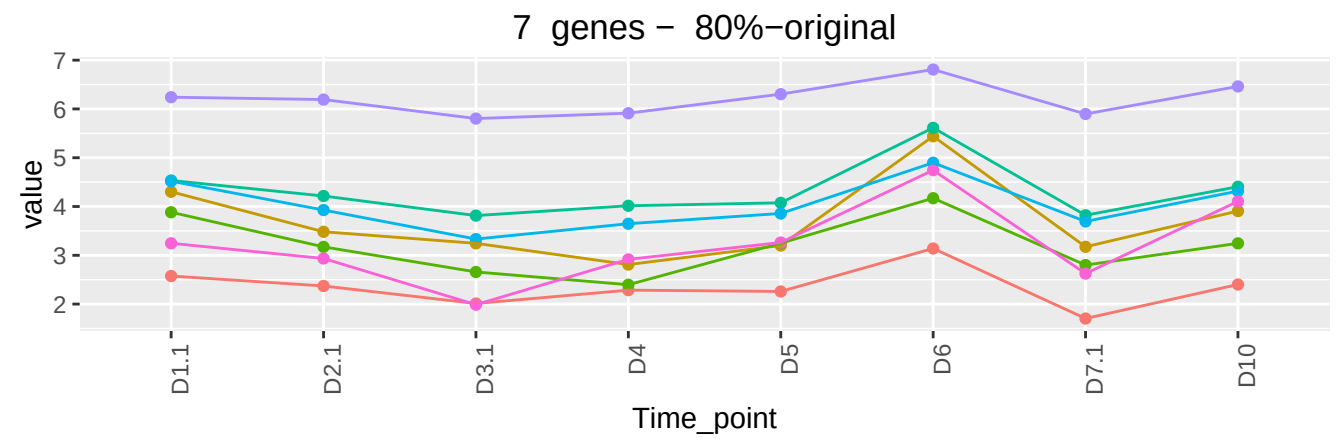
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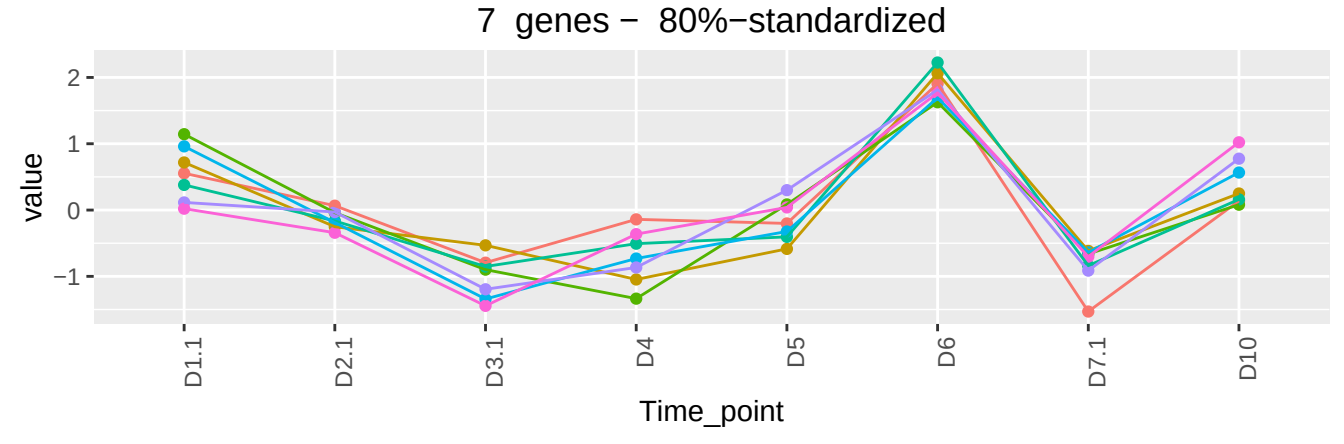
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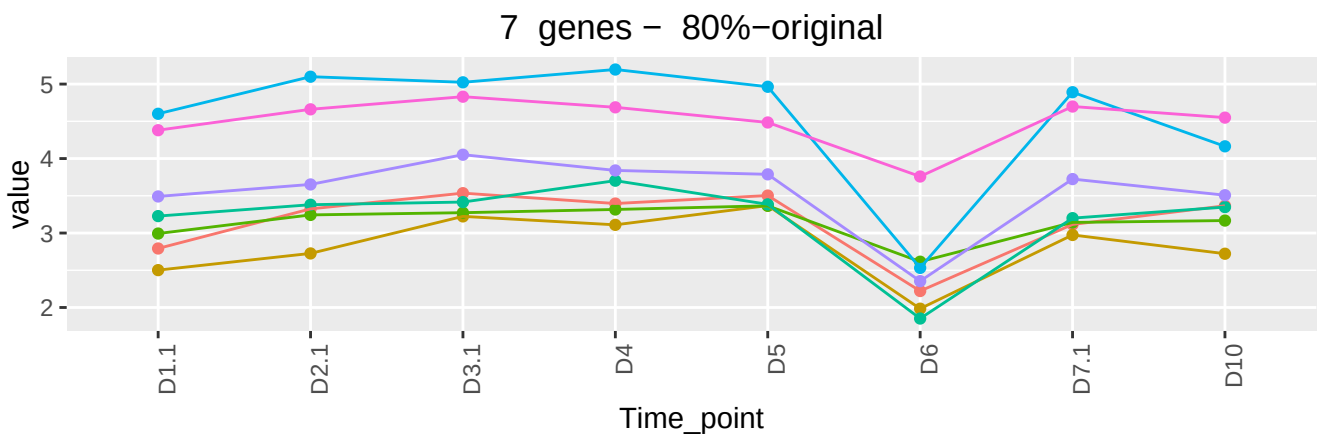
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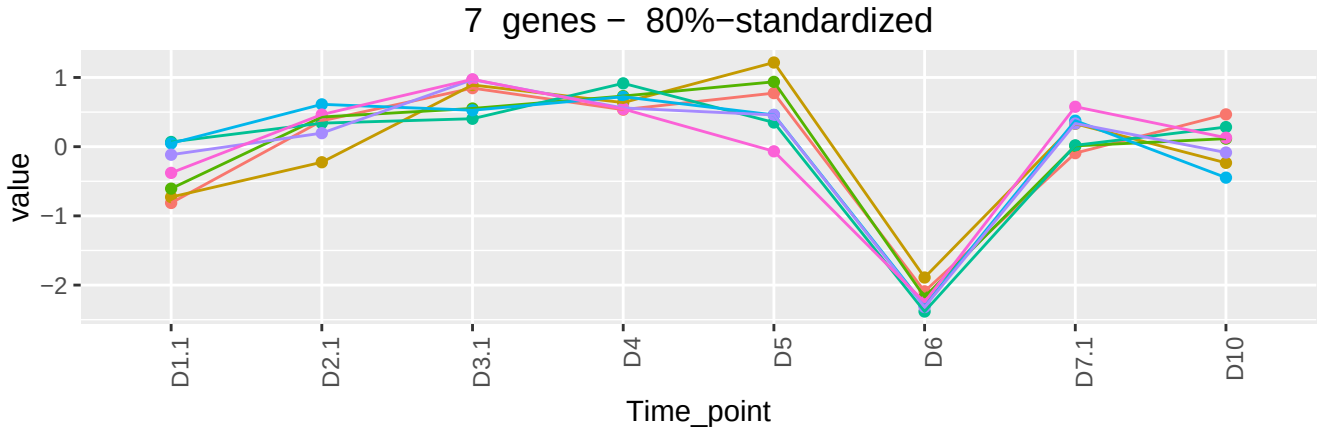
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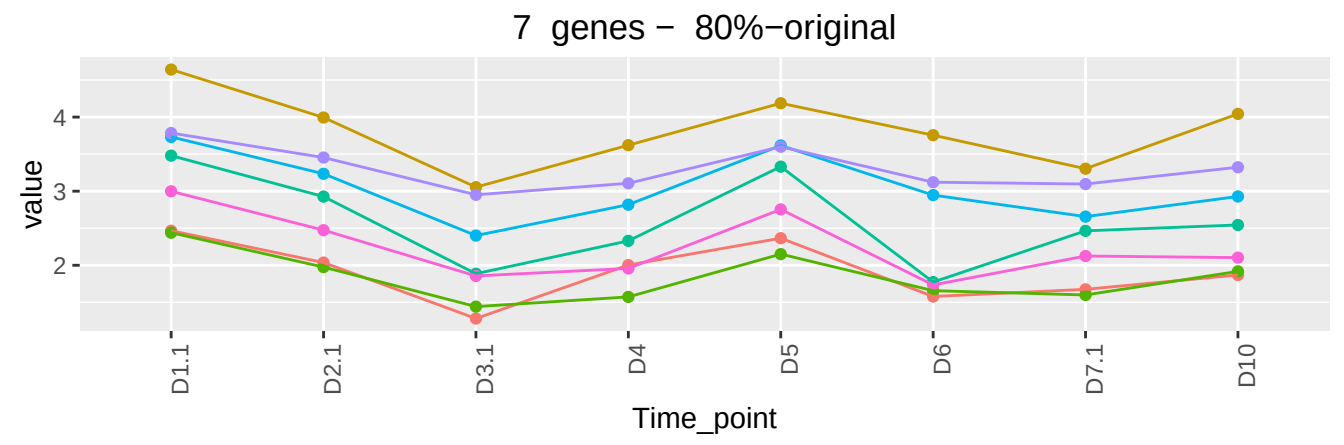
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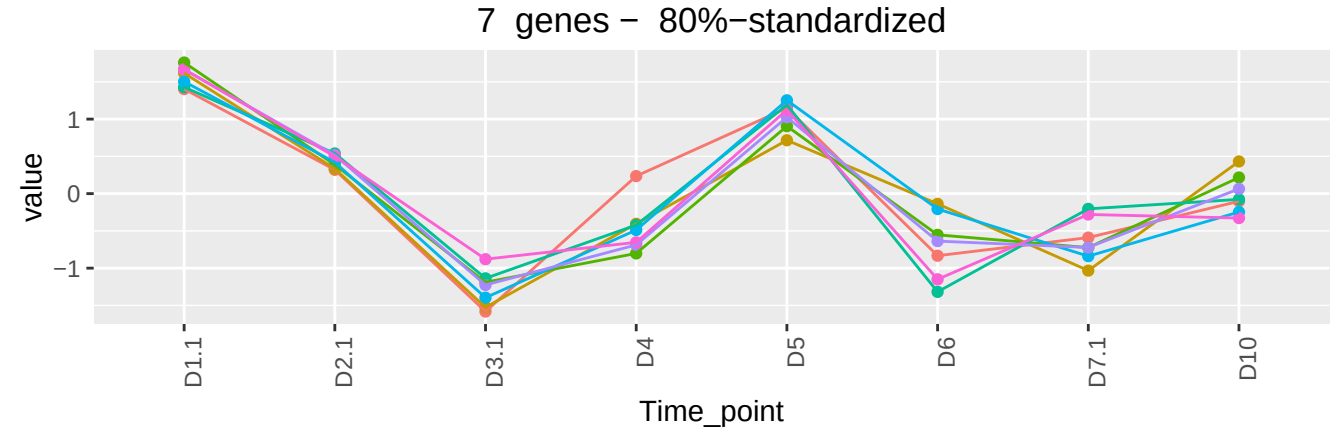
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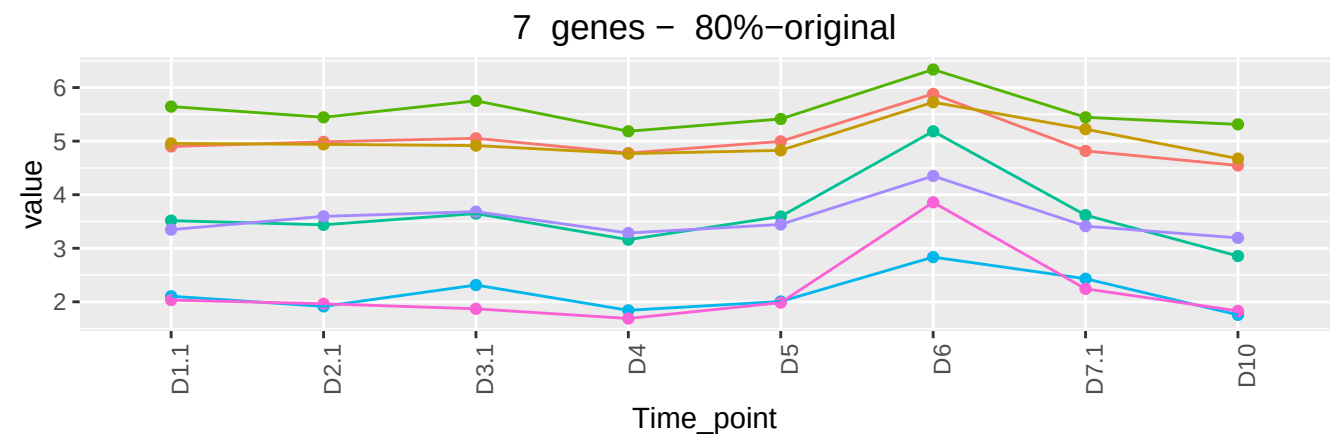
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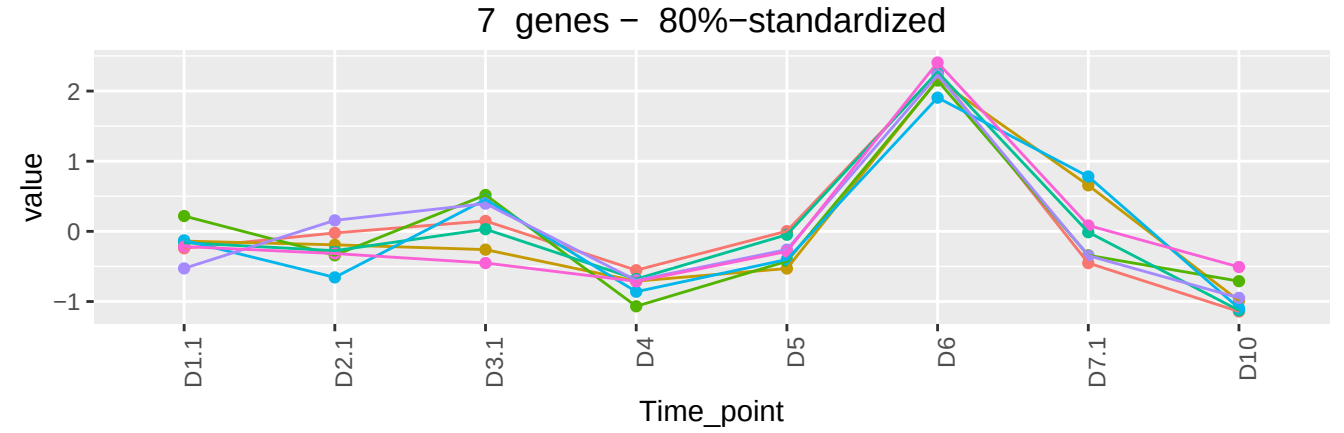
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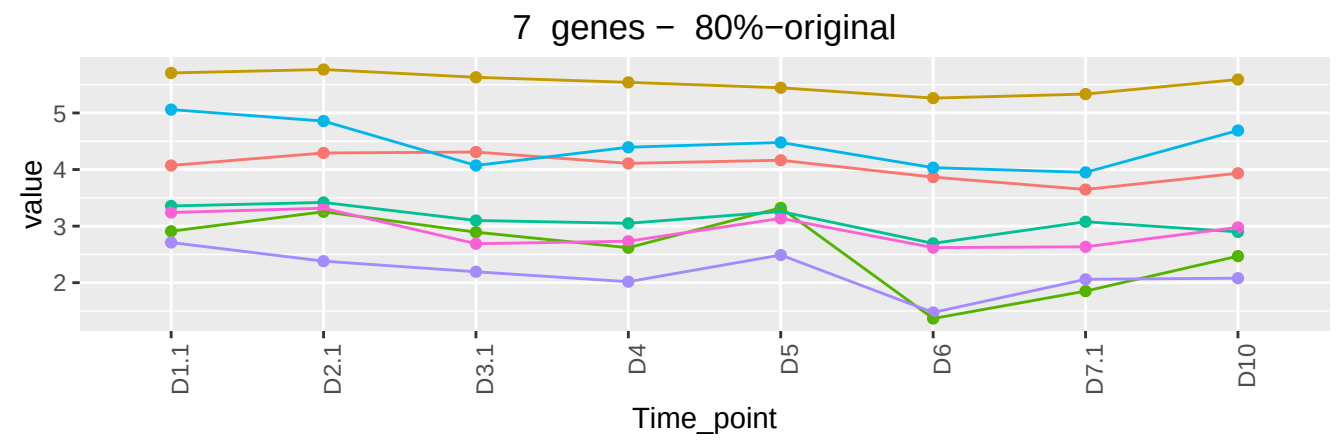
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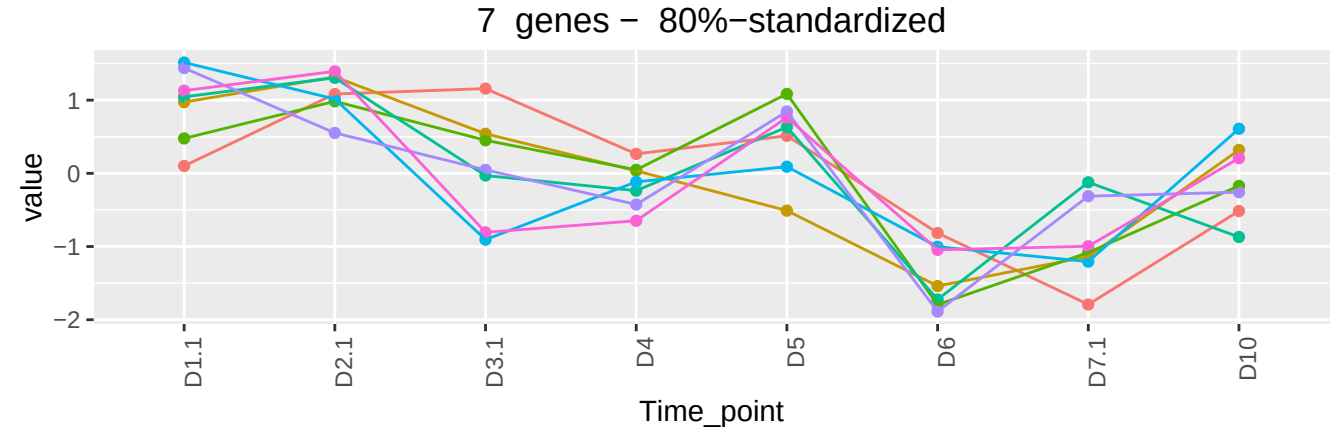
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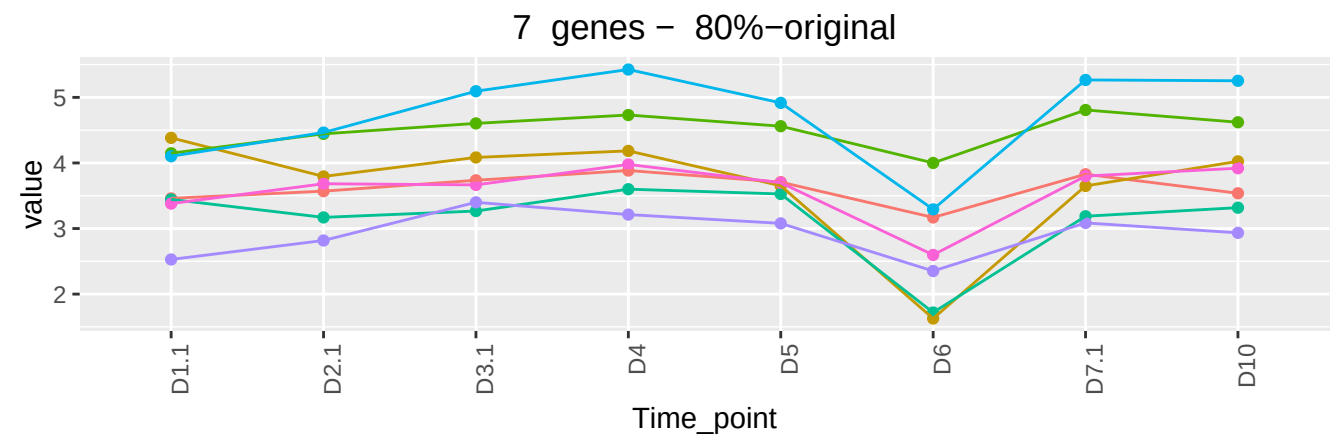
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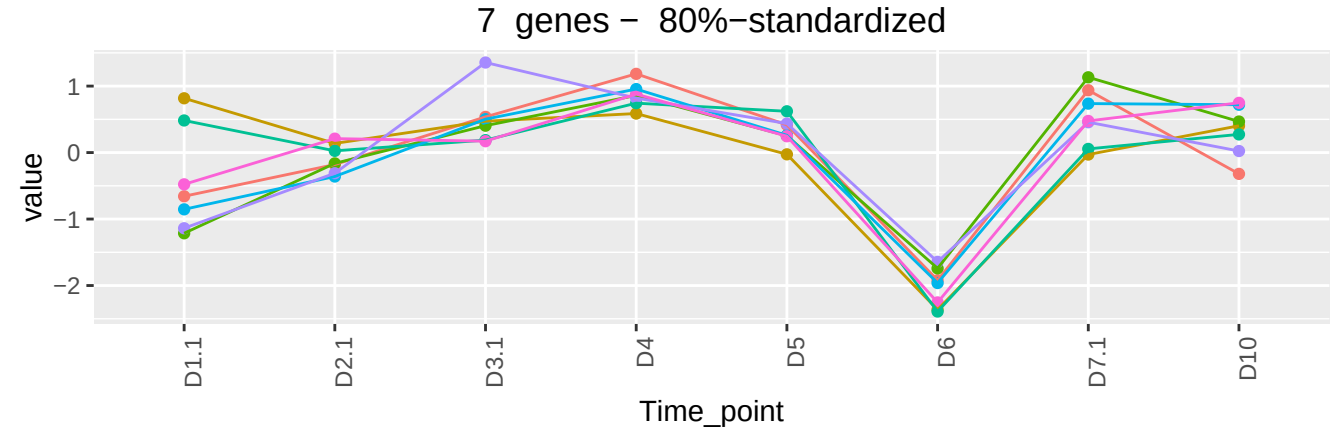
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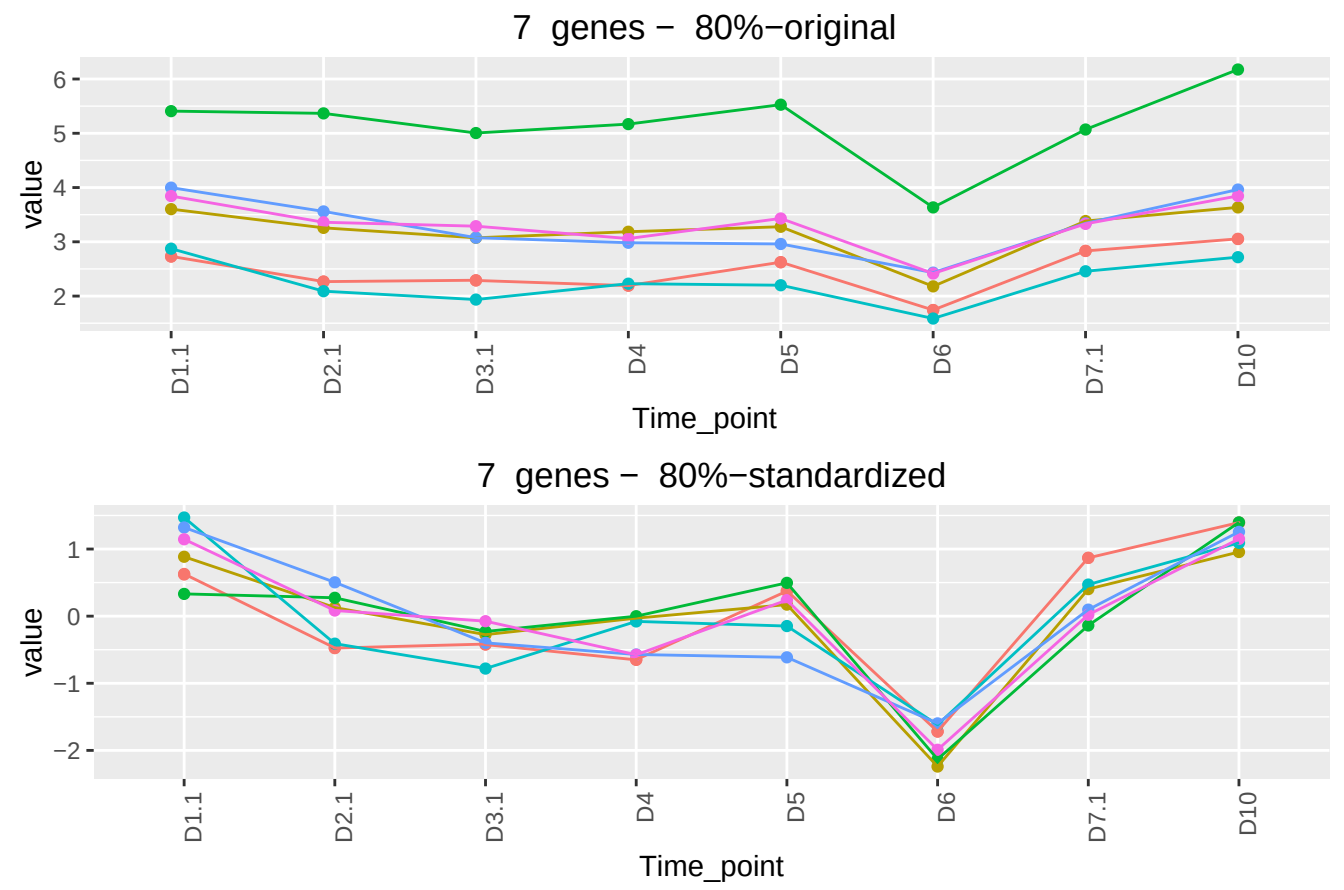
