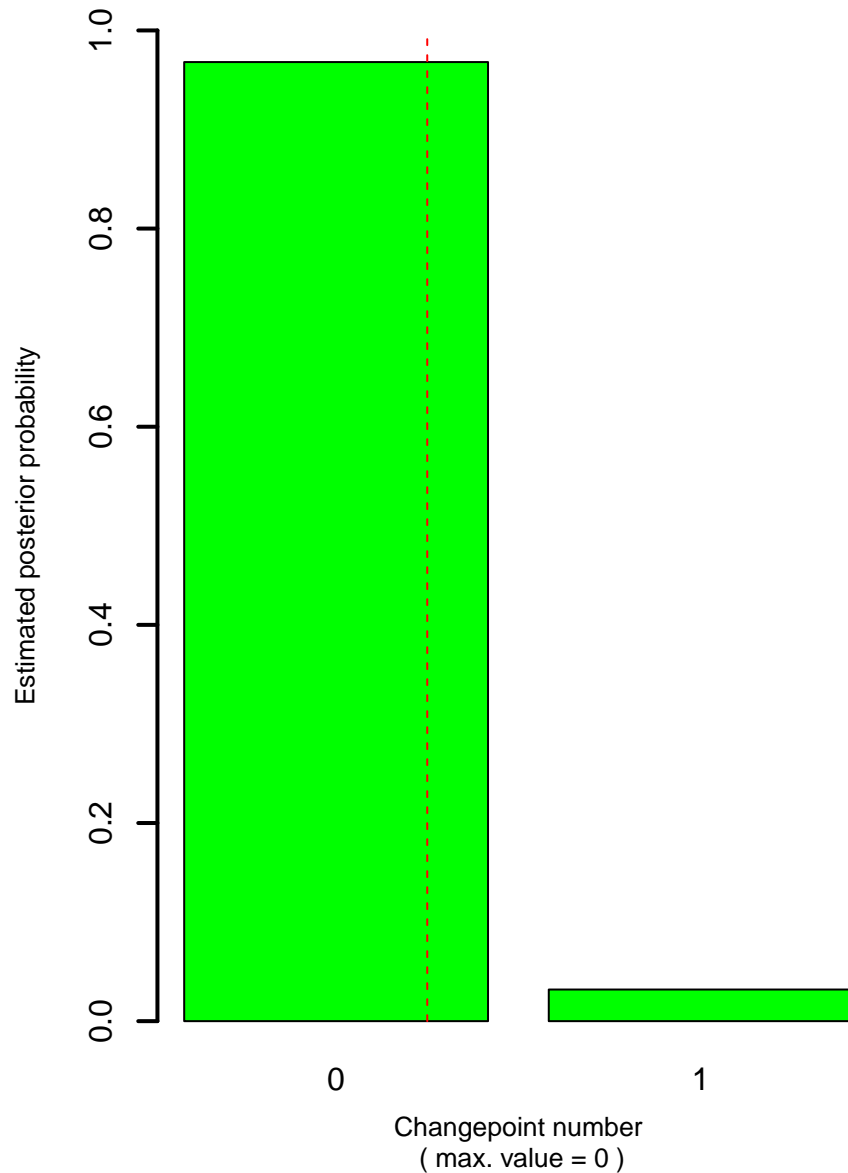


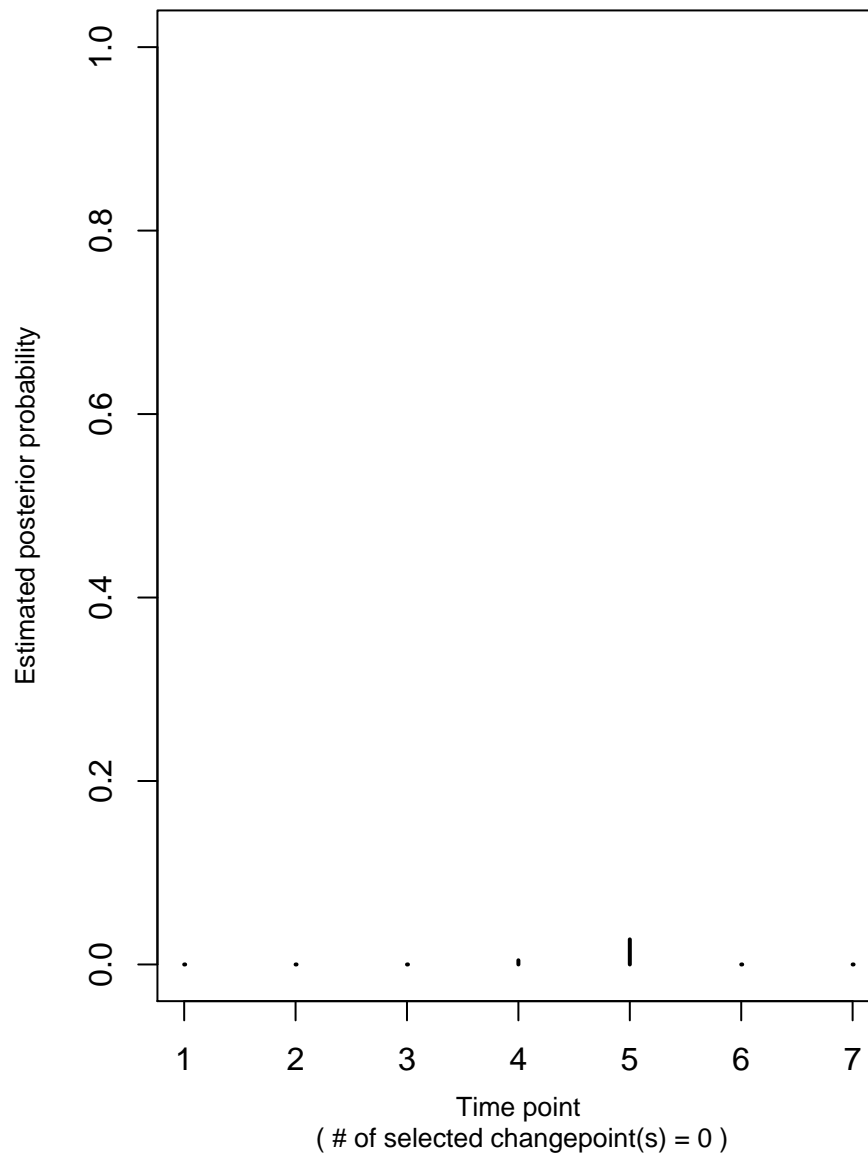
The graph displays a function $f(t)$ over 30 time points. The function is defined as follows:

