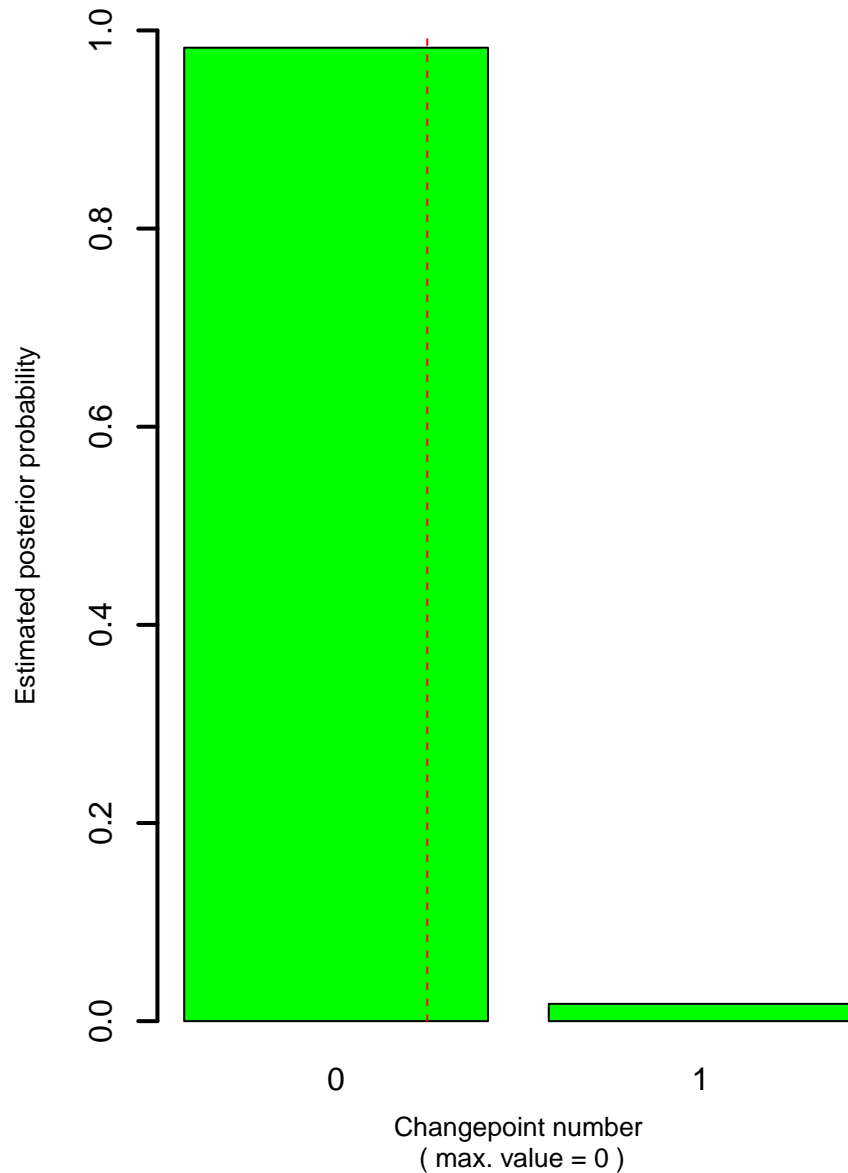


The graph displays a periodic signal over 30 time points. The signal is at a high level (approx. 2.0) for the first 6 time points, then drops to a low level (approx. 1.0) for the next 3 time points. This pattern repeats three times, with the signal returning to the high level for 6 time points and dropping to the low level for 3 time points in each cycle. The x-axis is labeled 'Time point' and ranges from 0 to 30. The y-axis ranges from 1.0 to 2.0.