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Time_point

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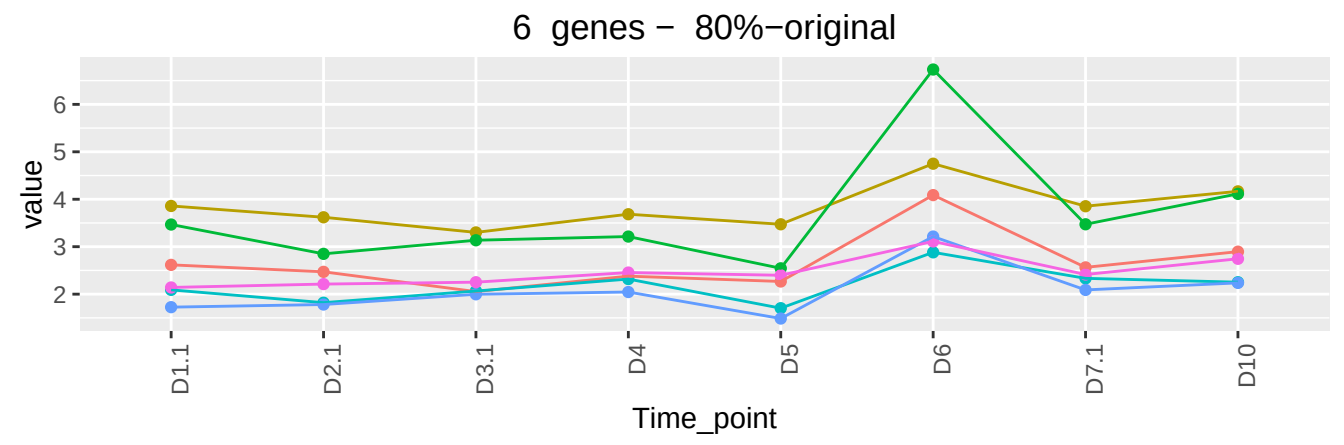
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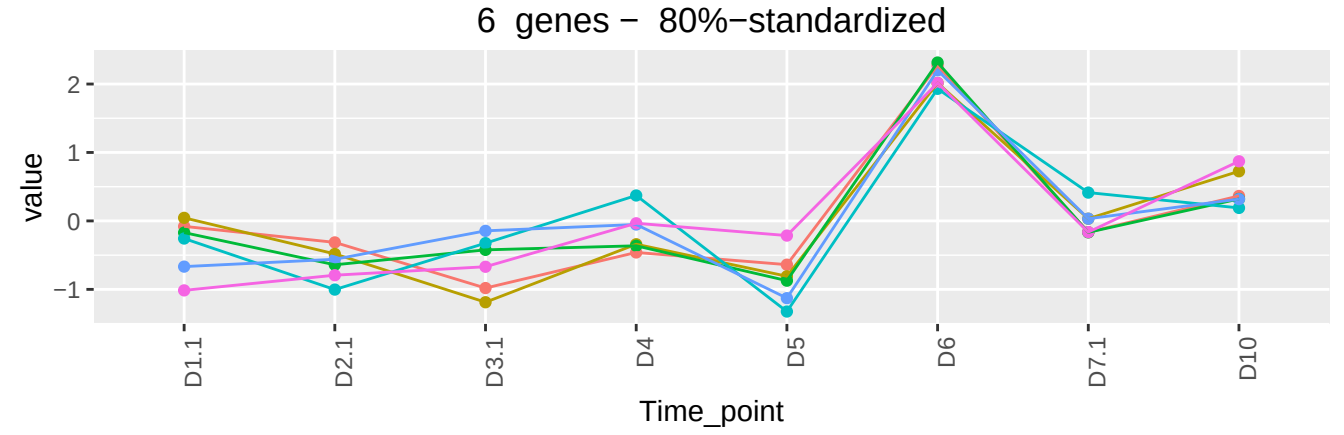
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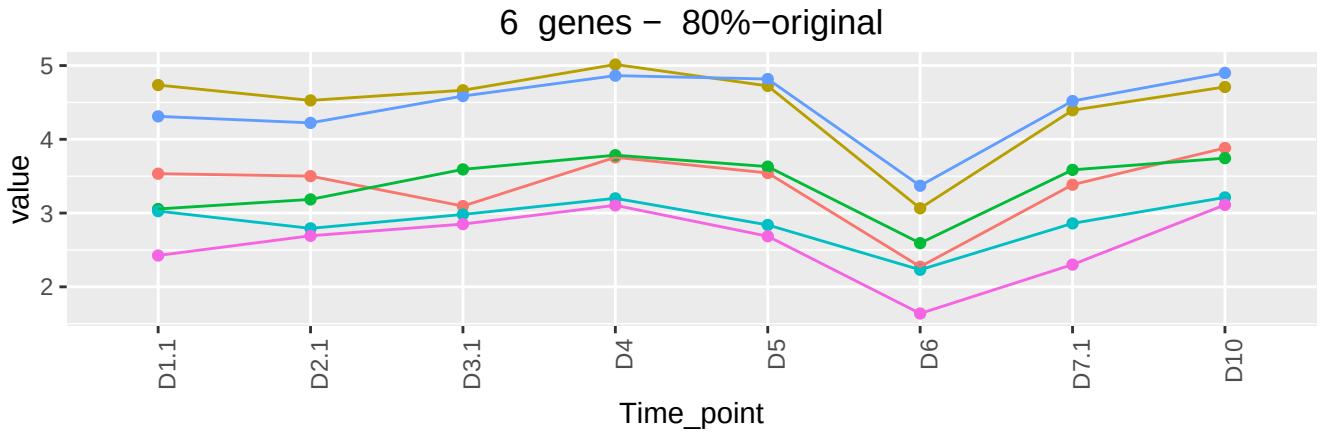
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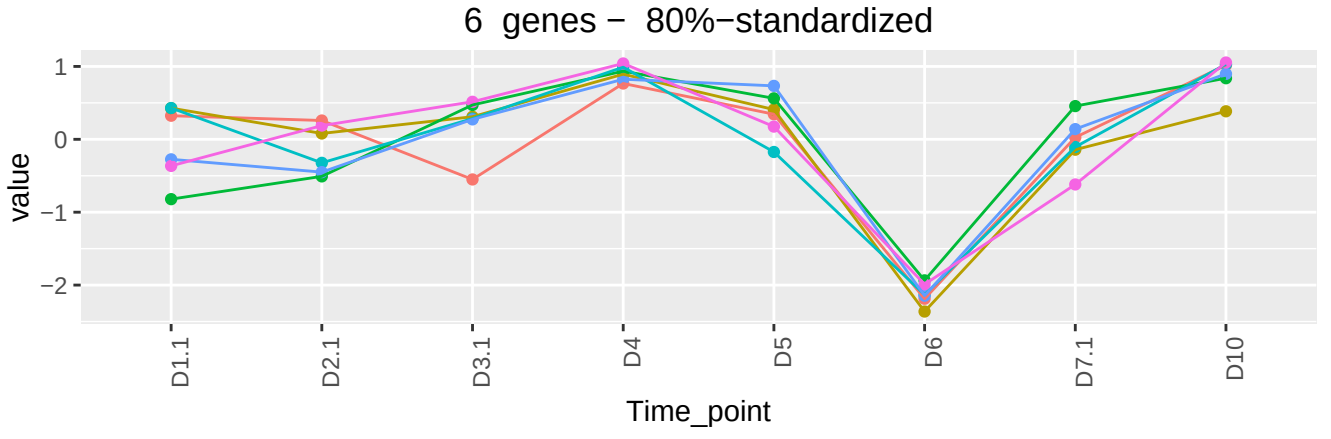
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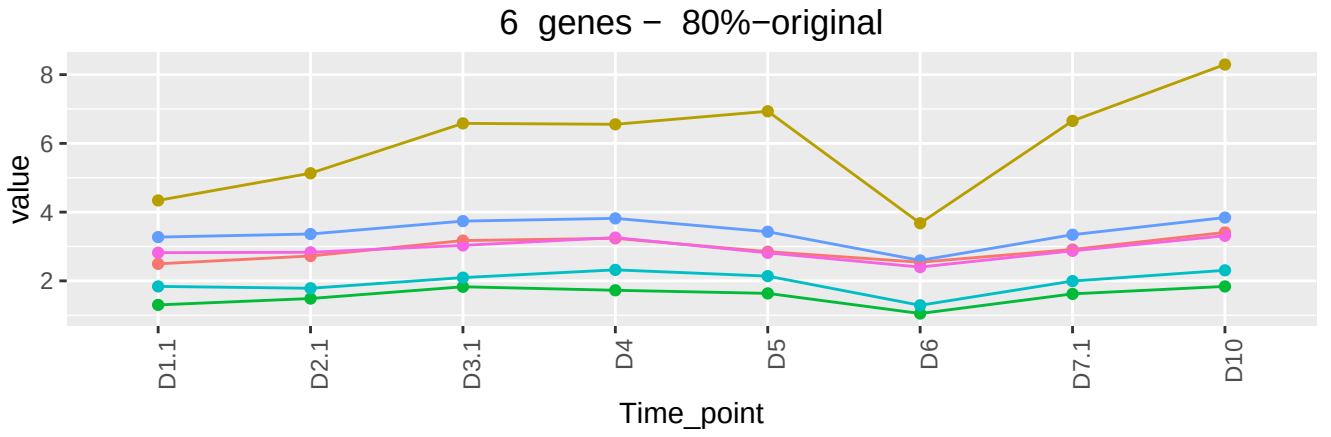
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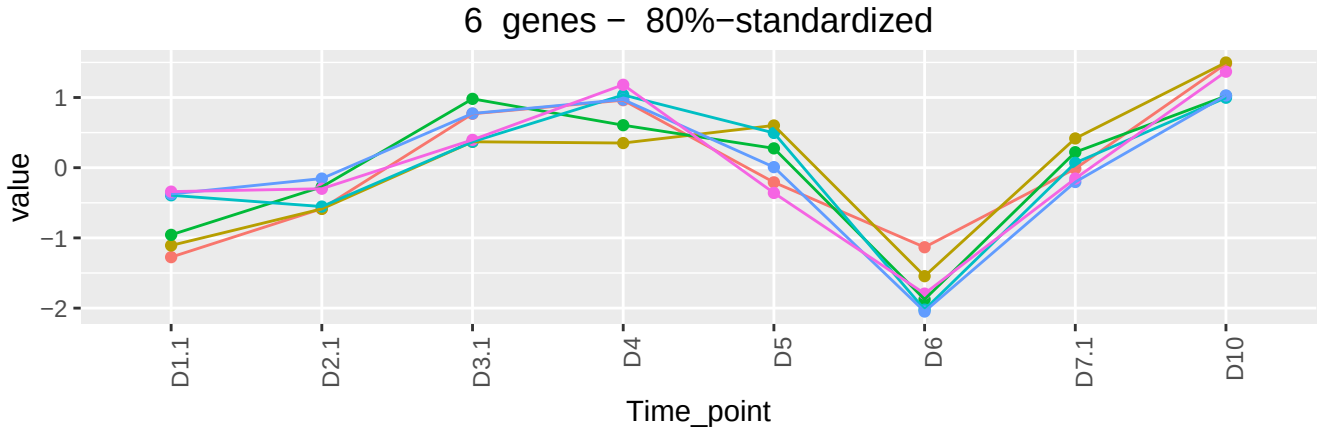
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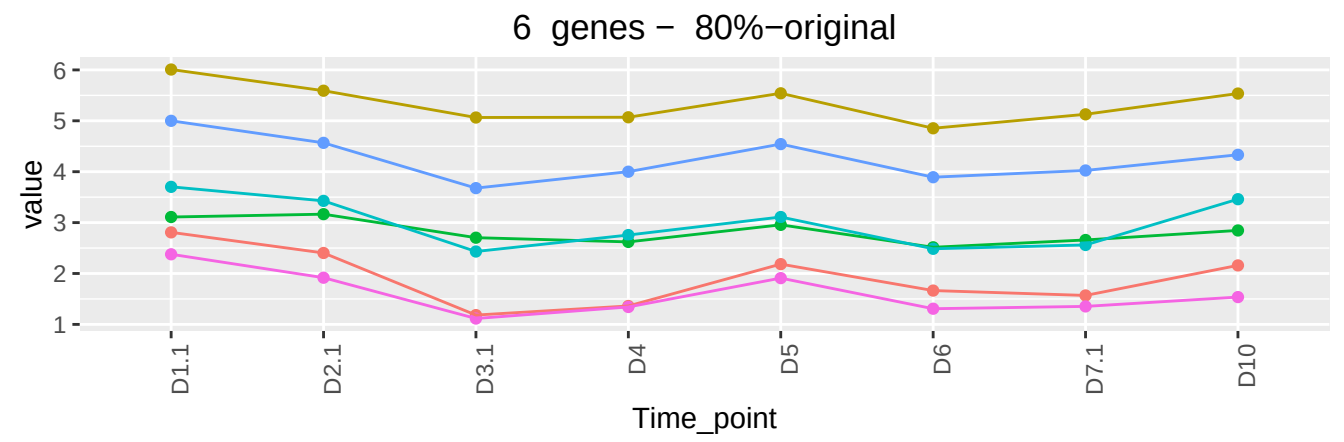
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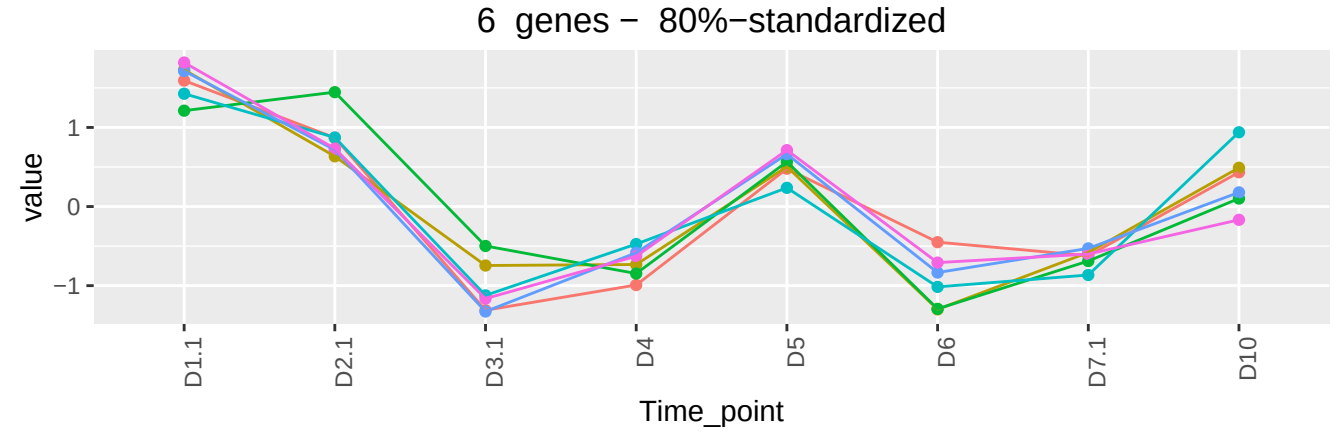
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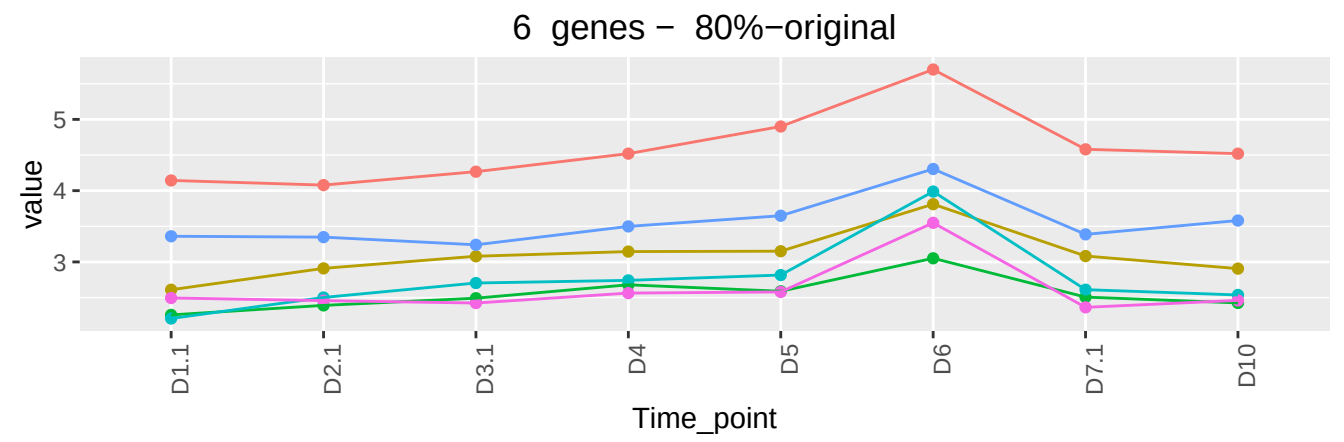
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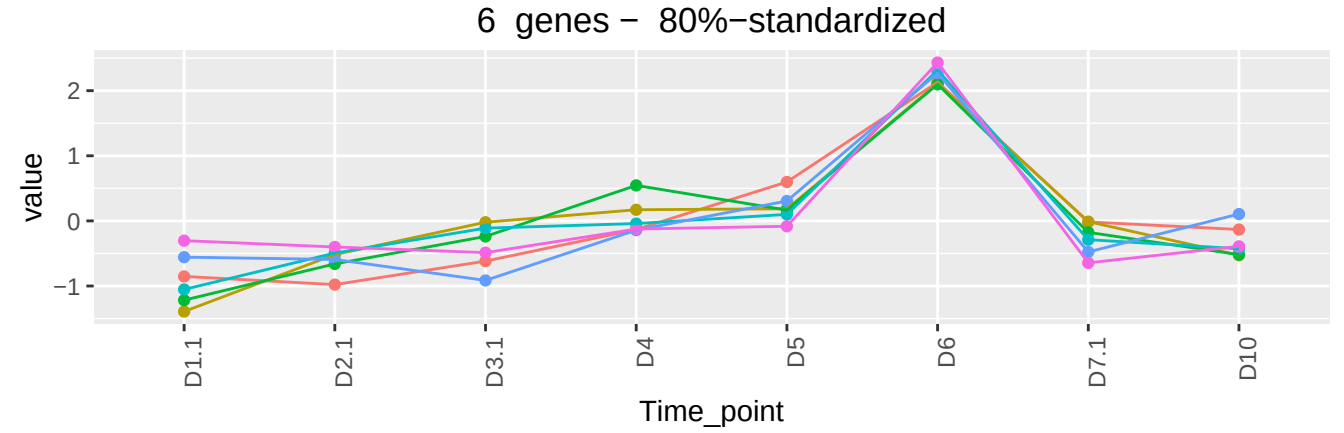
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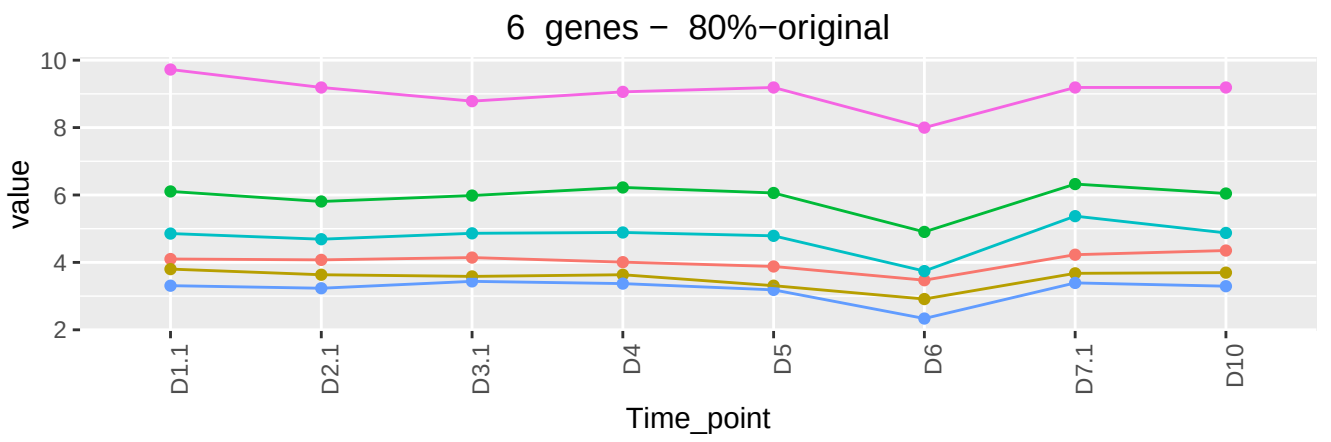
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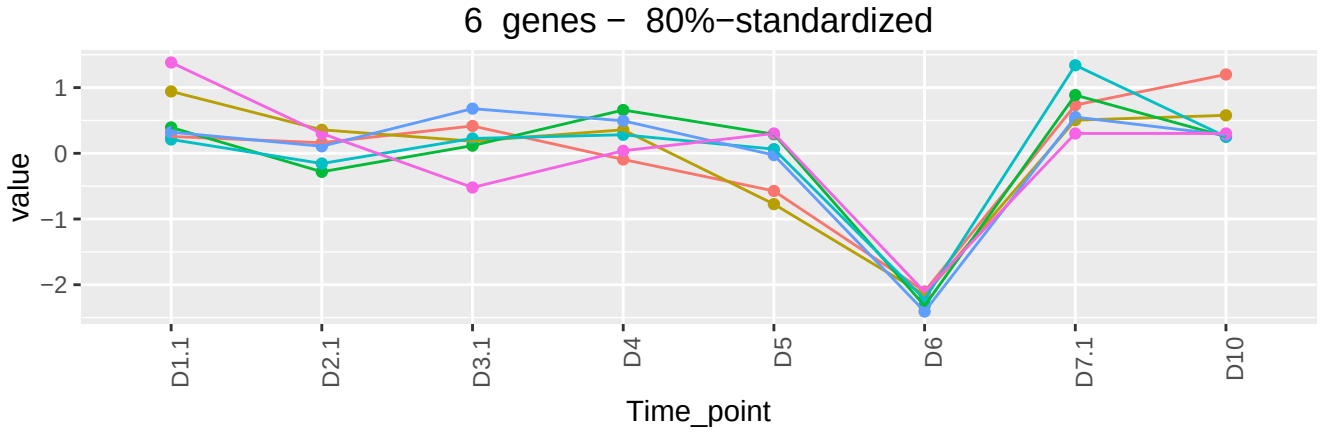