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

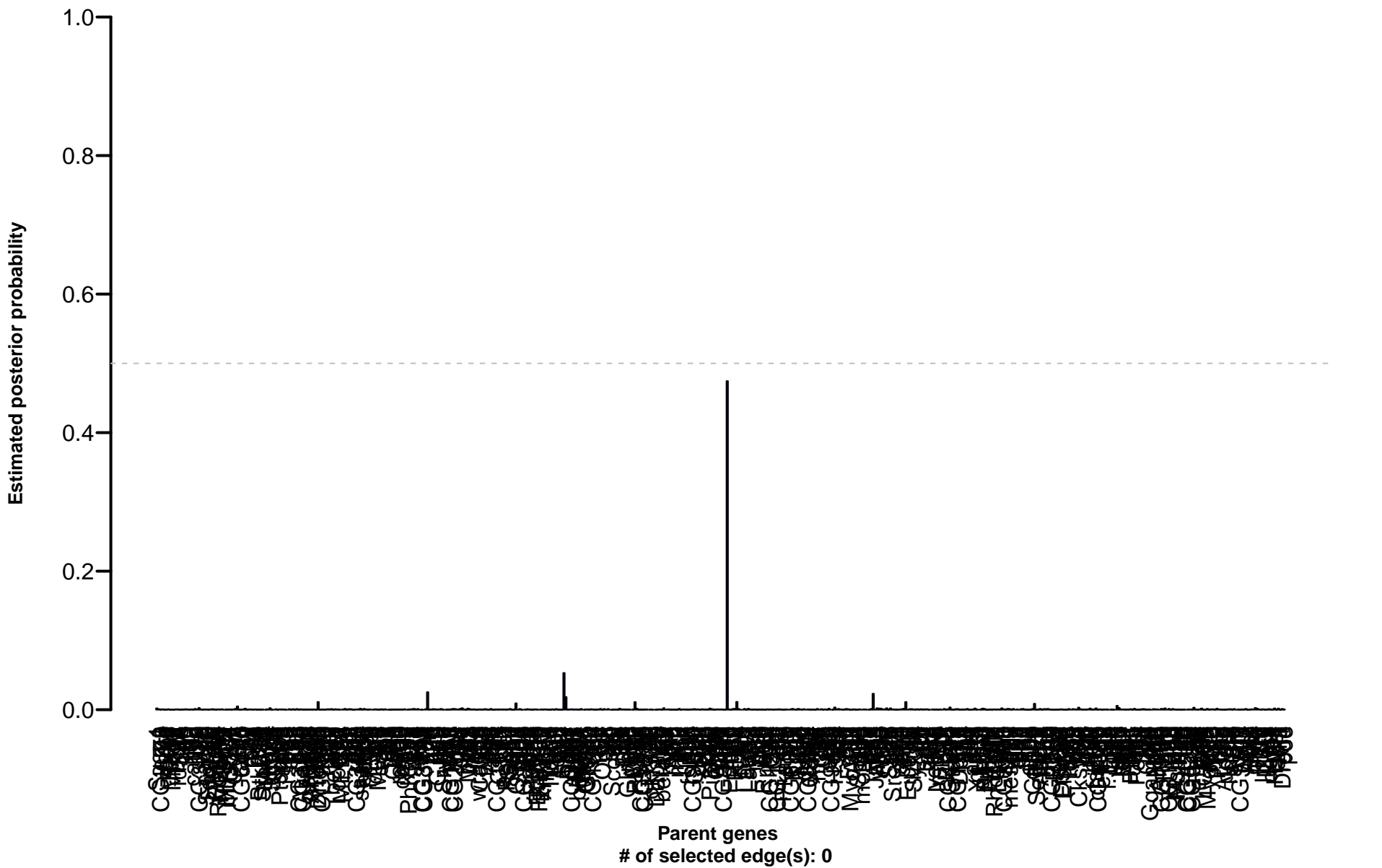
Number of changepoint
– target gene: Patj –