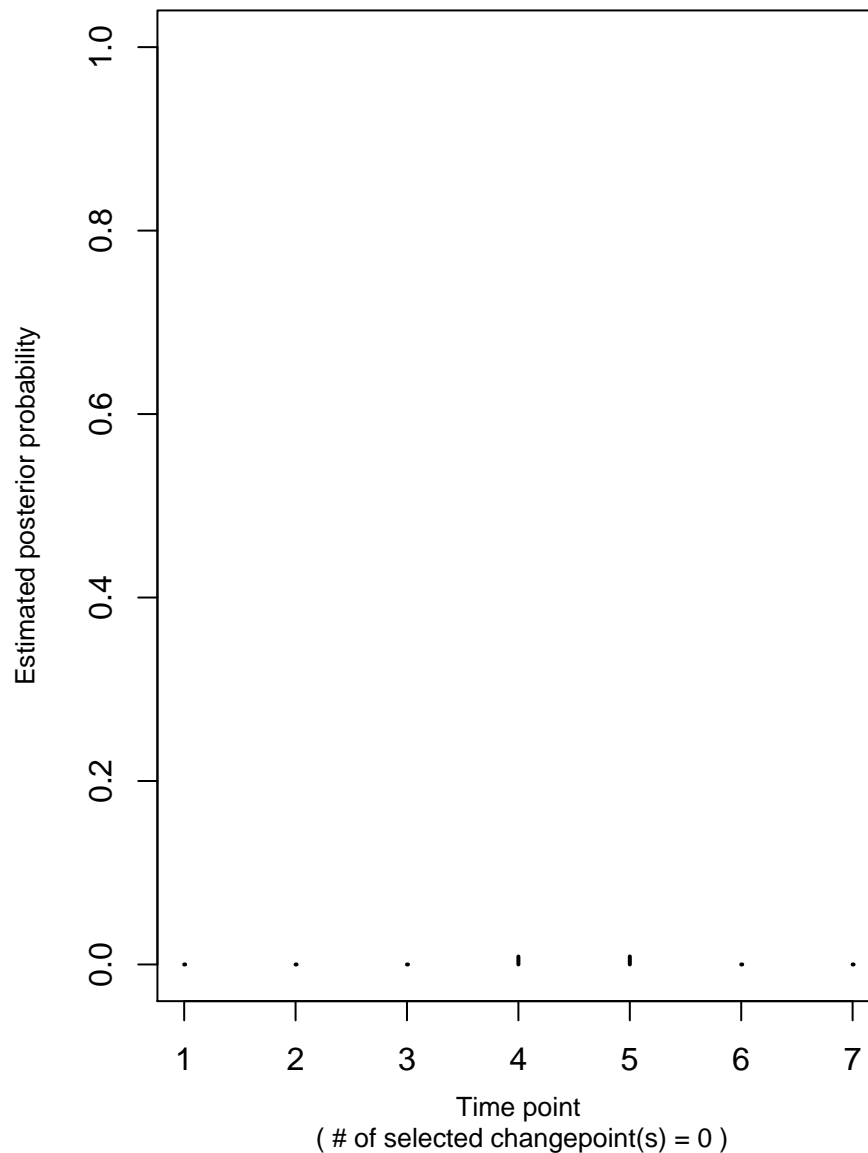
Time point	Signal Level
1	2.0
2	2.0
3	2.0
4	2.0
5	2.0
6	2.0
7	1.0
8	1.0
9	1.0
10	2.0
11	2.0
12	2.0
13	2.0
14	2.0
15	1.0
16	1.0
17	1.0
18	2.0
19	2.0
20	2.0
21	1.0
22	1.0
23	1.0
24	2.0
25	2.0
26	2.0
27	2.0
28	1.0
29	1.0
30	1.0

The graph displays two oscillating expression profiles over 30 time points. The y-axis, labeled 'Expression value', ranges from 1.0 to 1.8. The x-axis, labeled 'Time point', ranges from 0 to 30. Both genes show a periodic pattern, alternating between a minimum value of 1.0 and a maximum value of 1.8. The two profiles are out of phase, with one gene reaching its peak when the other is at its minimum.

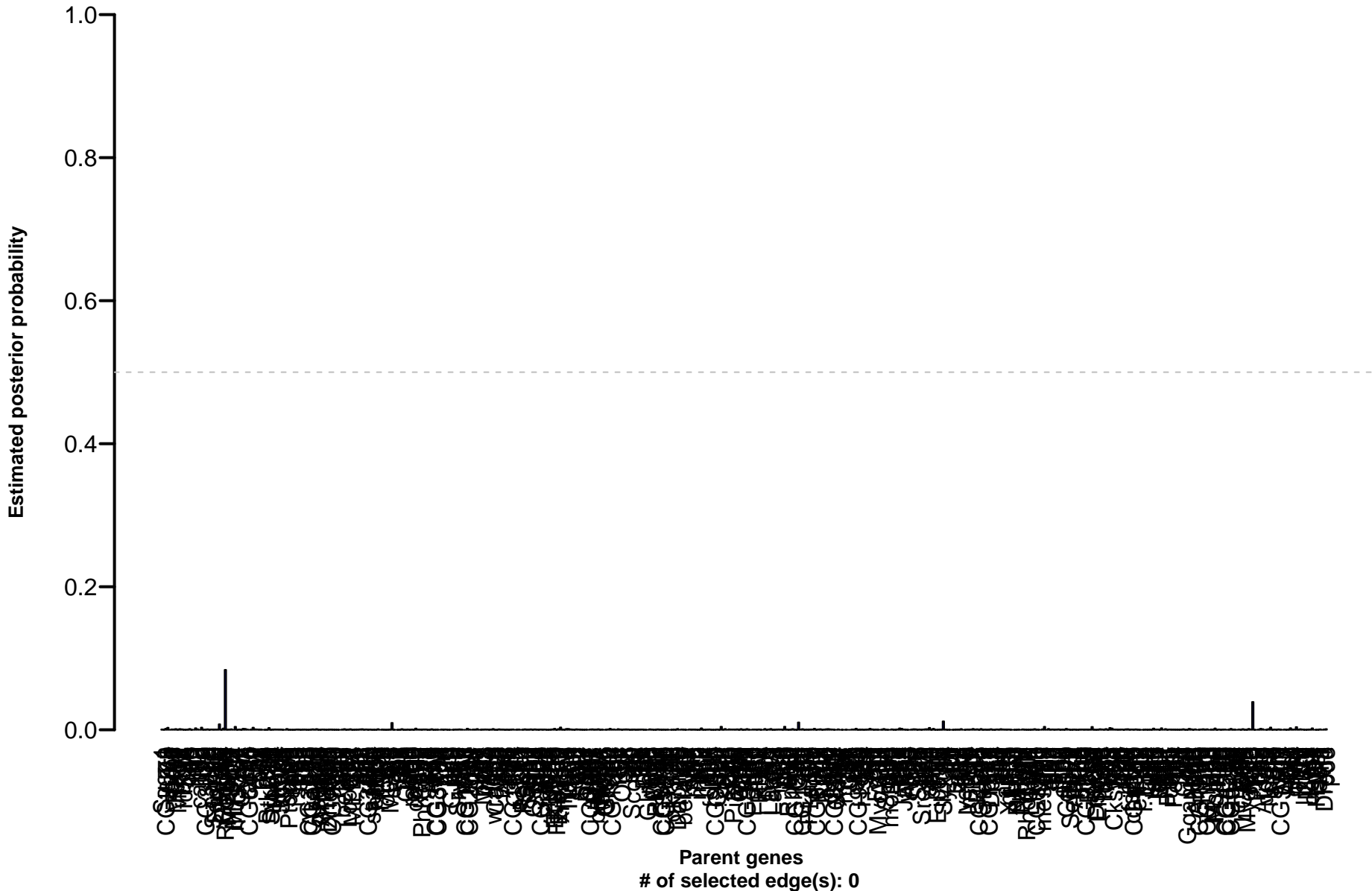
Number of changepoint
– target gene: Apc2 –