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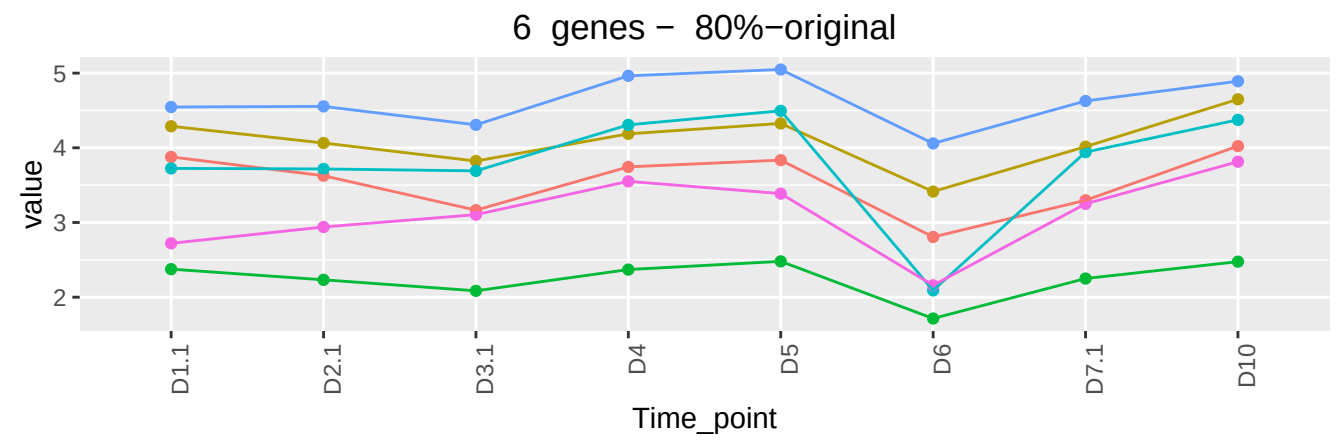
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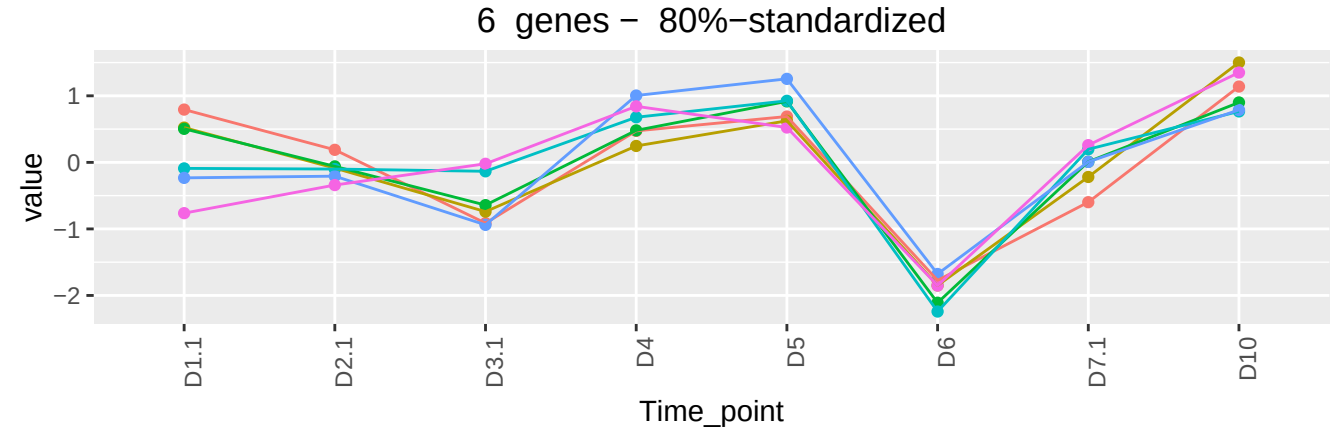
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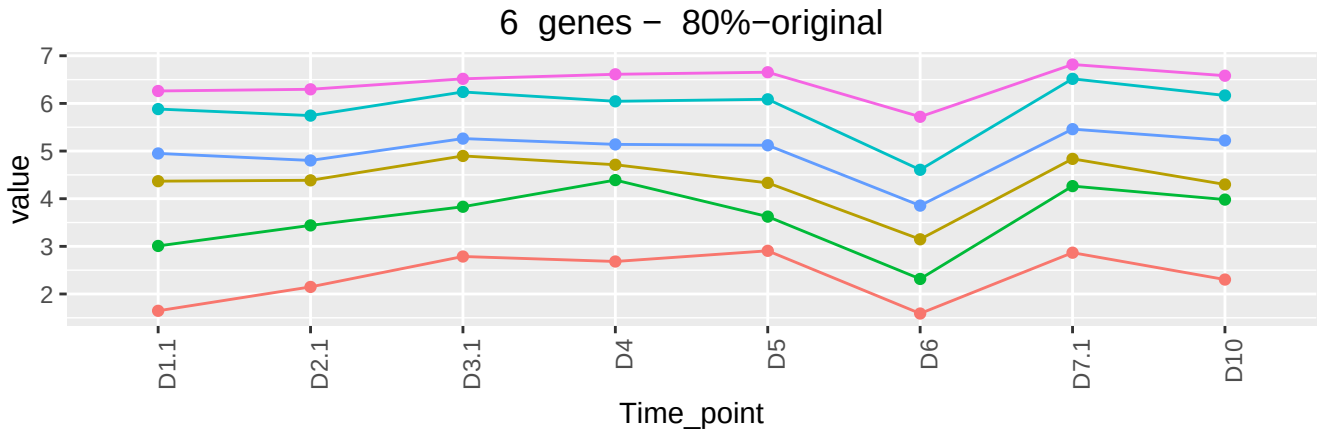
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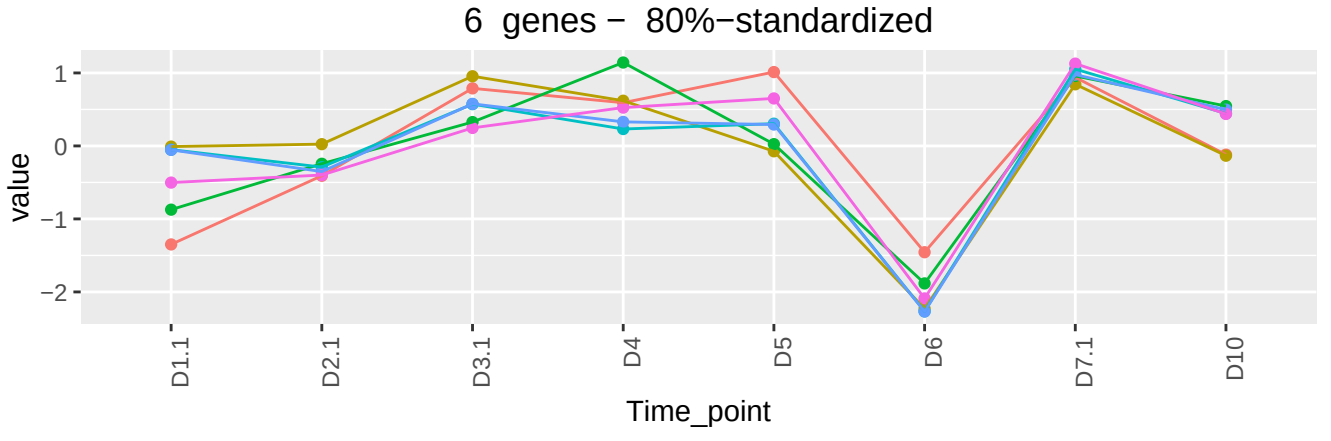
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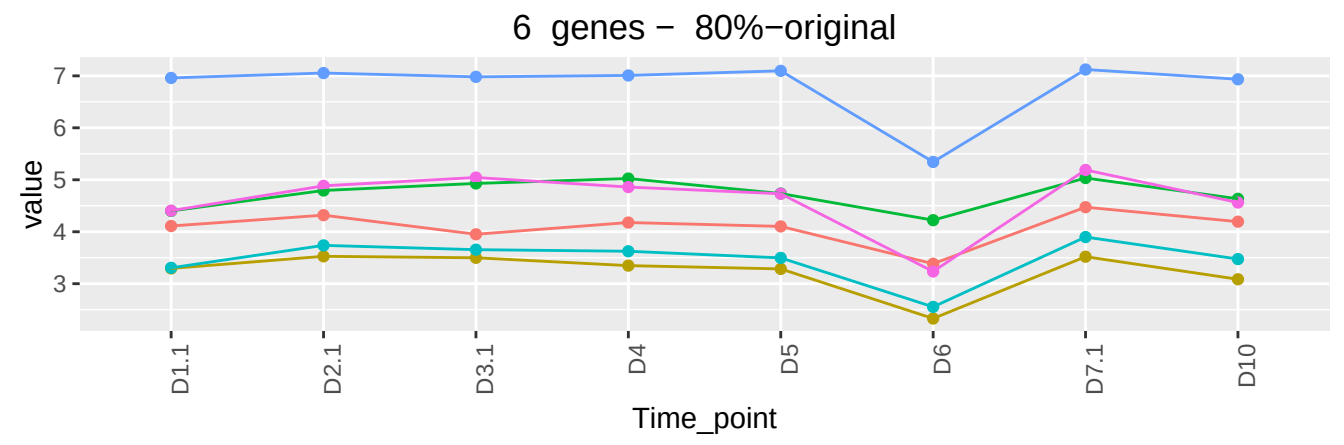
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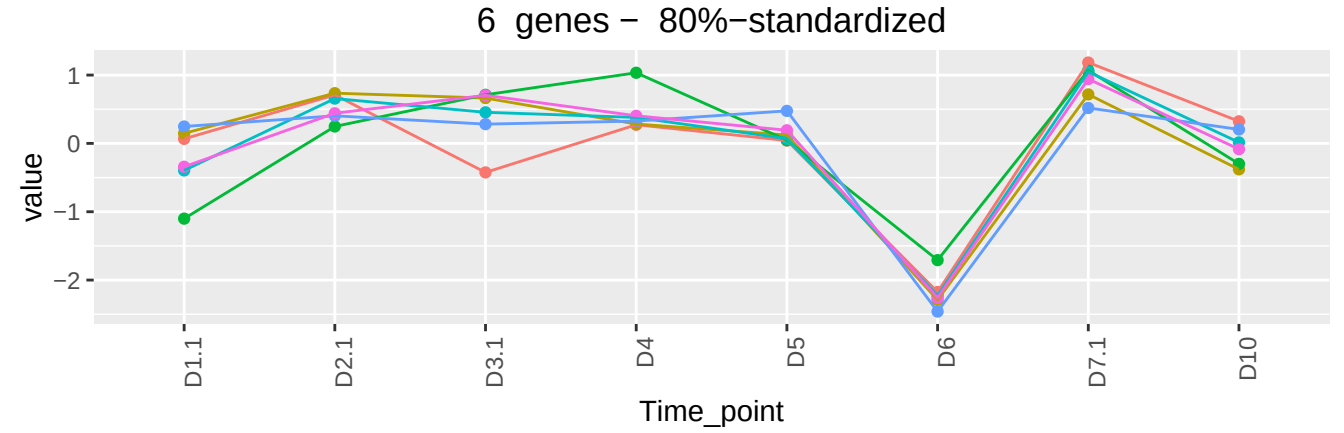
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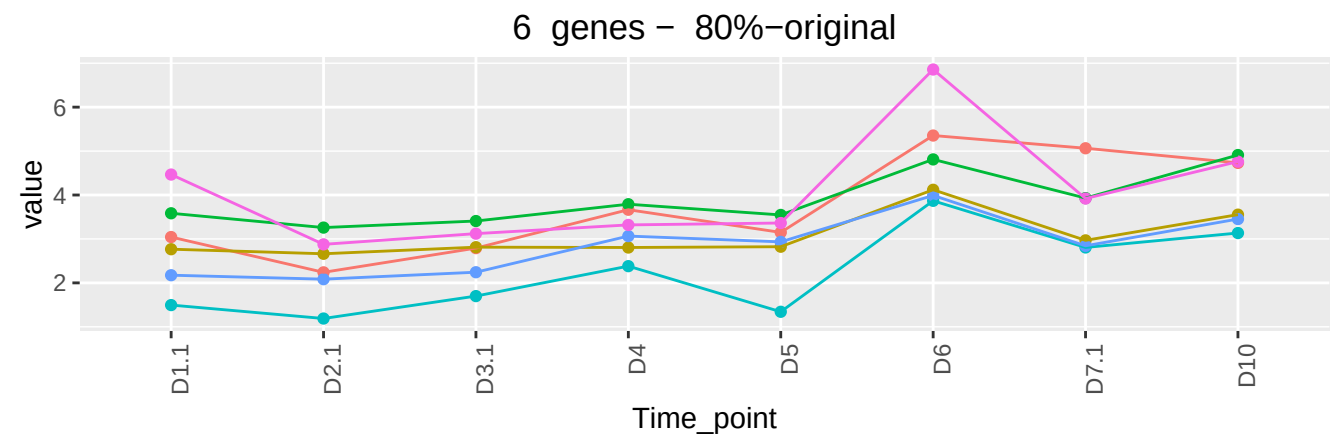
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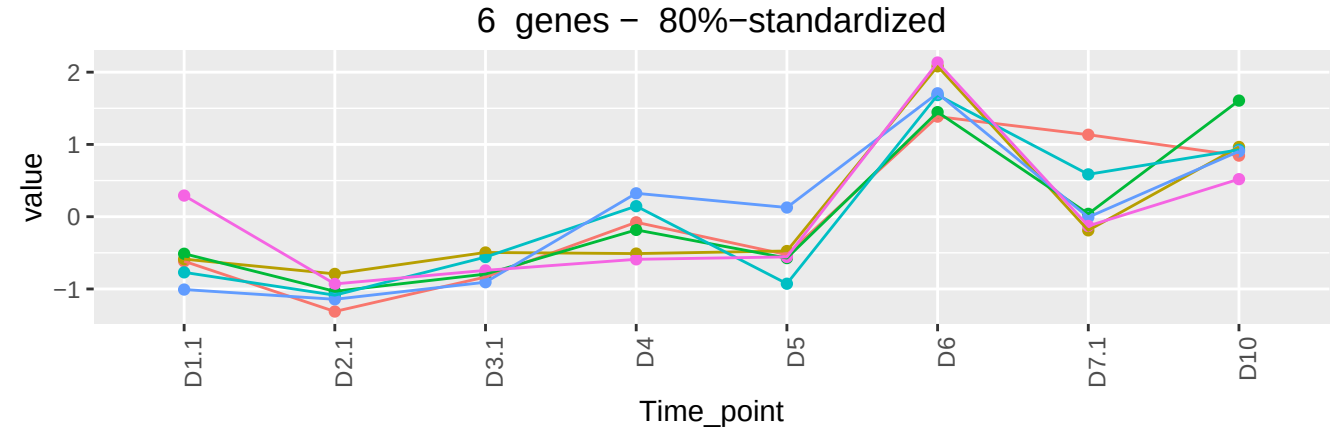
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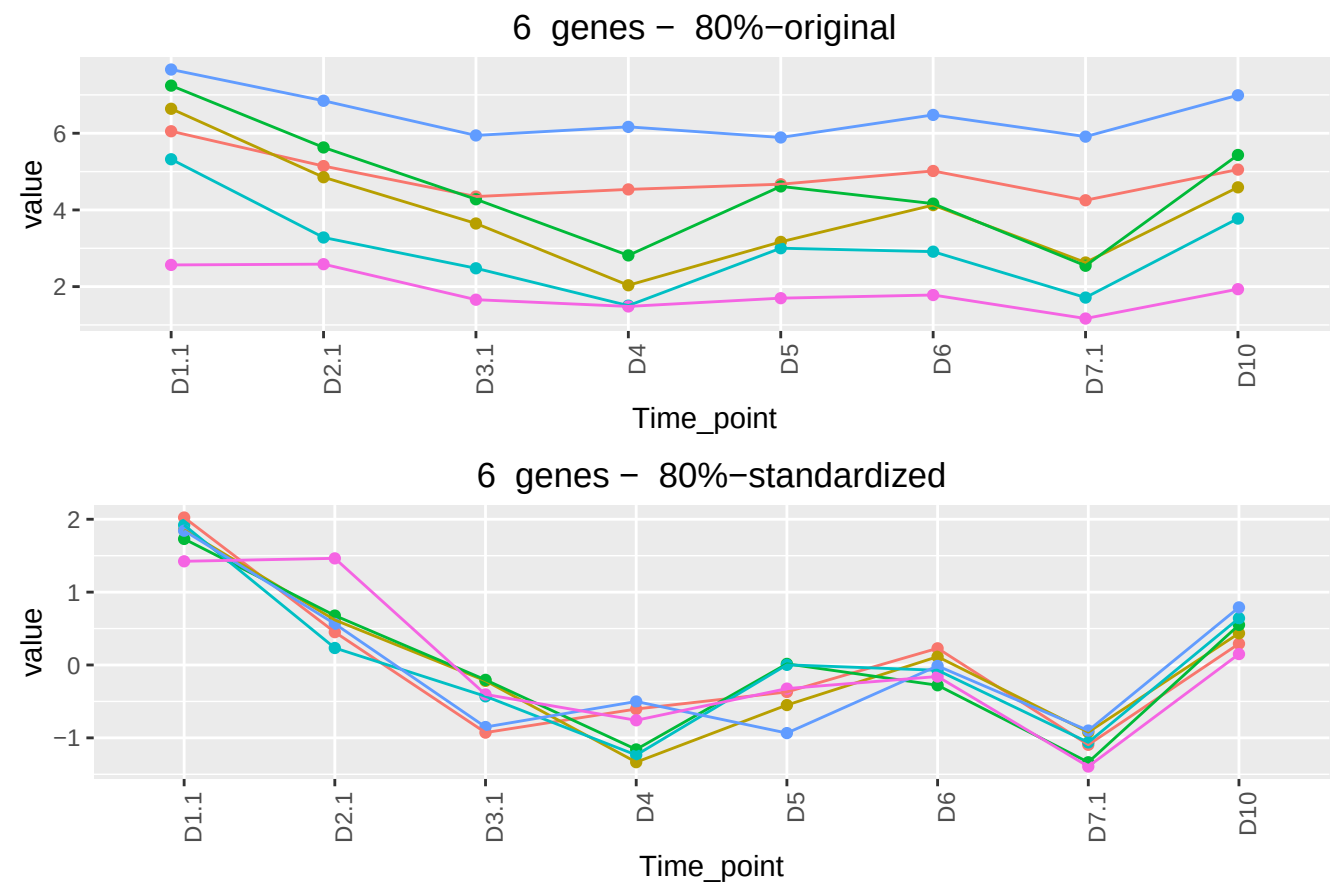
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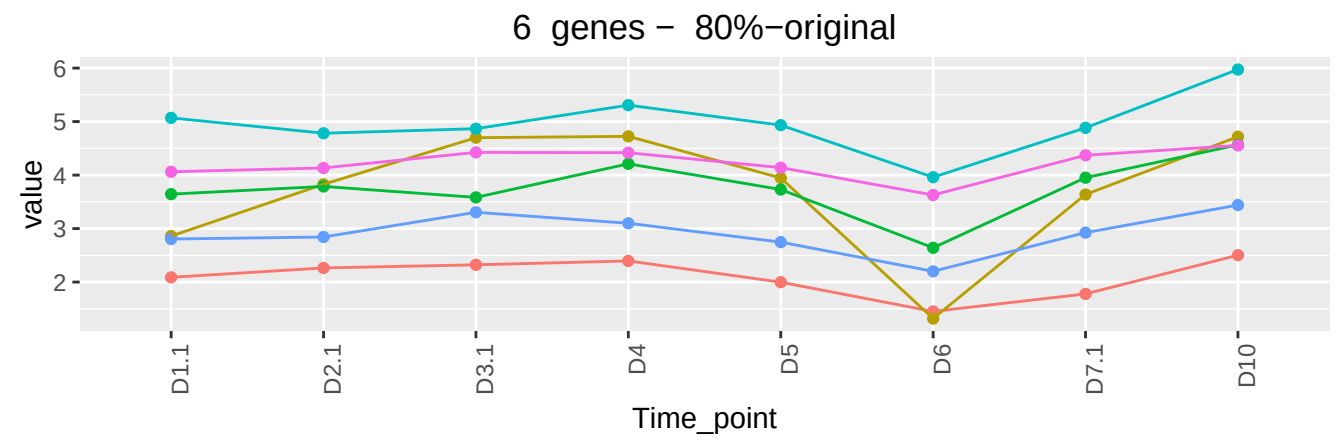
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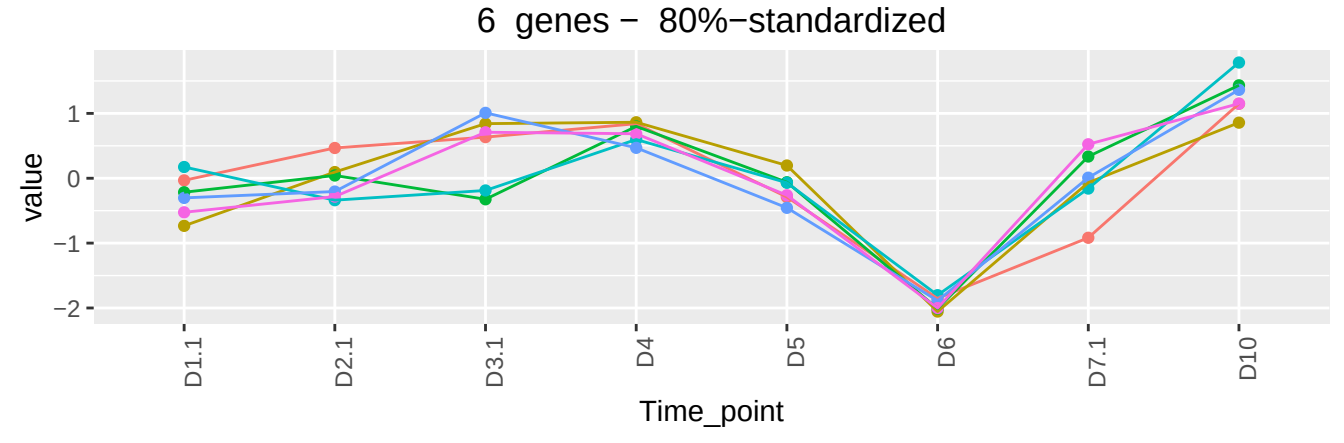
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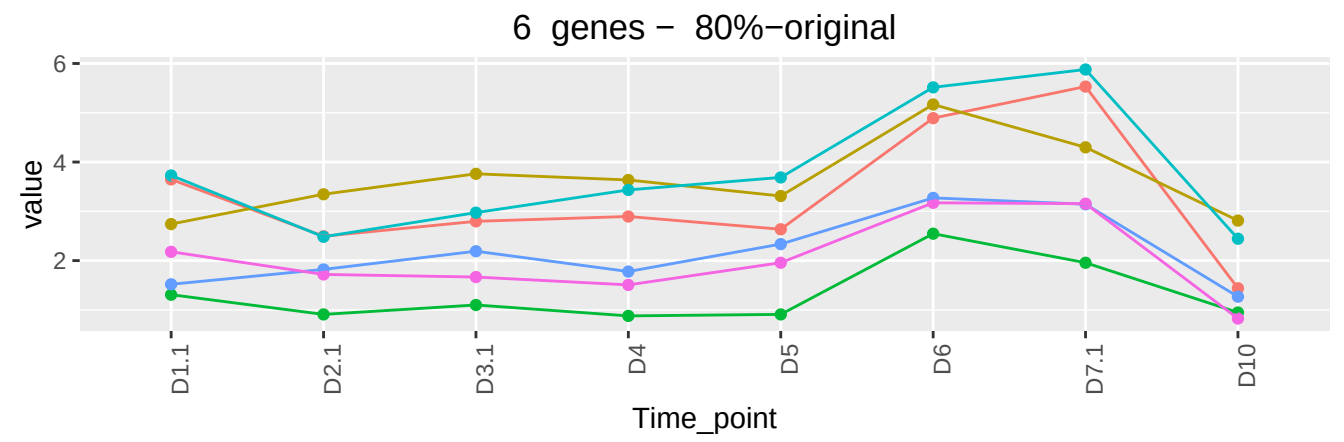
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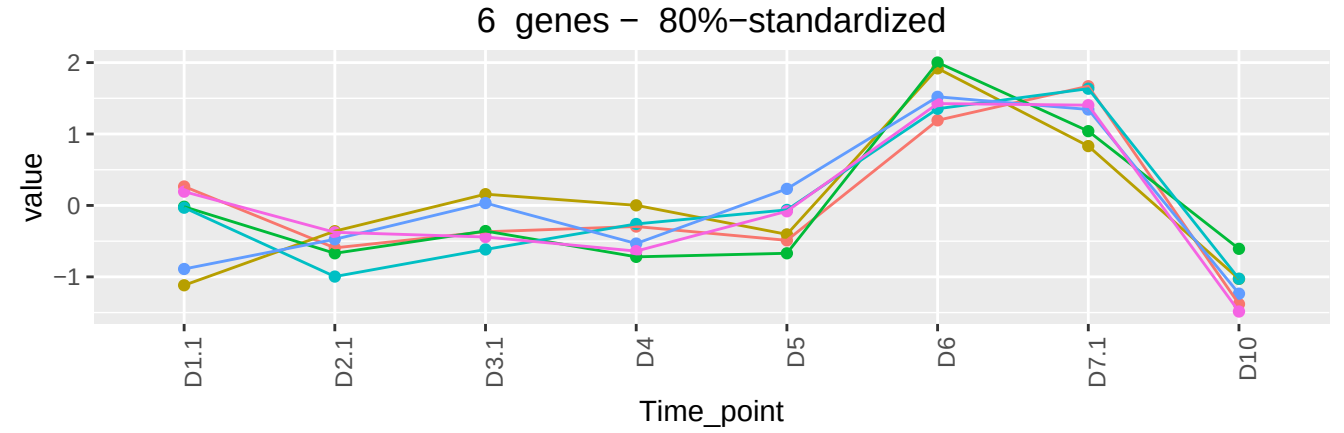