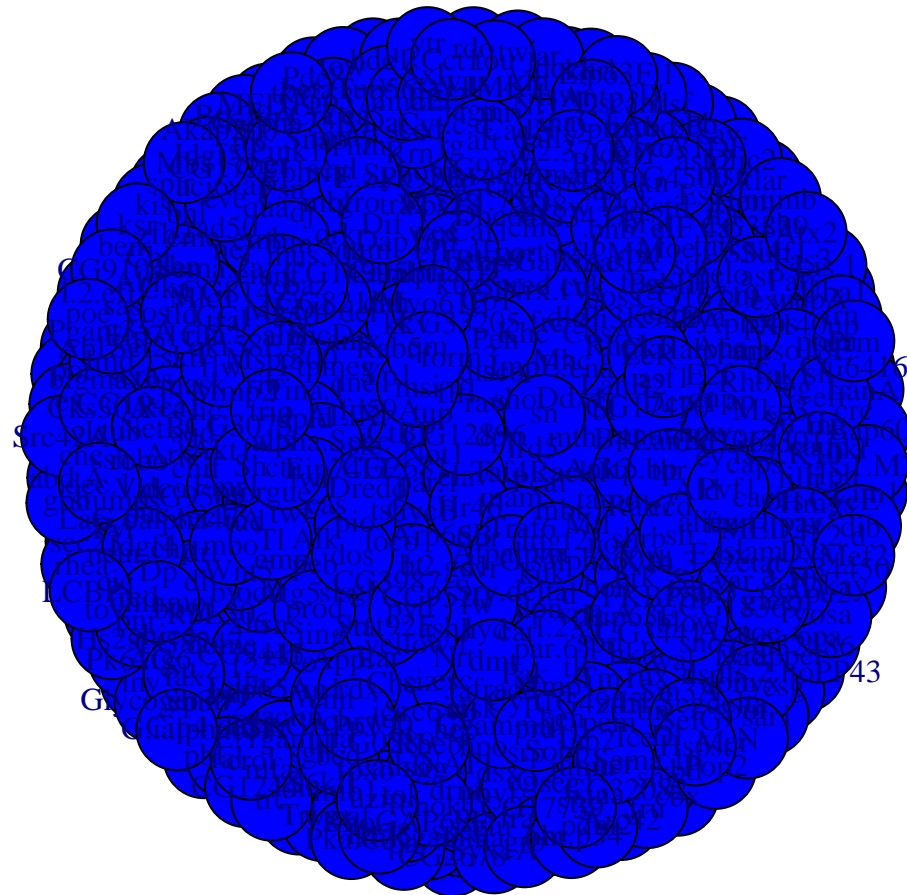
Changepoint position
– target gene: Patj –



Regulatory model for target gene: Patj
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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