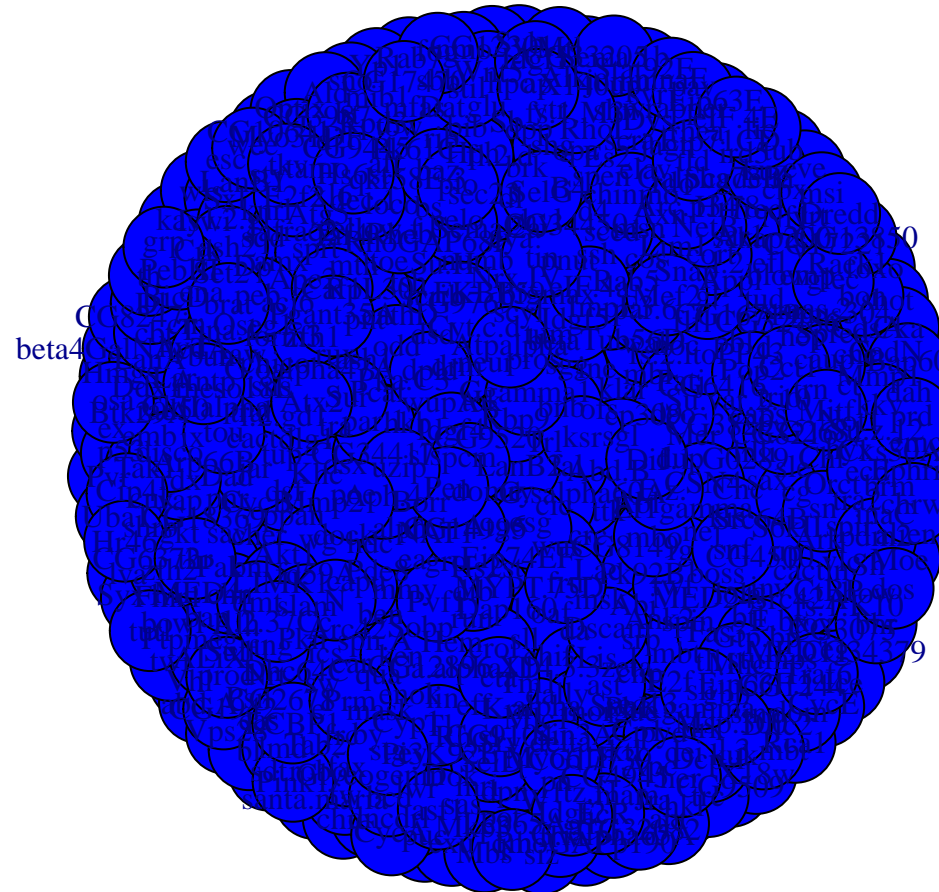
Changepoint position
– target gene: Apc2 –



Regulatory model for target gene: Apc2
Temporal segment # 1 : [2 , 6]



Sub-network # 1
(time point 2 to 6)



— Positive interaction
- - - Negative interaction

ARTIVA summary page
(interactions are arranged in order of decreasing confidence level)

parentGene	targetGene	postProb	CPstart	CPend	interactionSign
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