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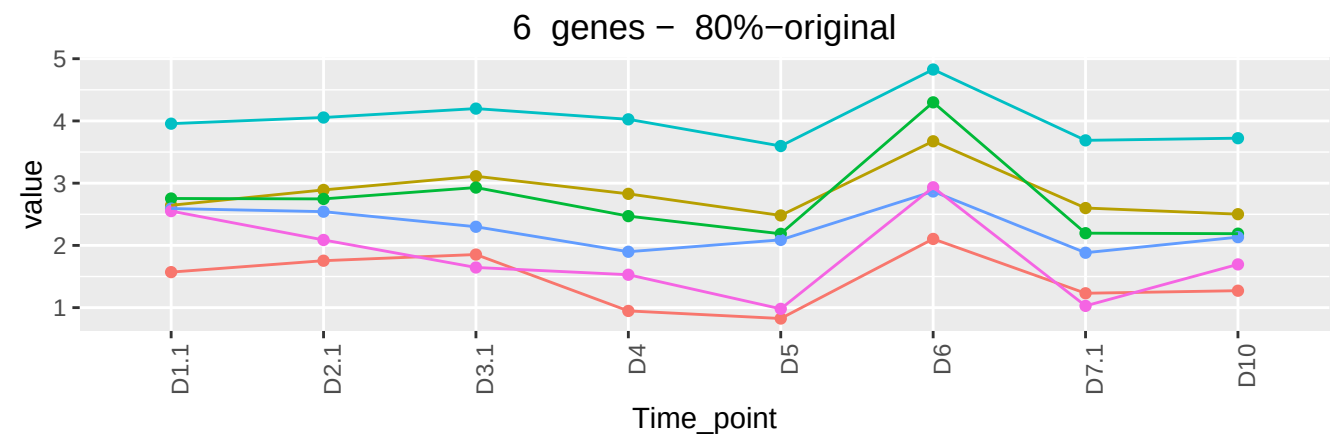
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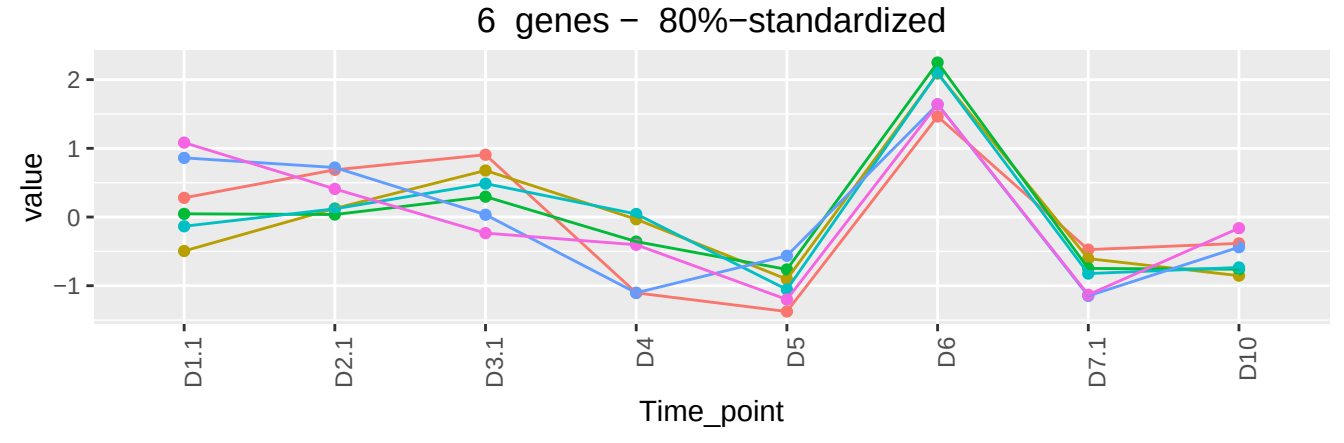
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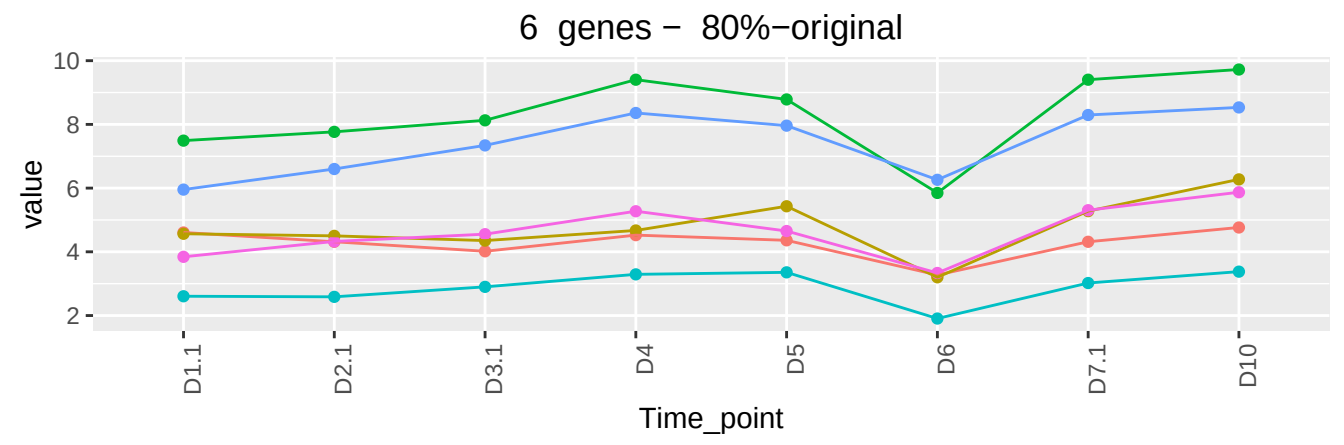
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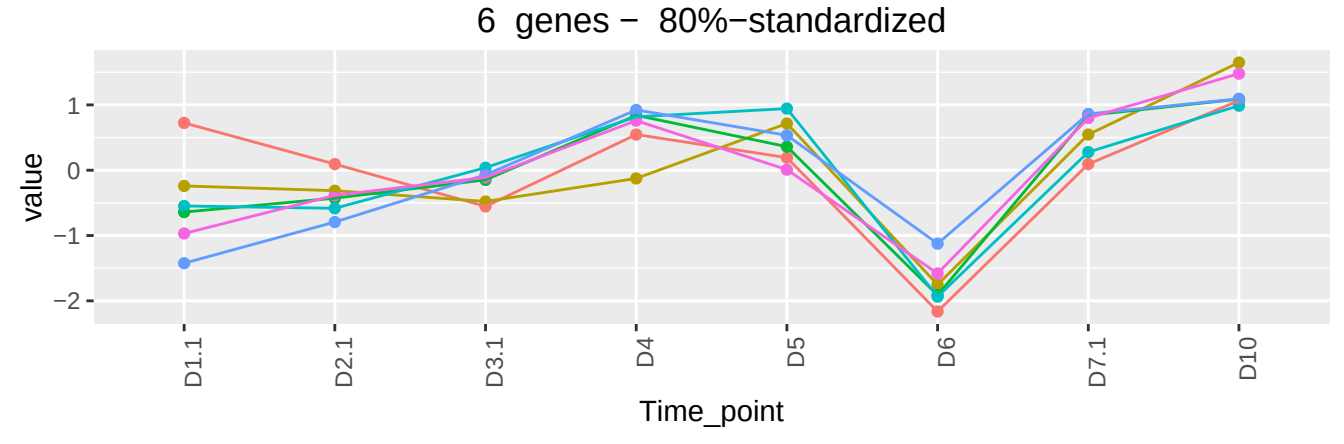
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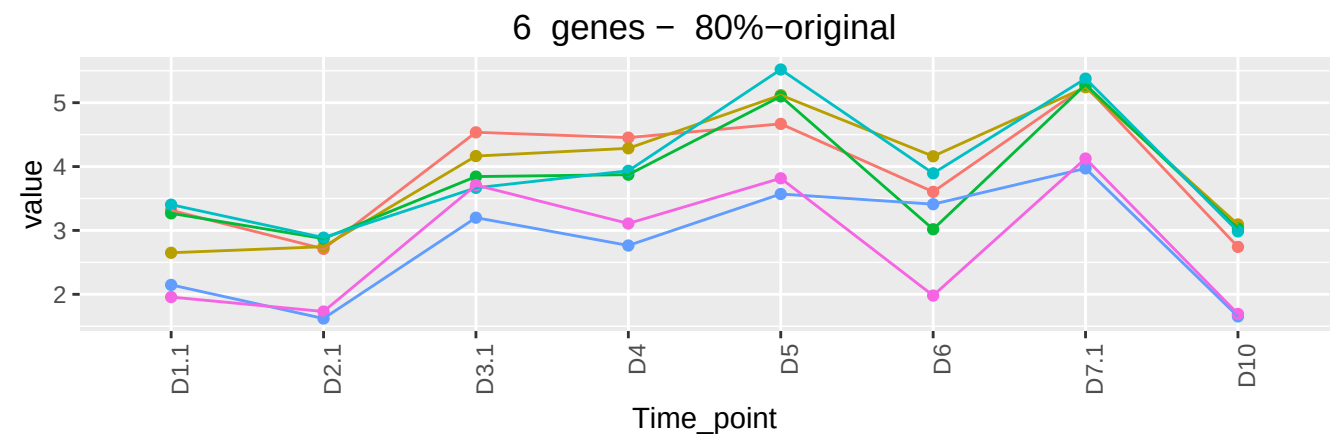
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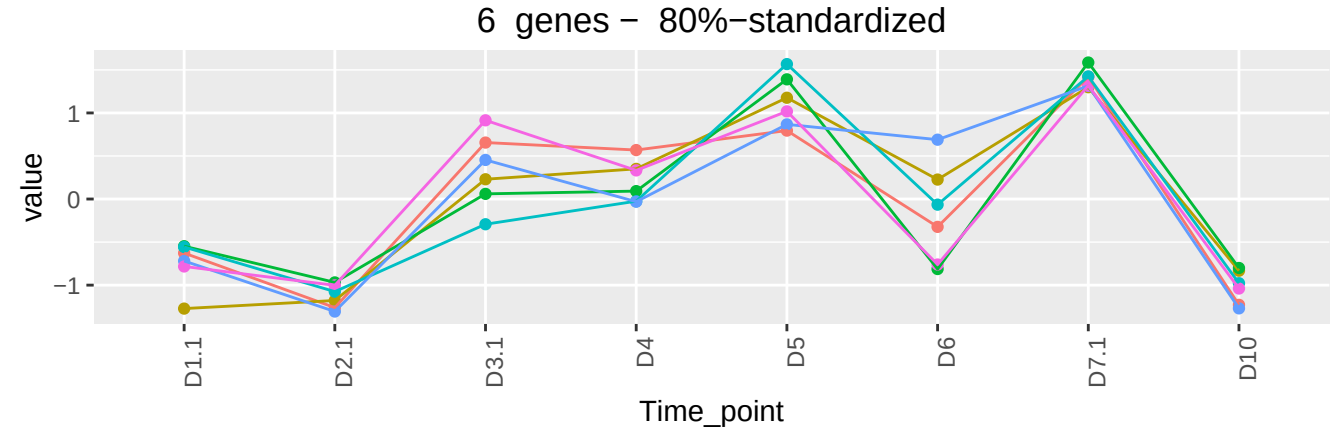
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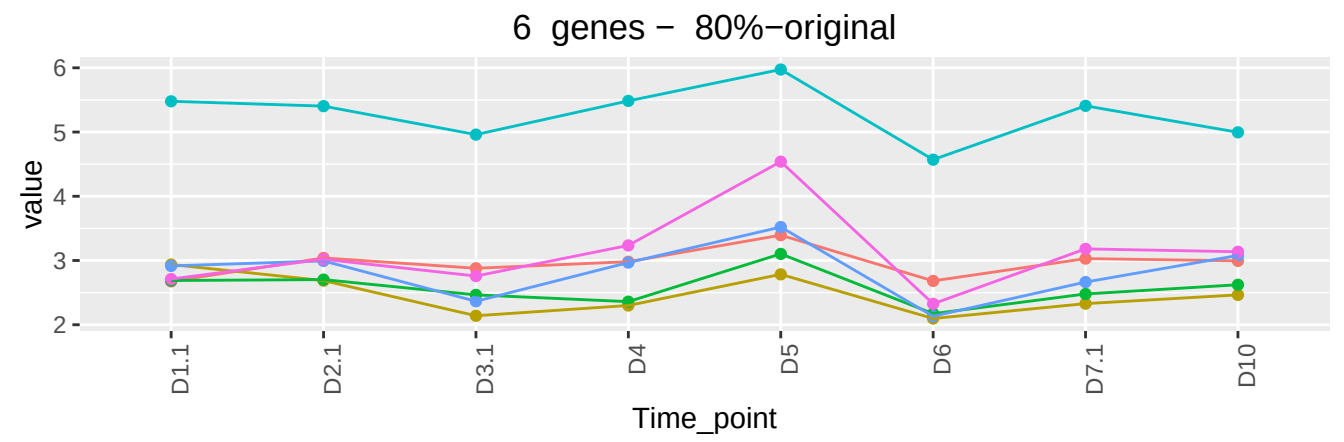
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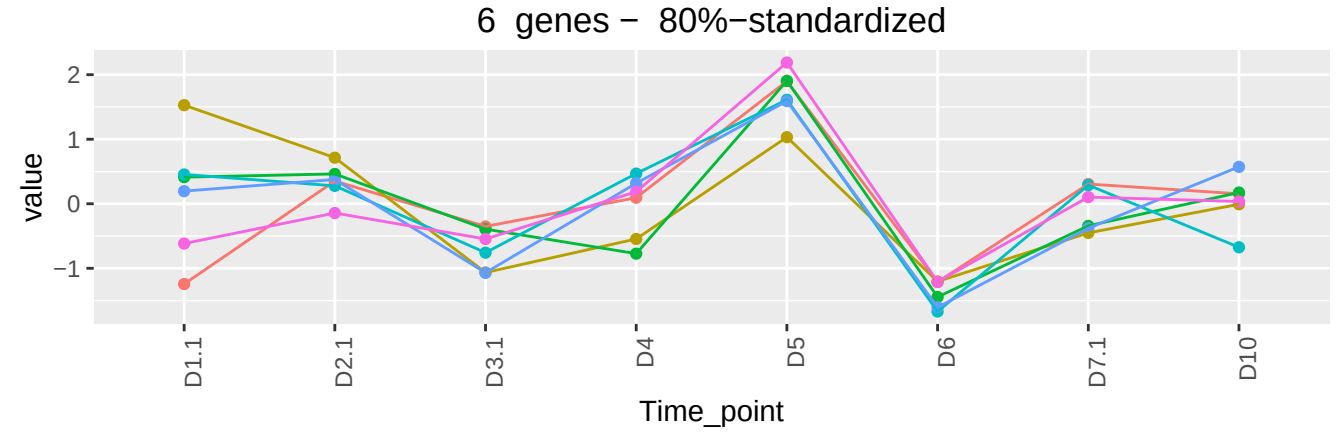
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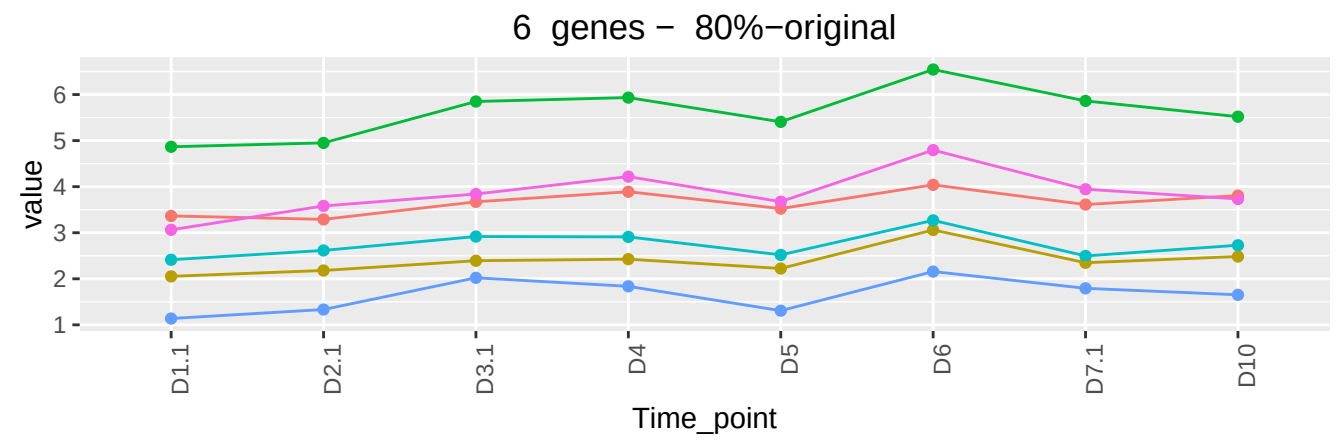
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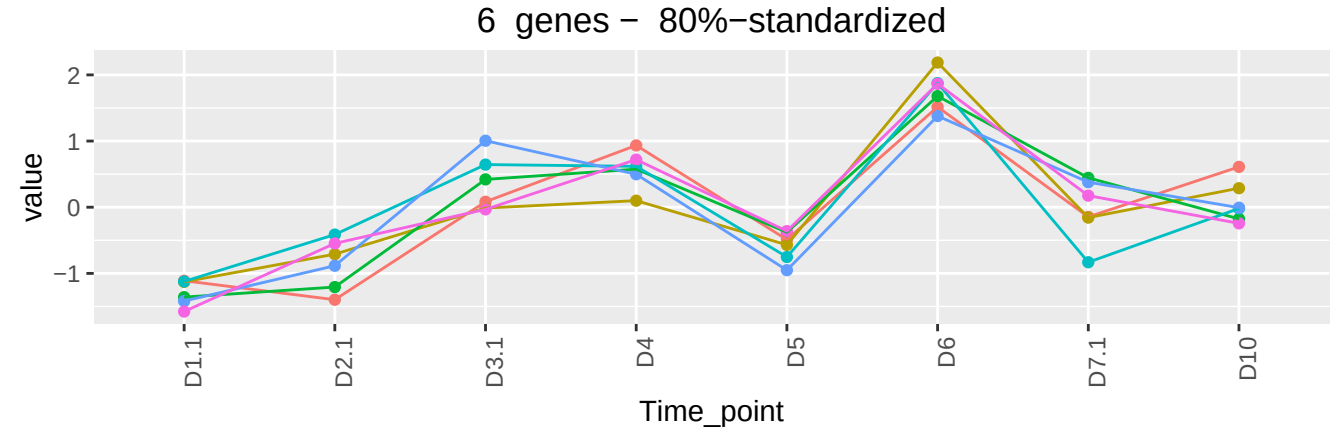
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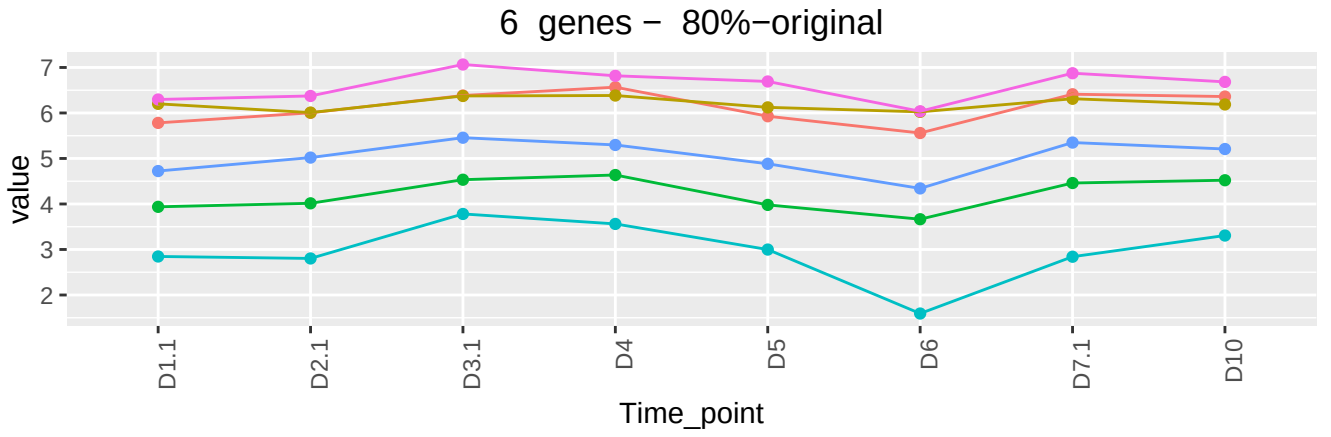
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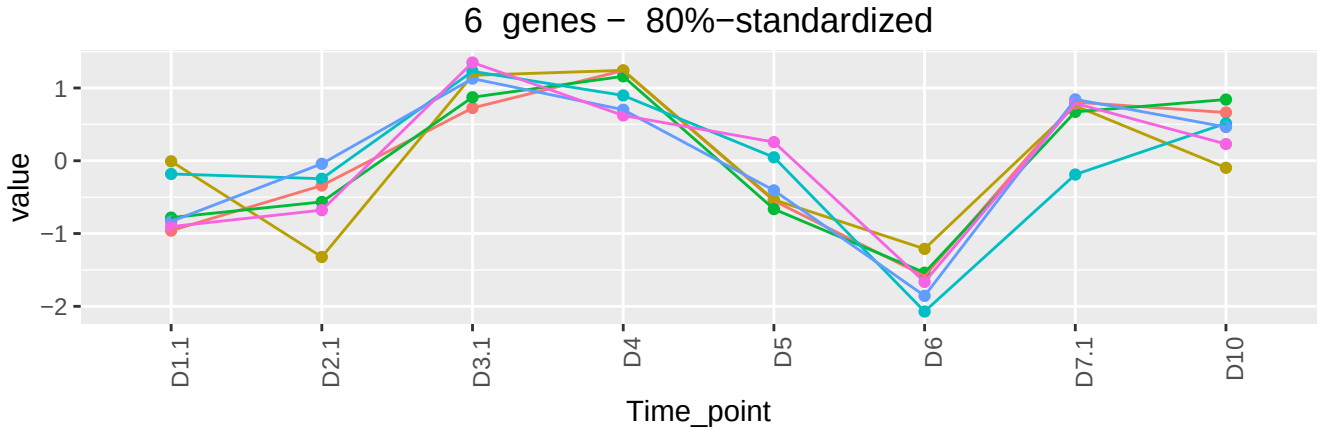
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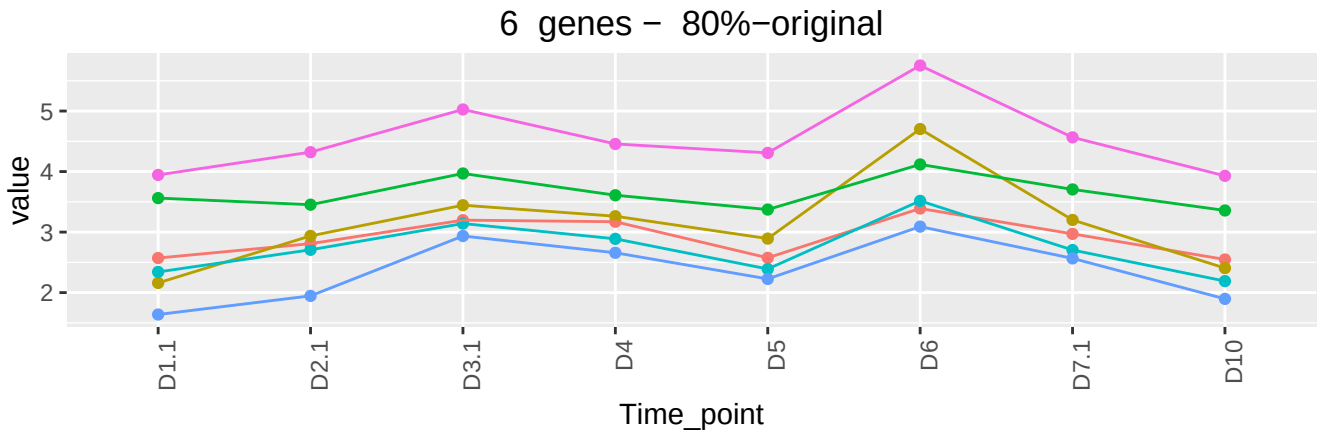
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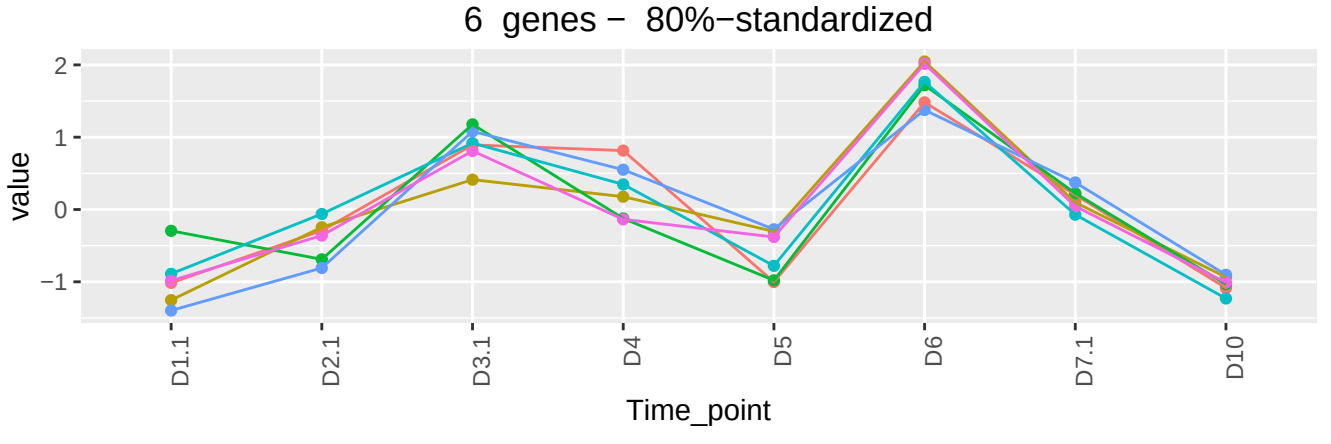
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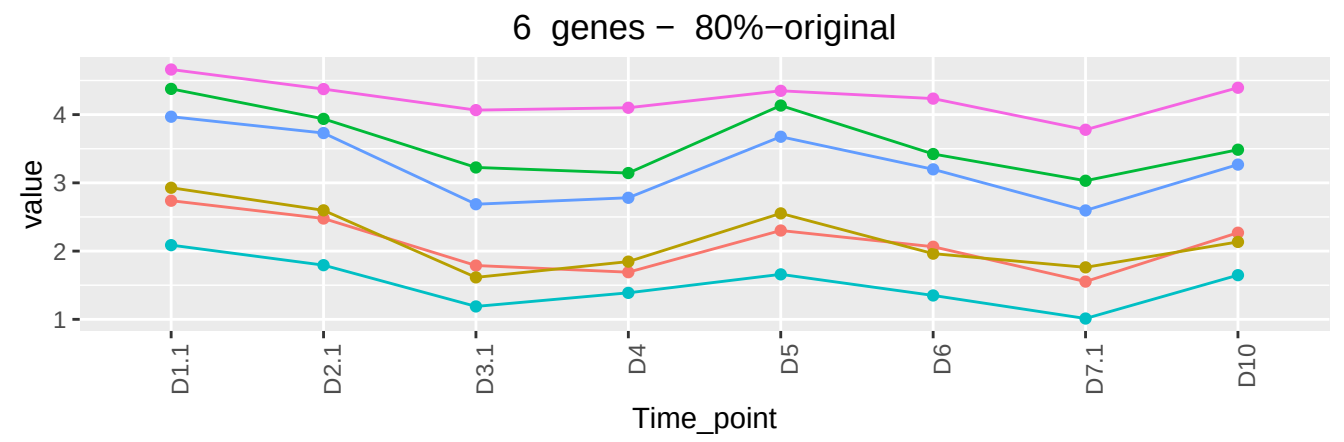
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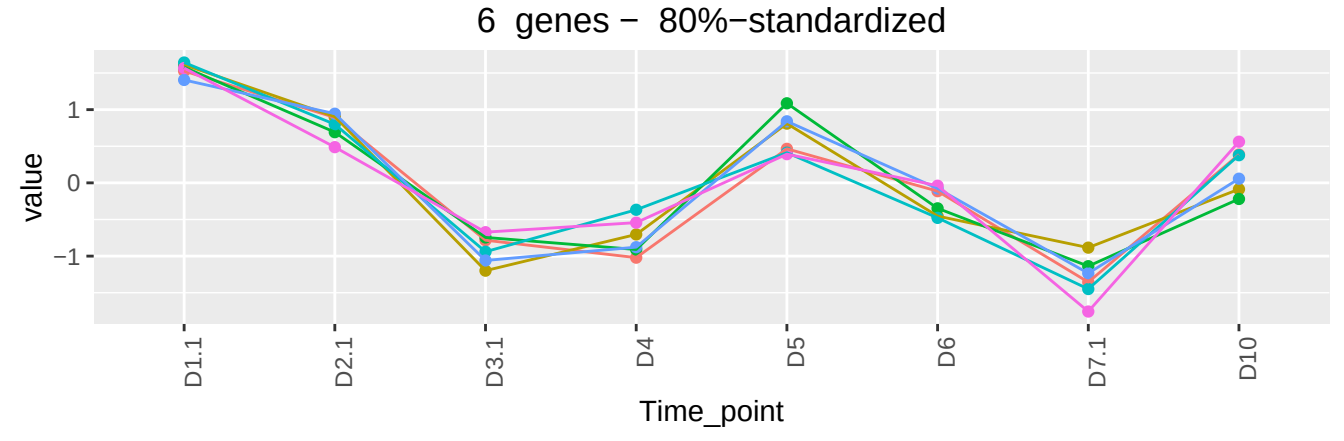
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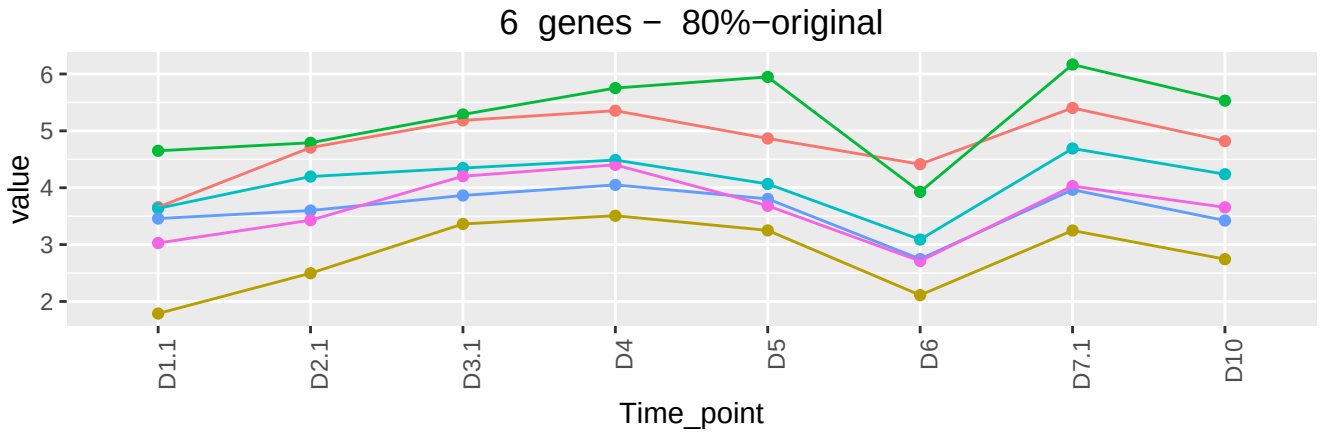
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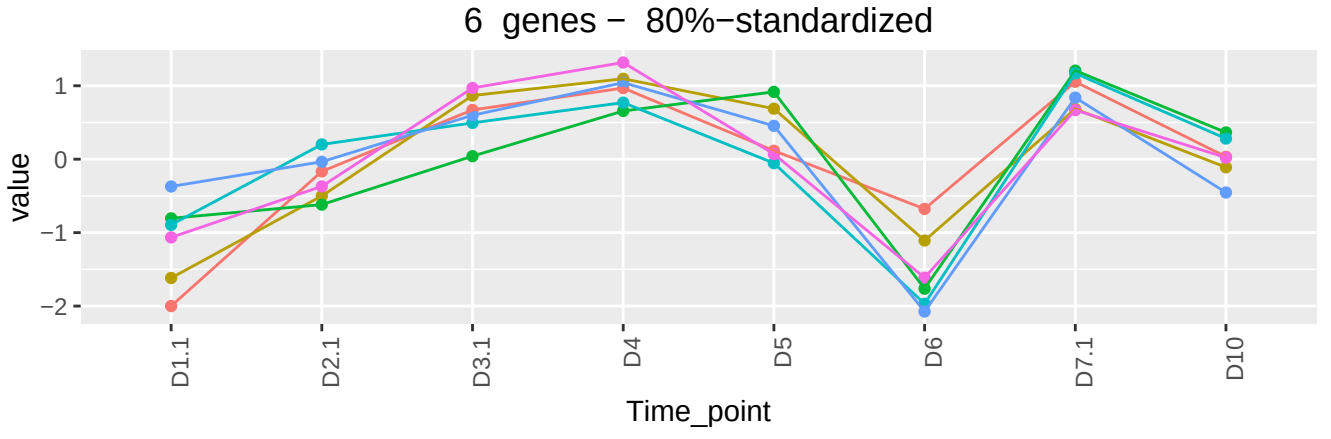
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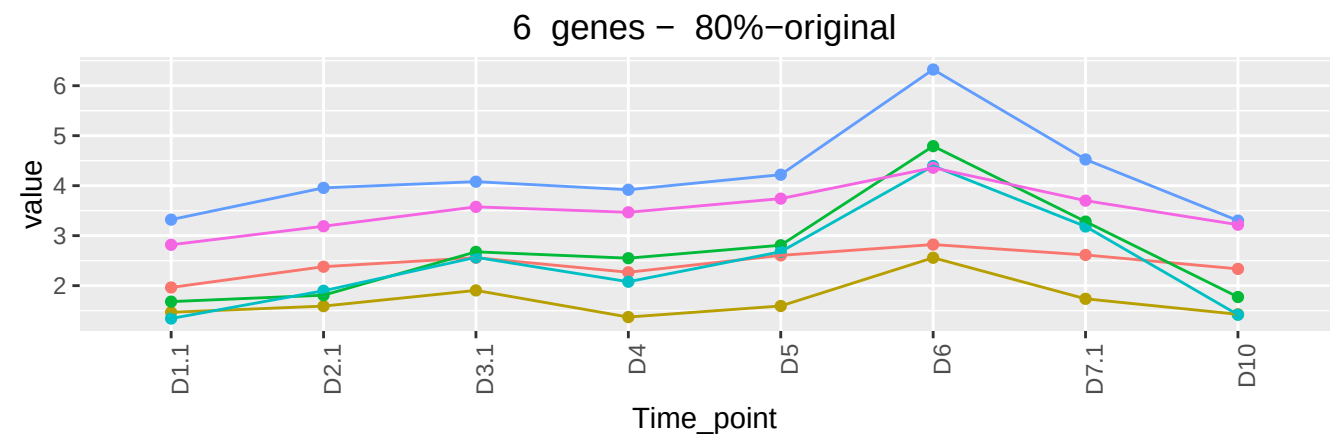
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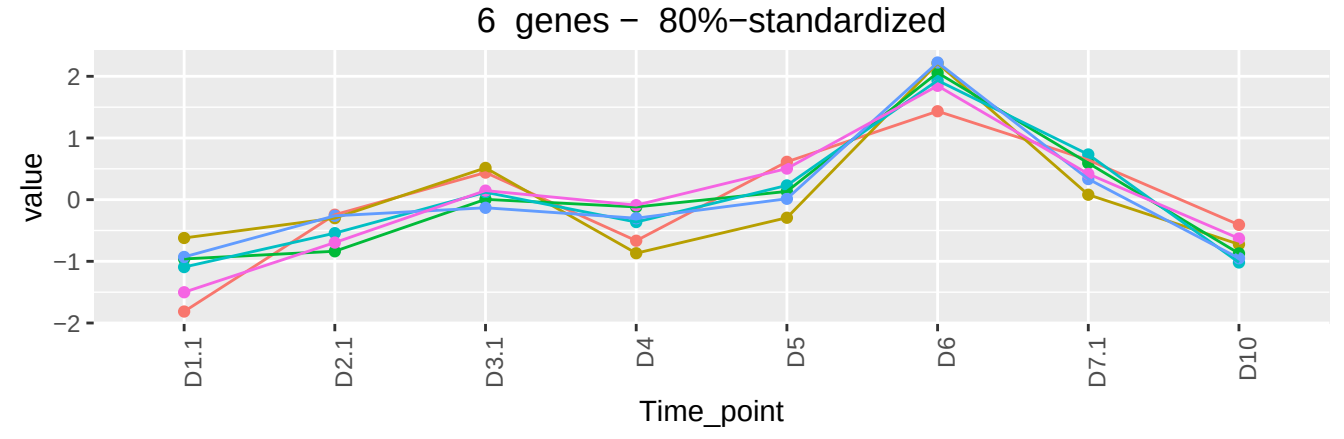
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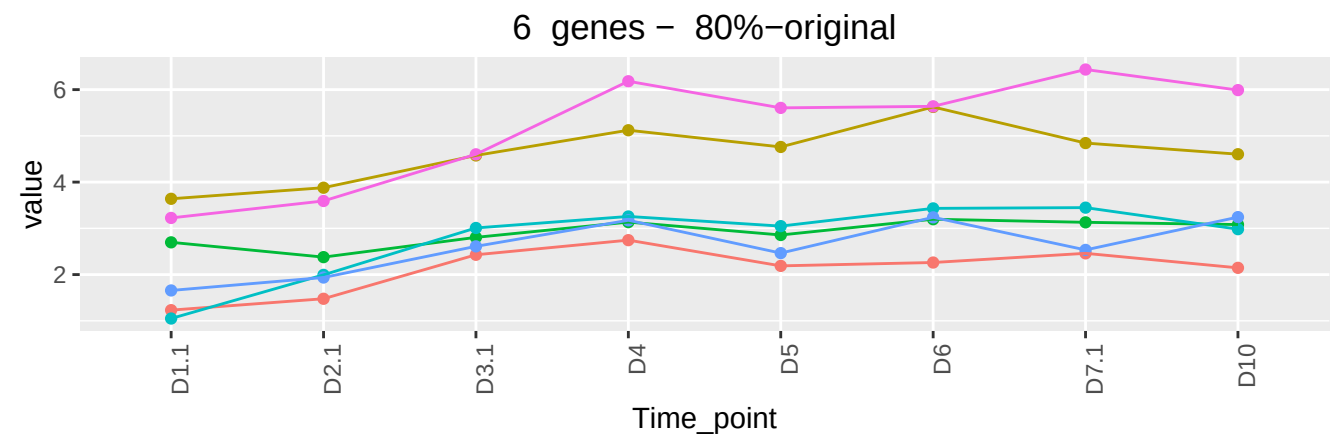
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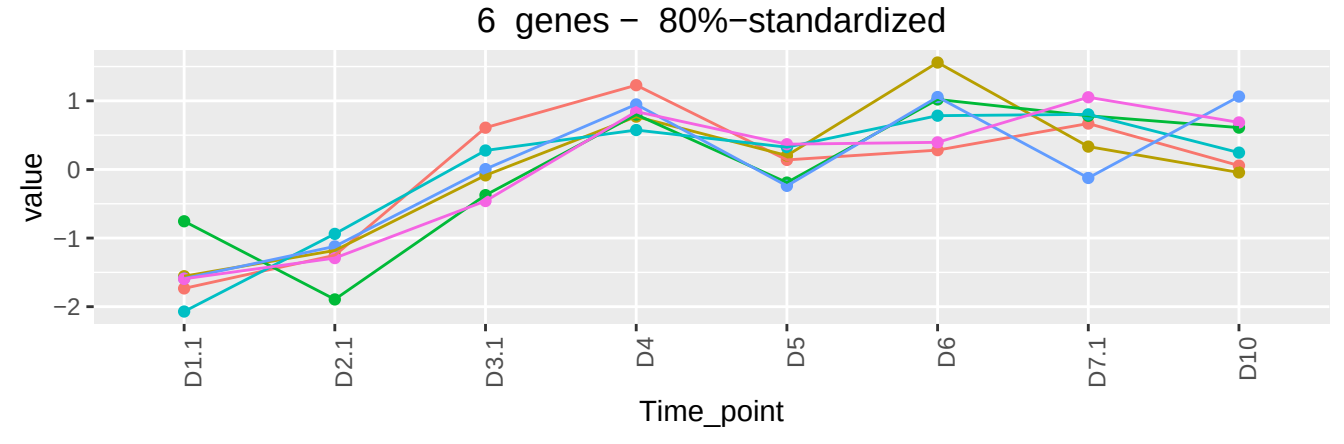
Time_point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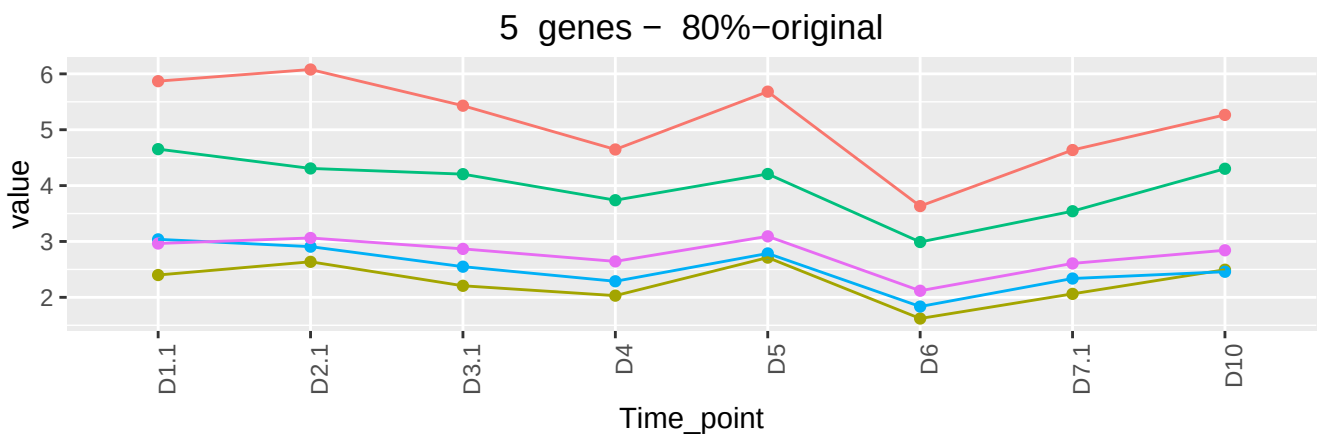
Time_point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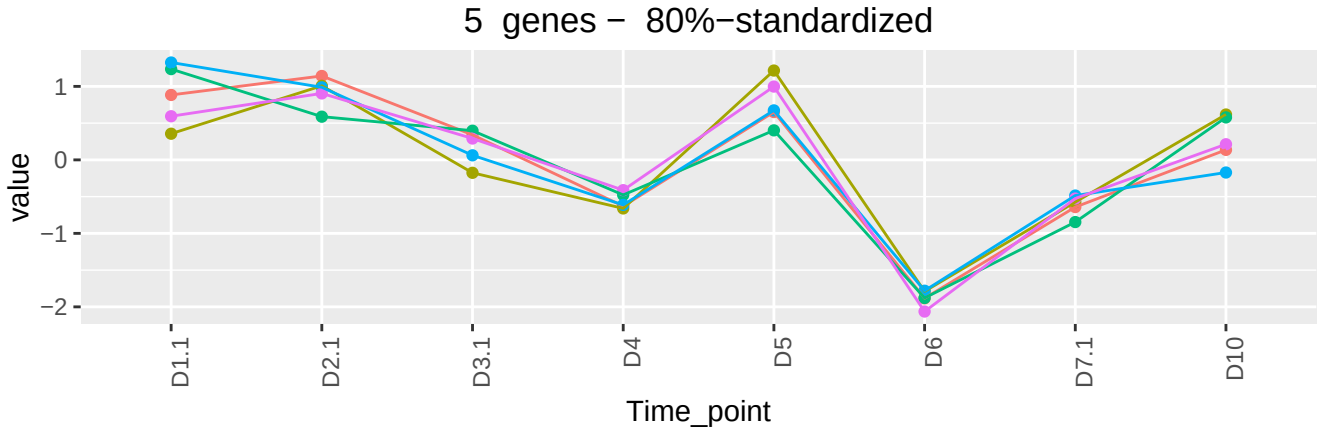
Time_point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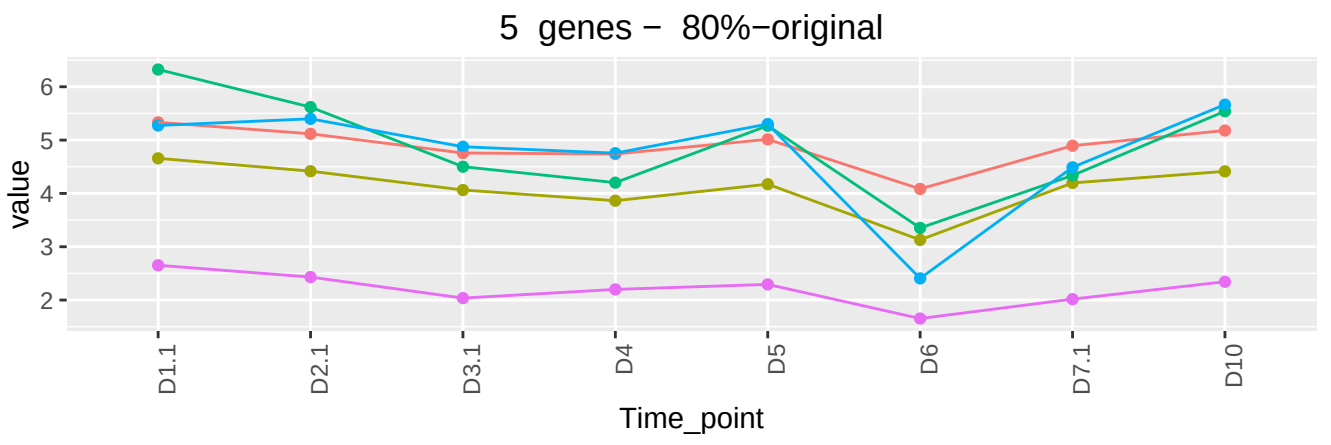
Time_point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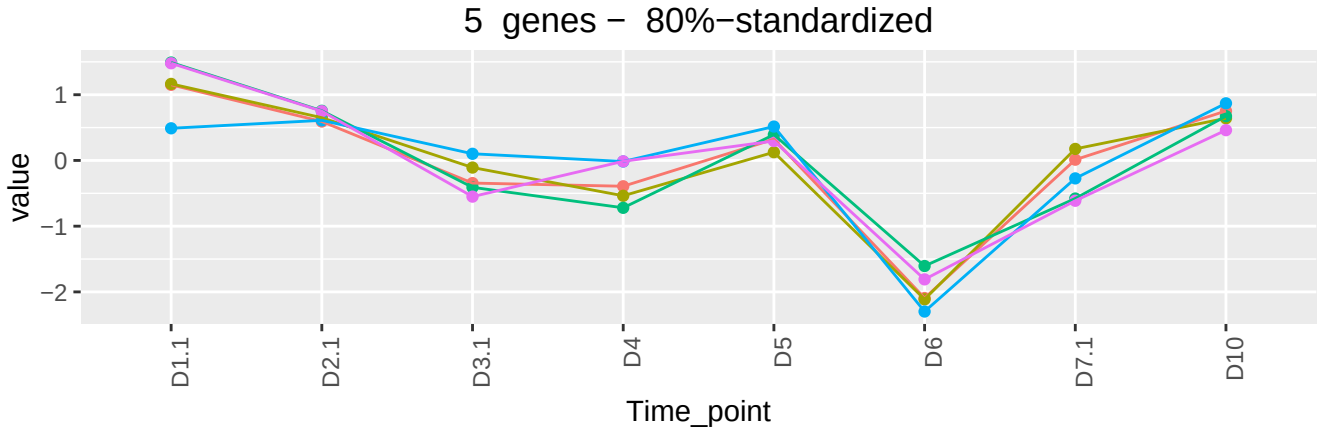
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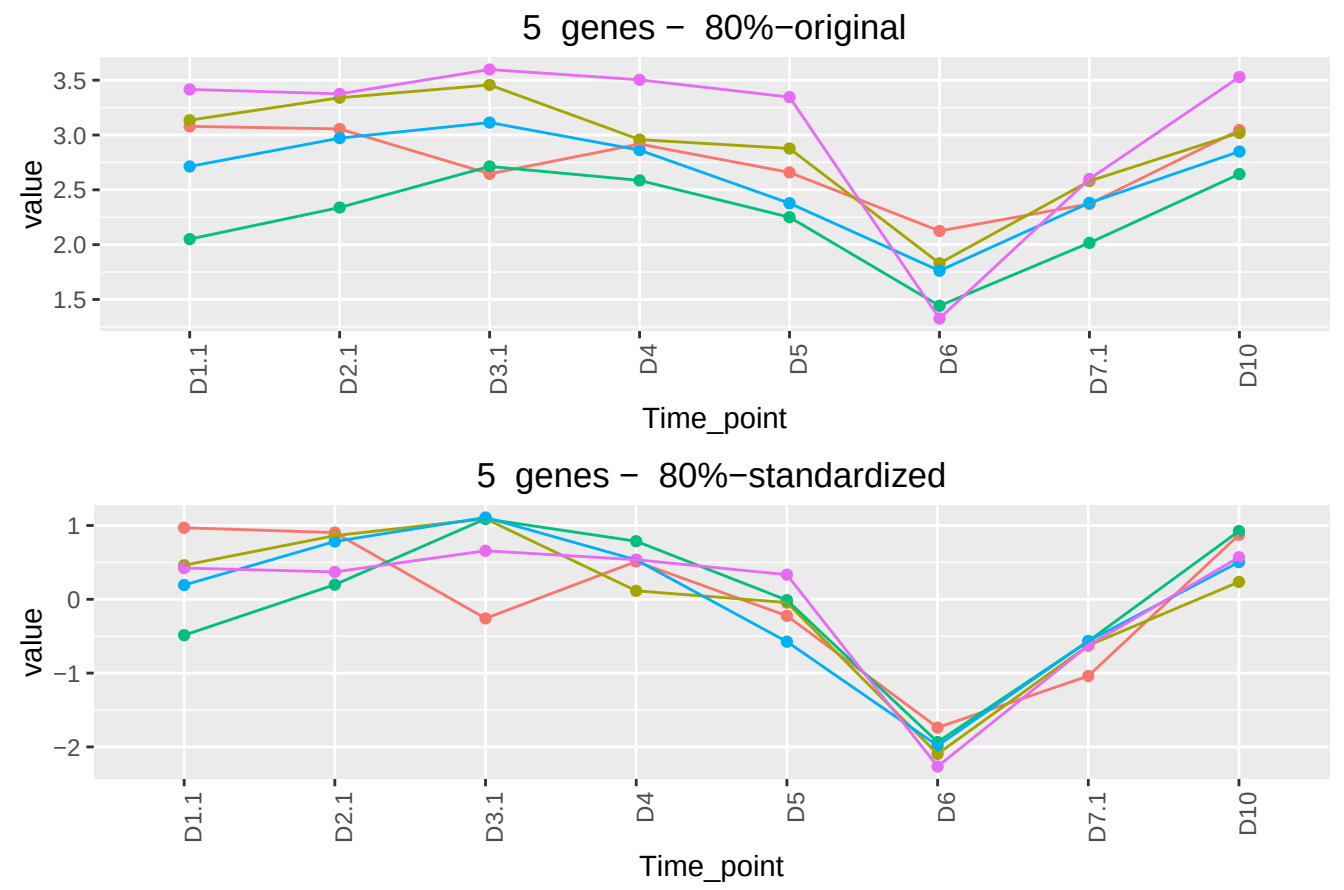
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D5
D6
D7.1
D10
Time_point
5 genes − 80%−standardized
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−2
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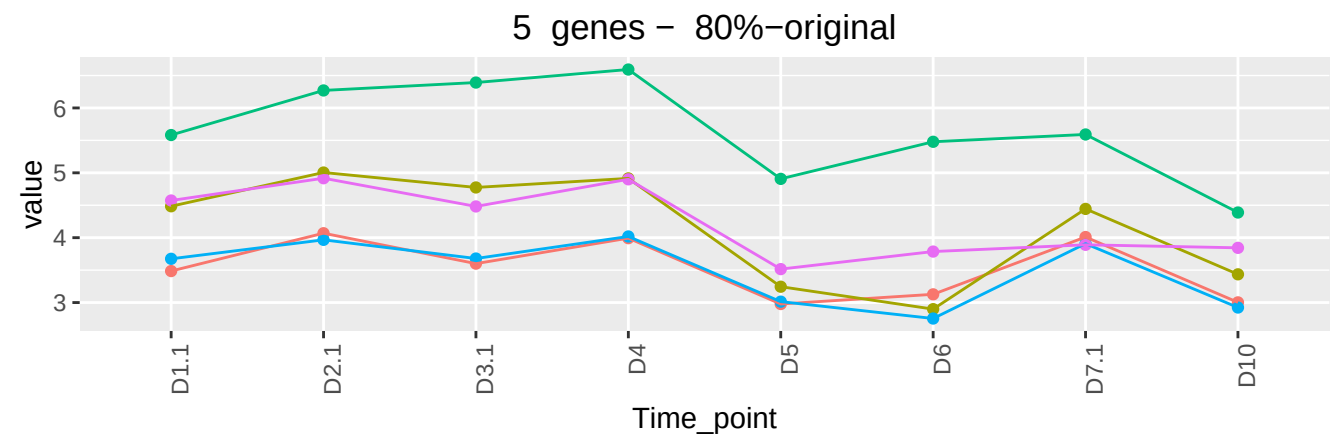
Time_point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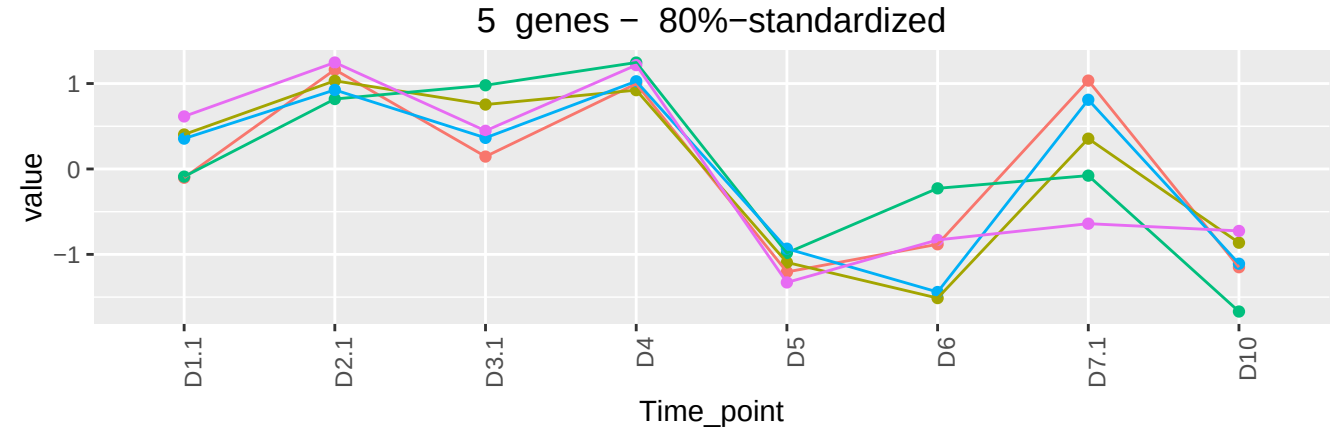
Time_point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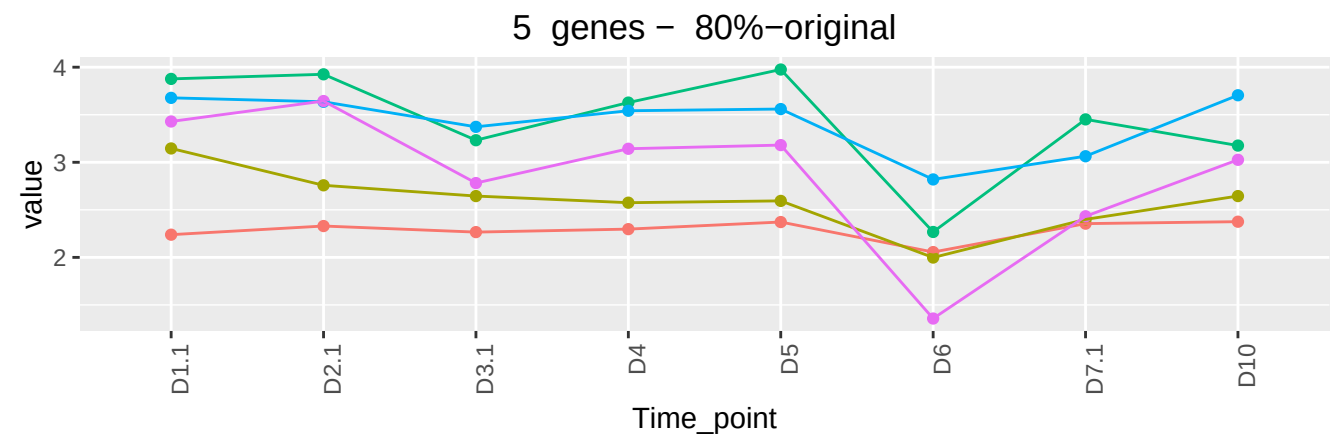
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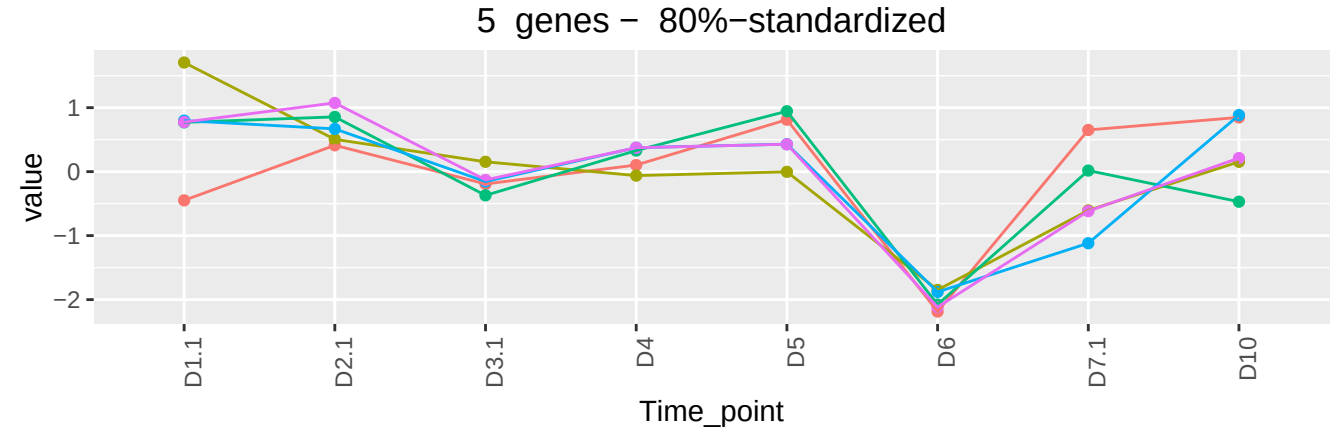
Time_point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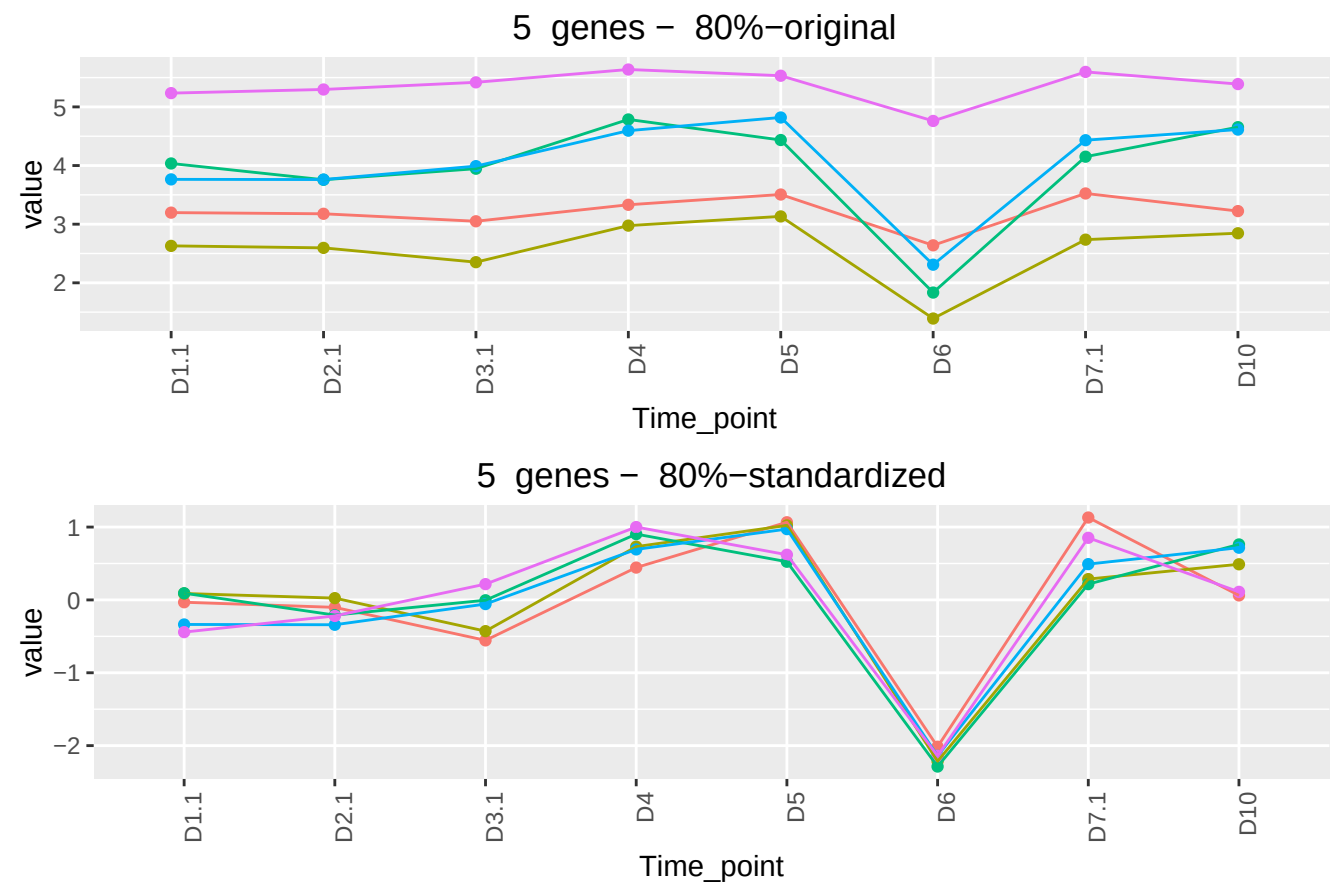
Time_point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Time_point
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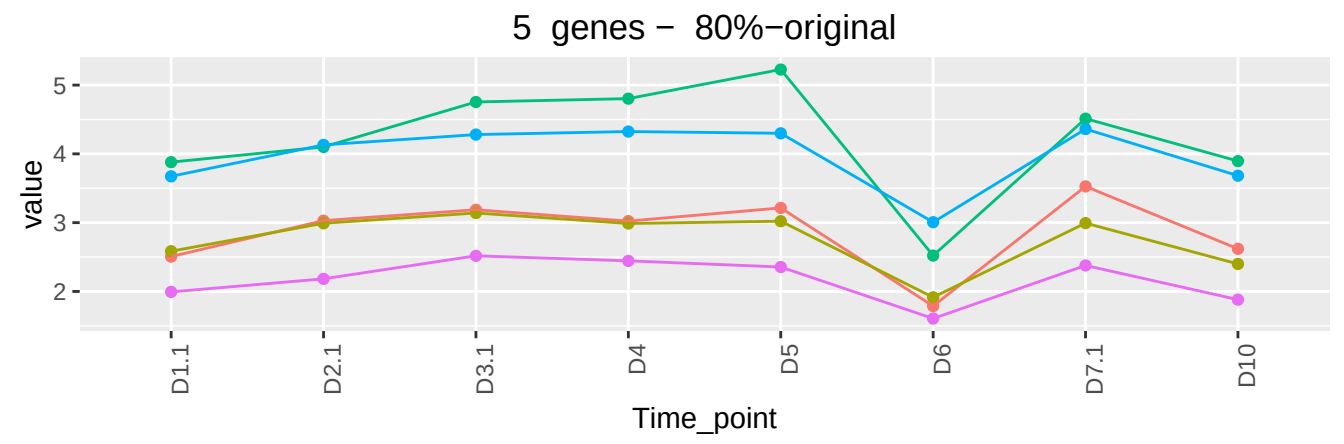
Time_point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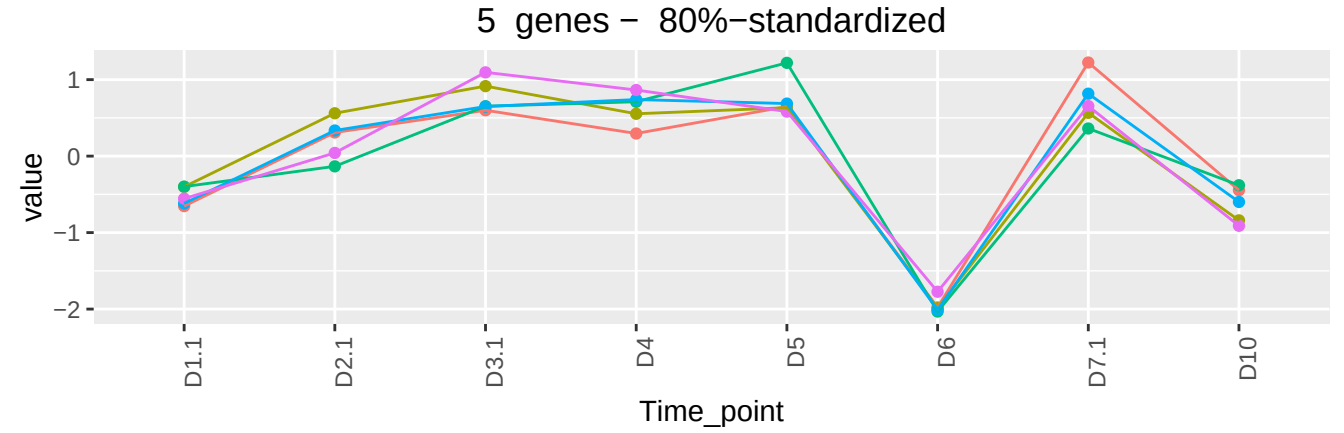
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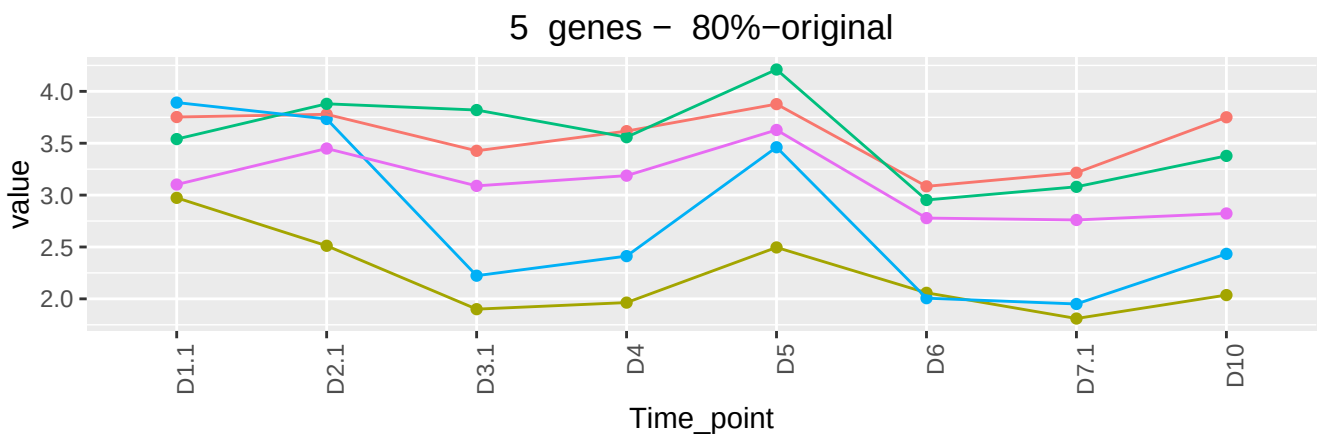
Time_point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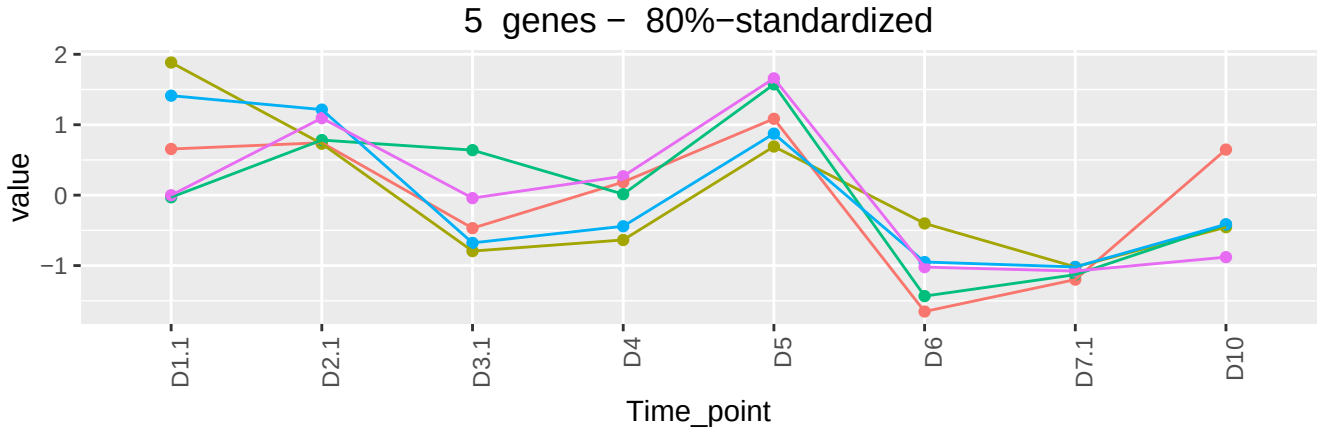
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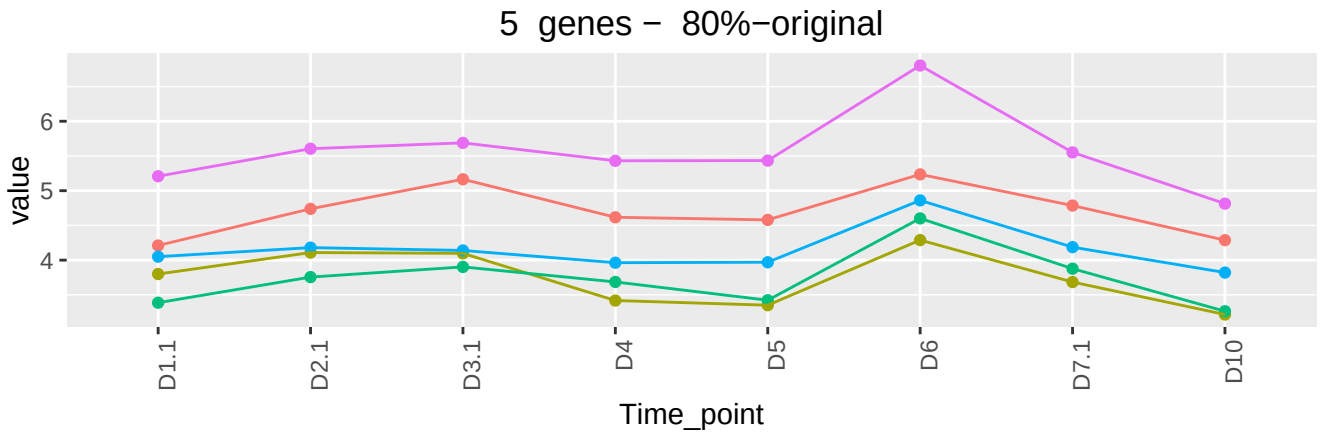
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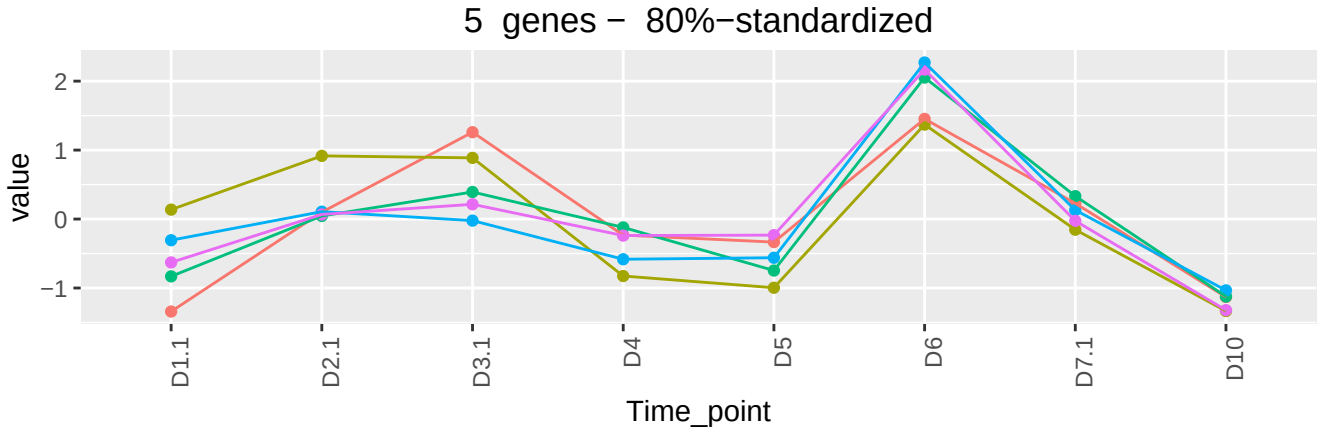
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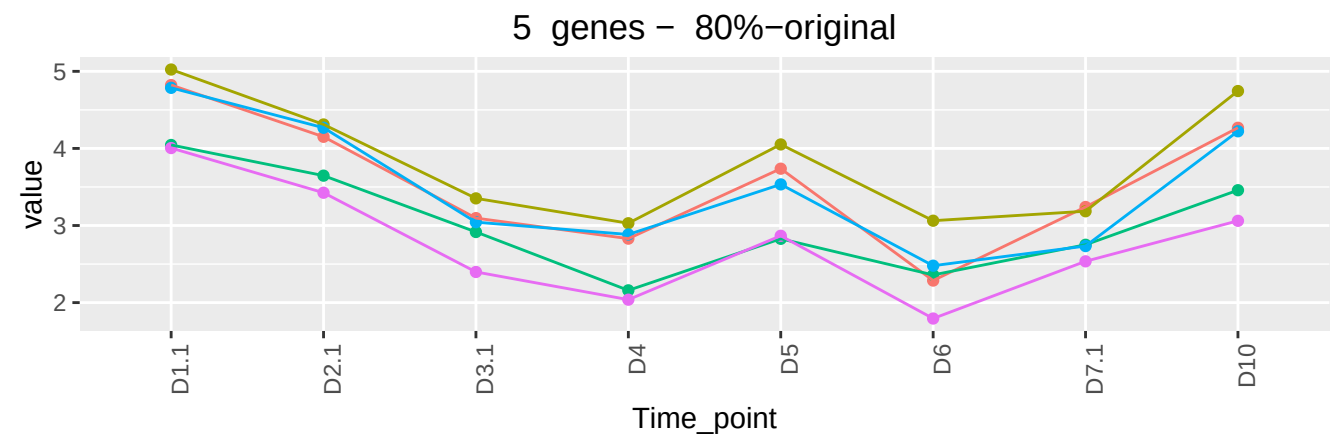
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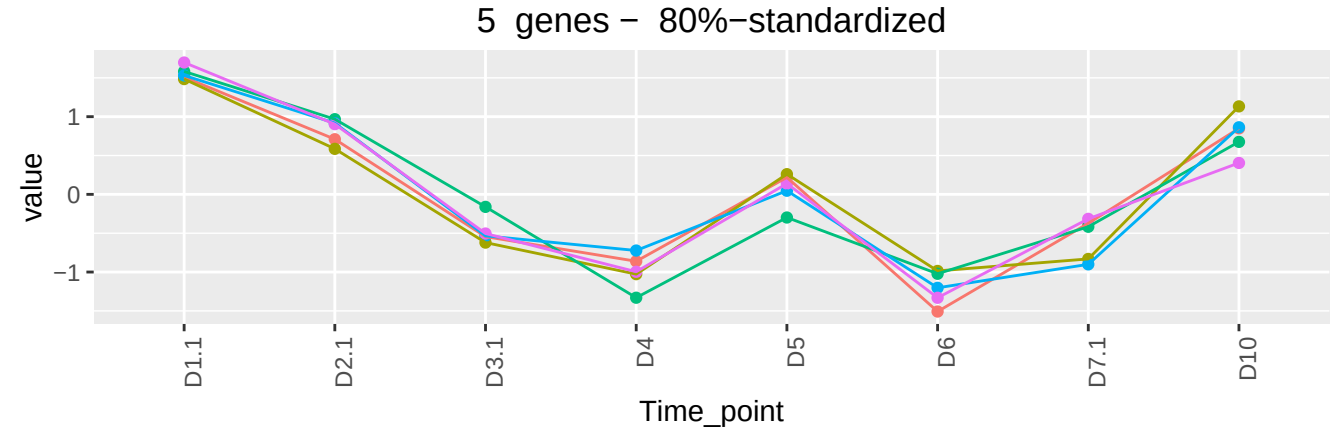
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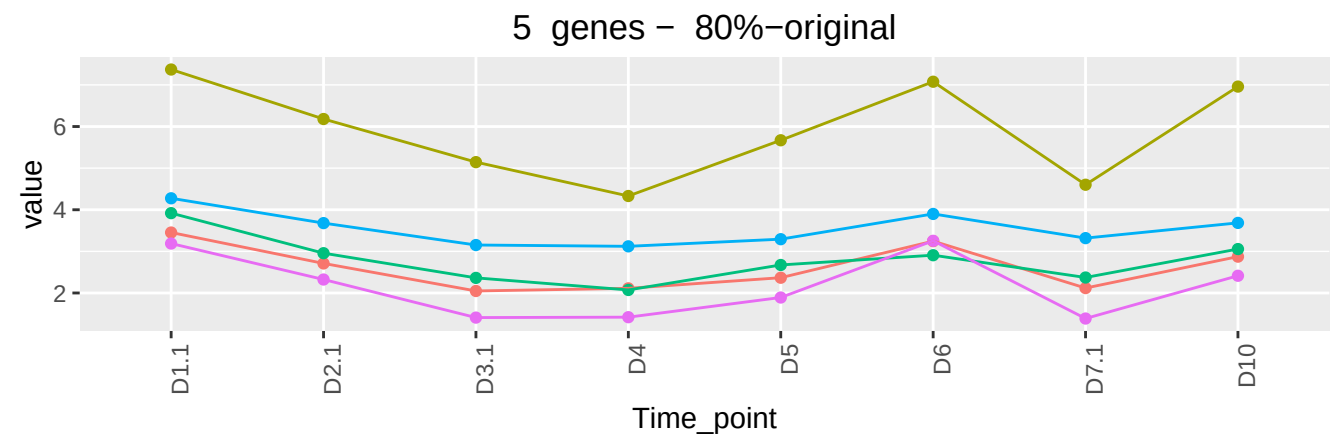
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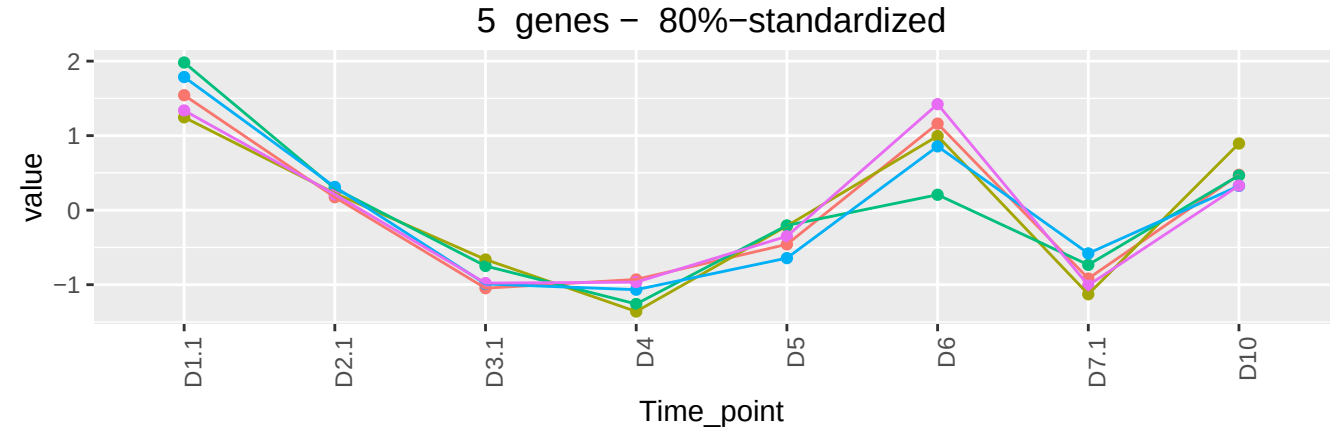
Time_point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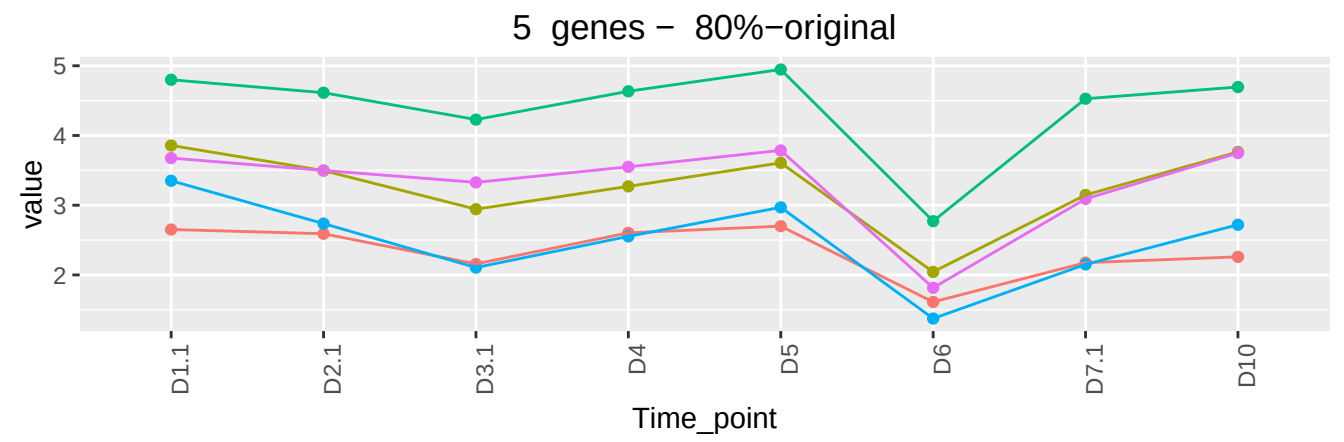
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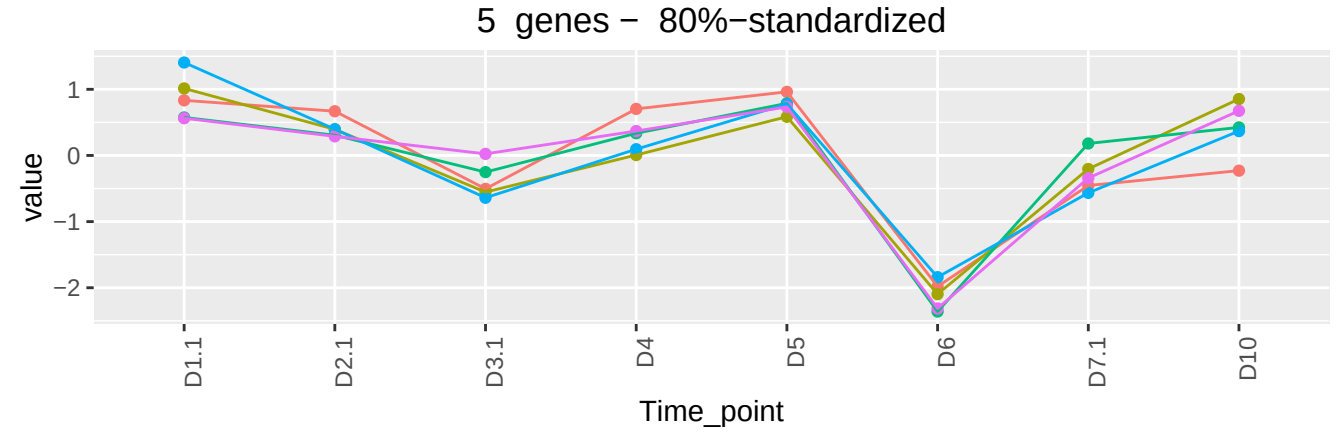
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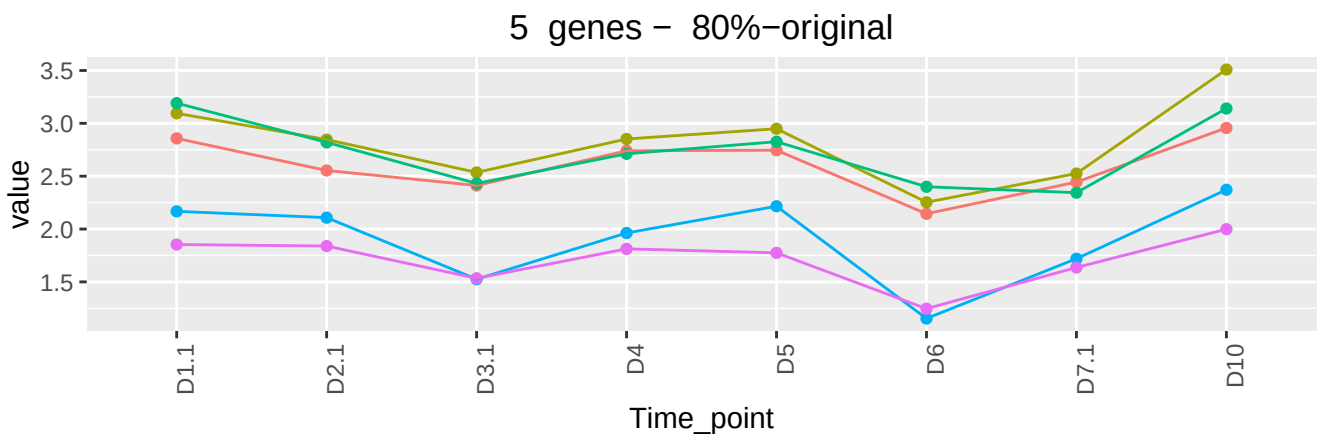
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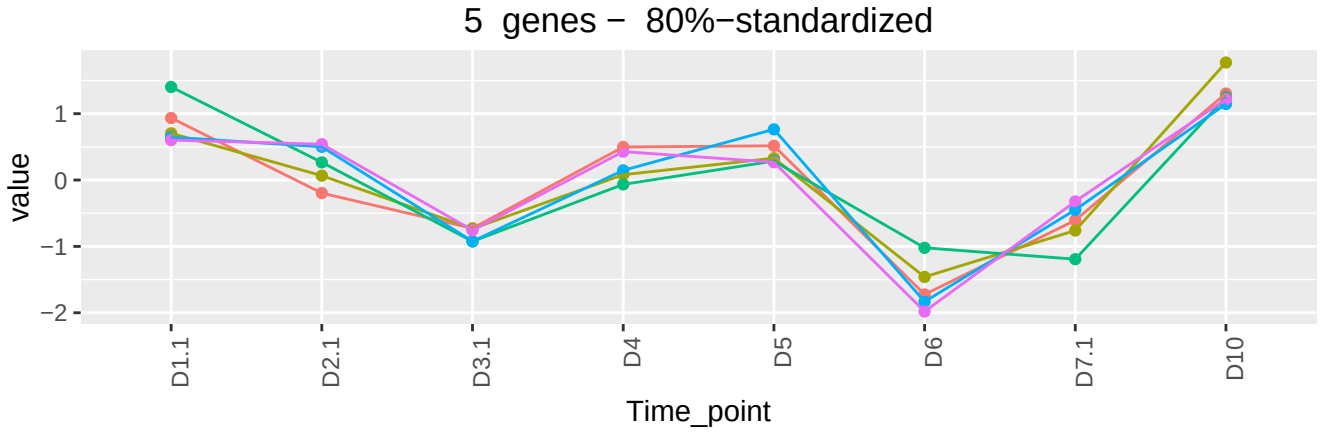
Time_point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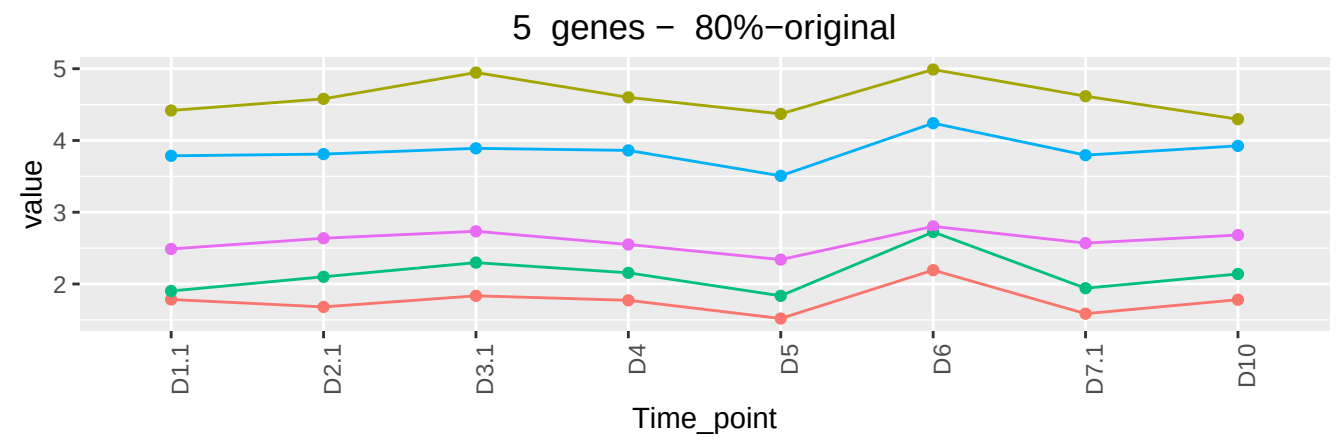
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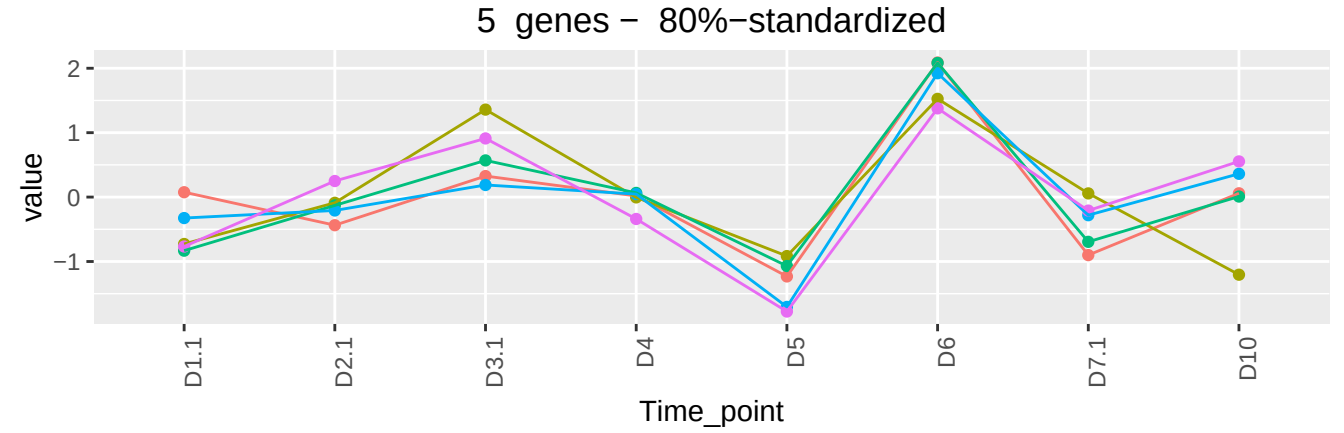
Time_point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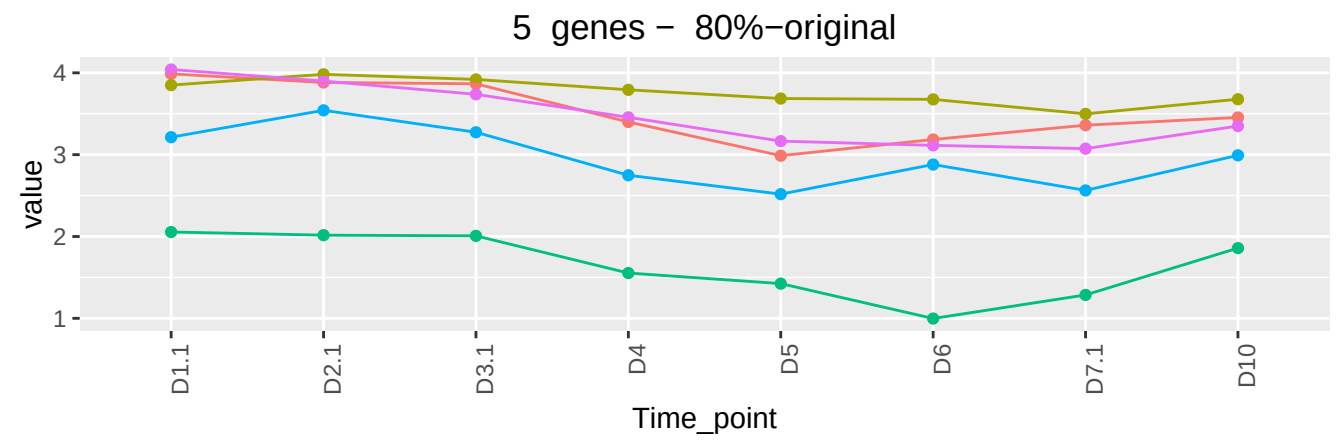
Time_point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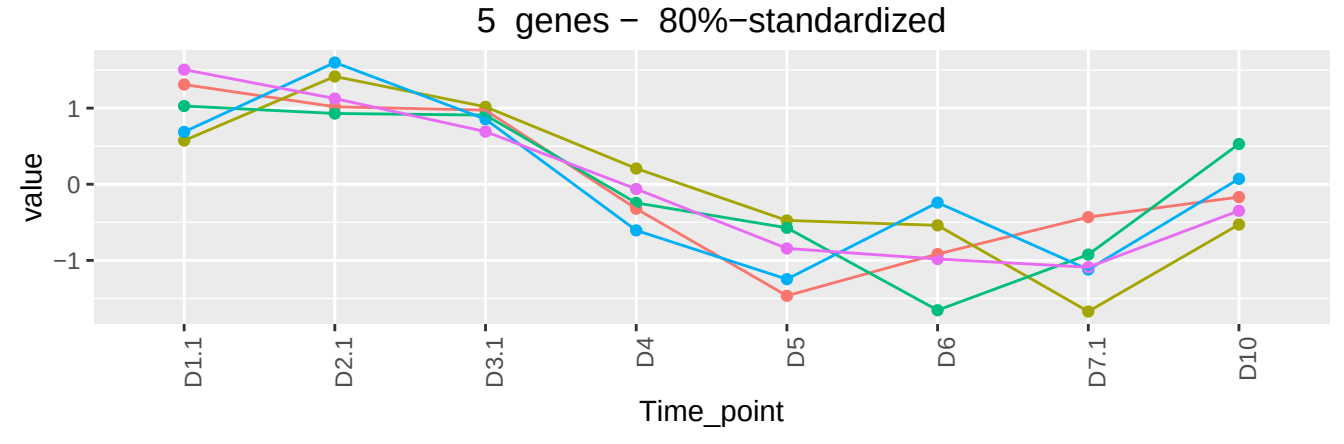
Time_point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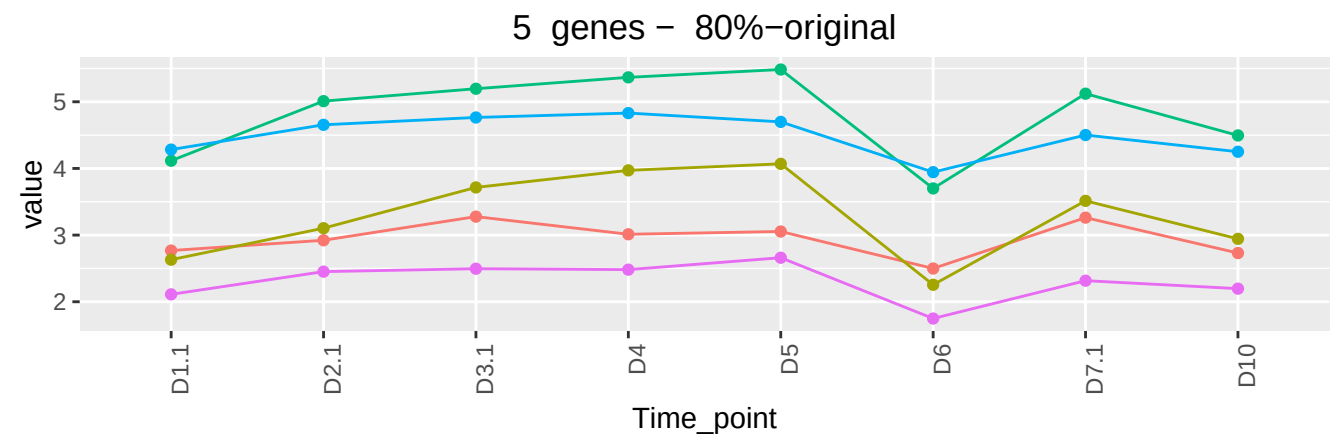
Time_point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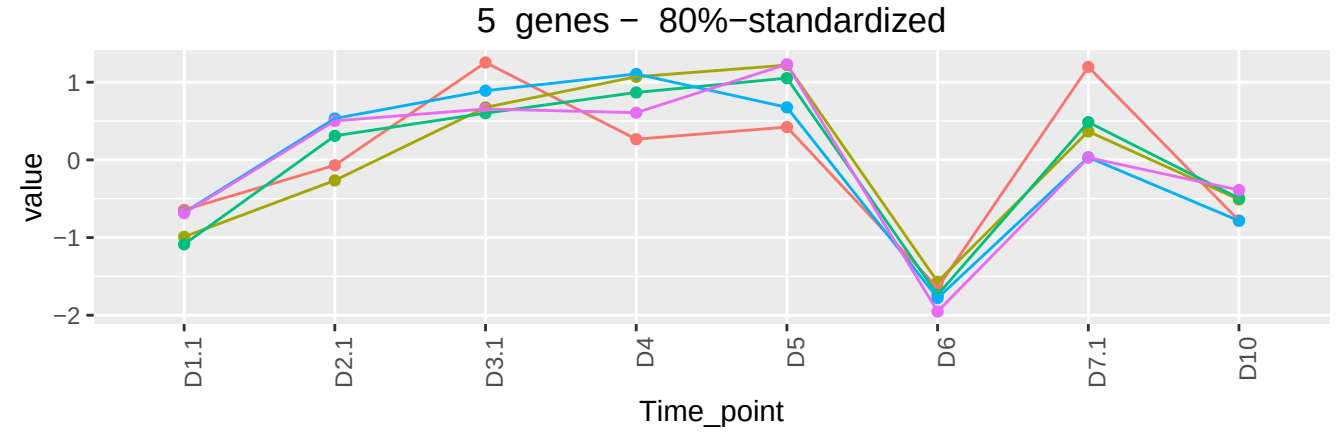
Time_point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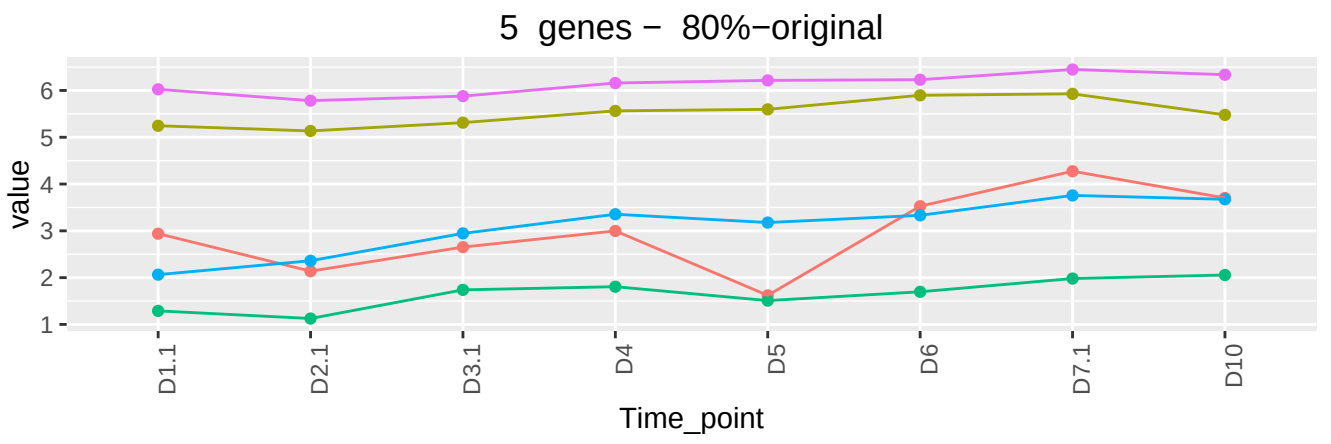
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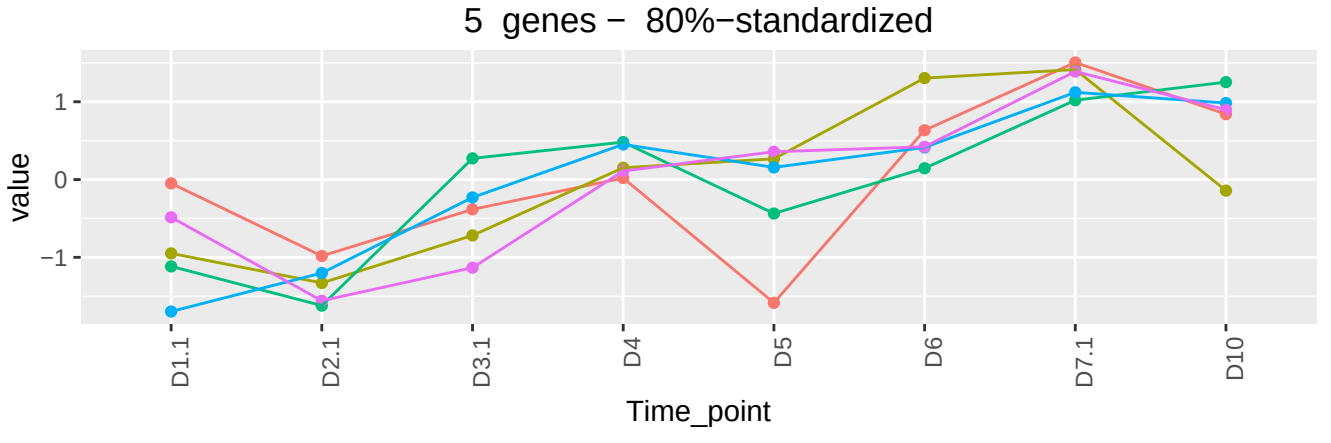
Time_point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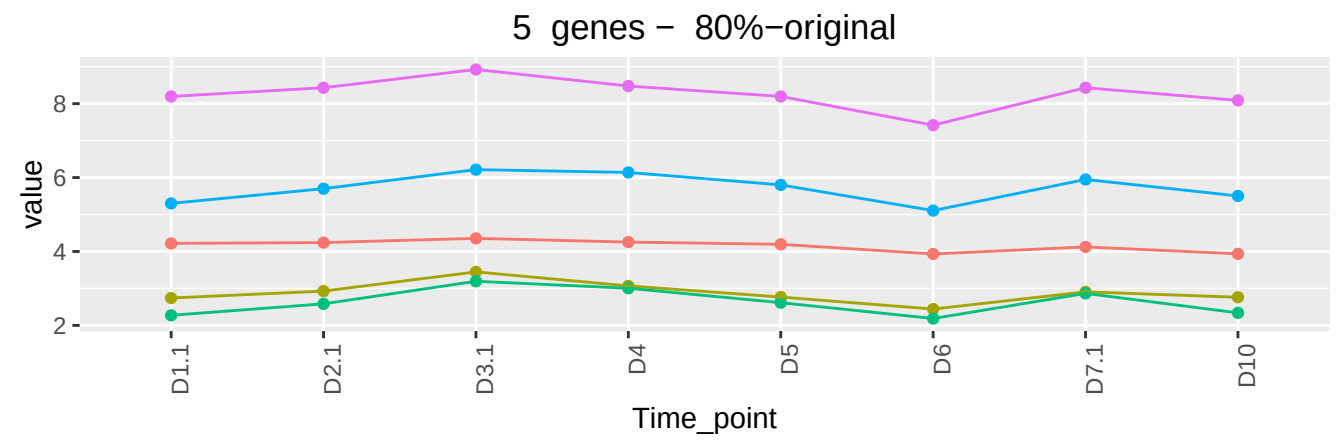
Time_point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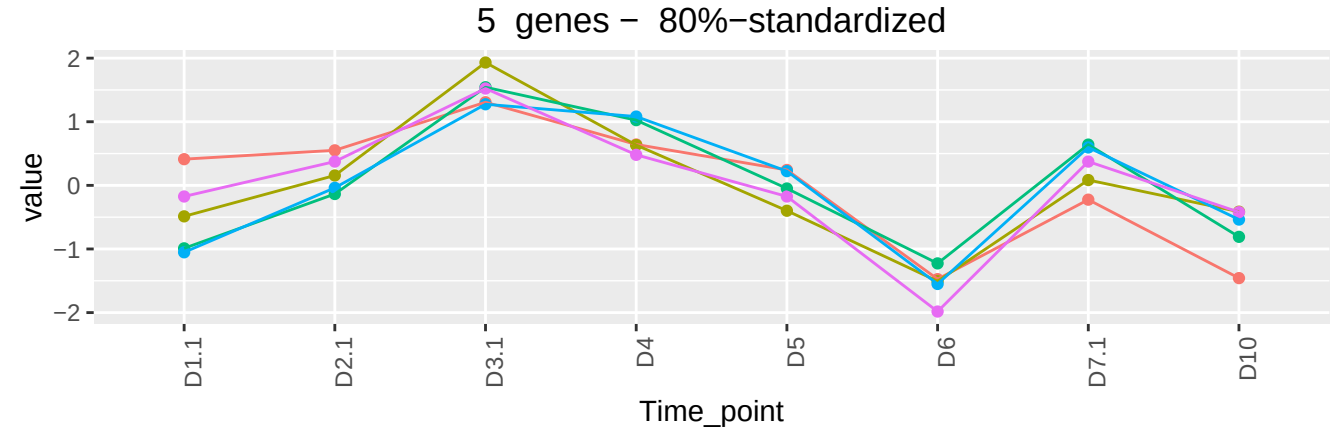
Time_point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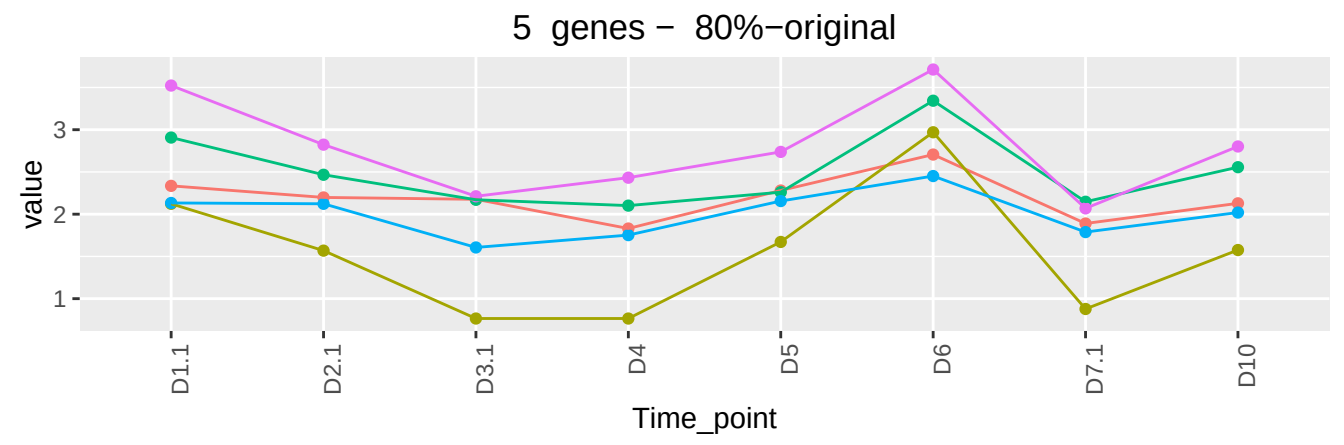
Time_point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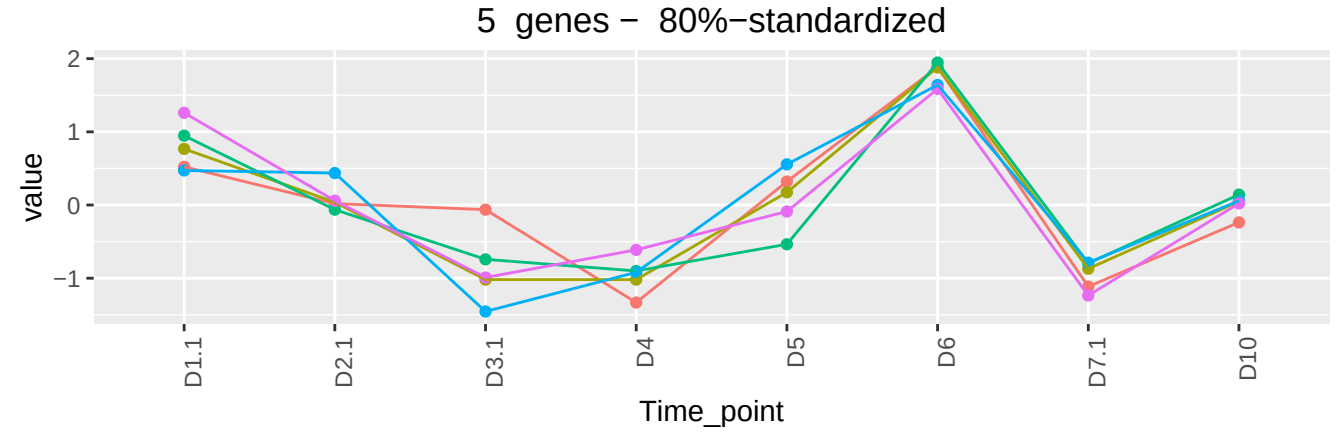
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D7.1
D10
Time_point