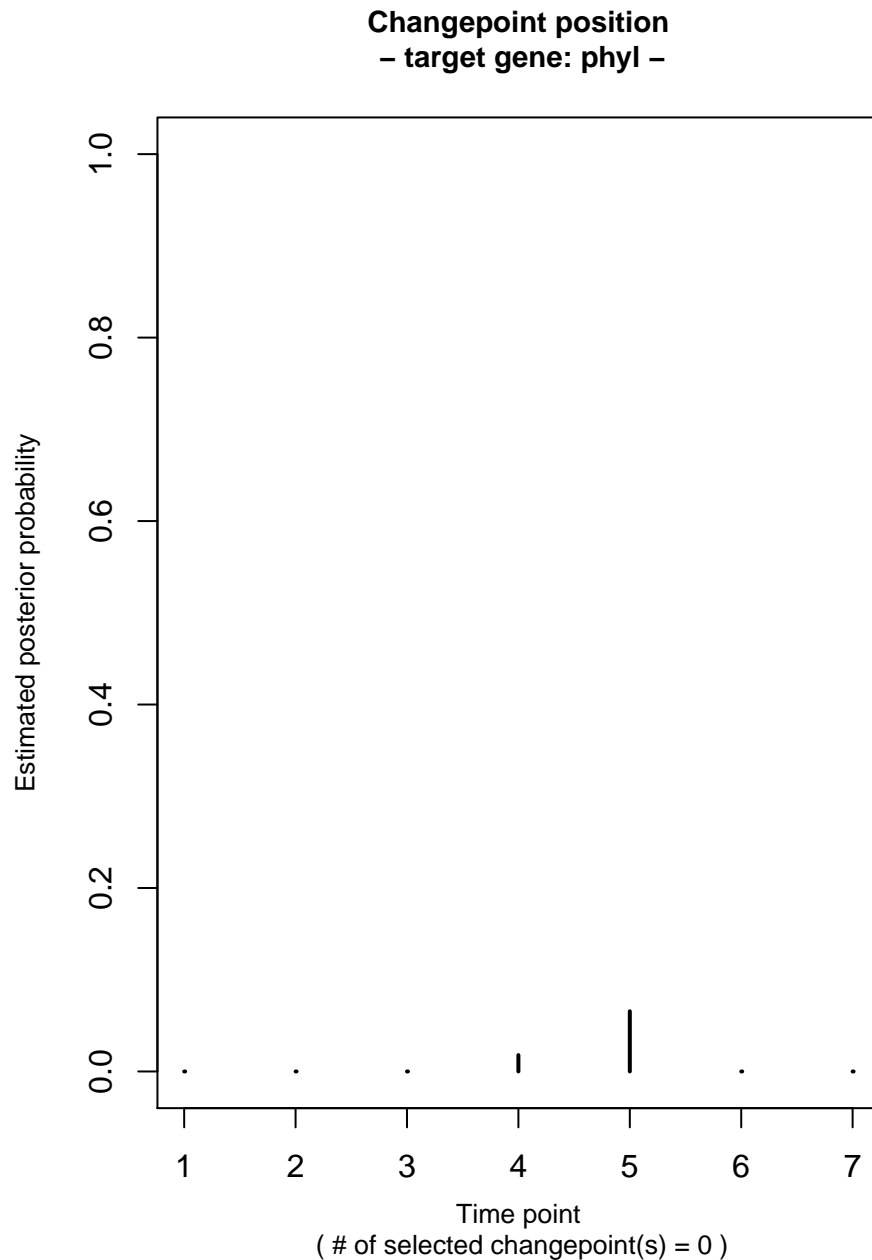
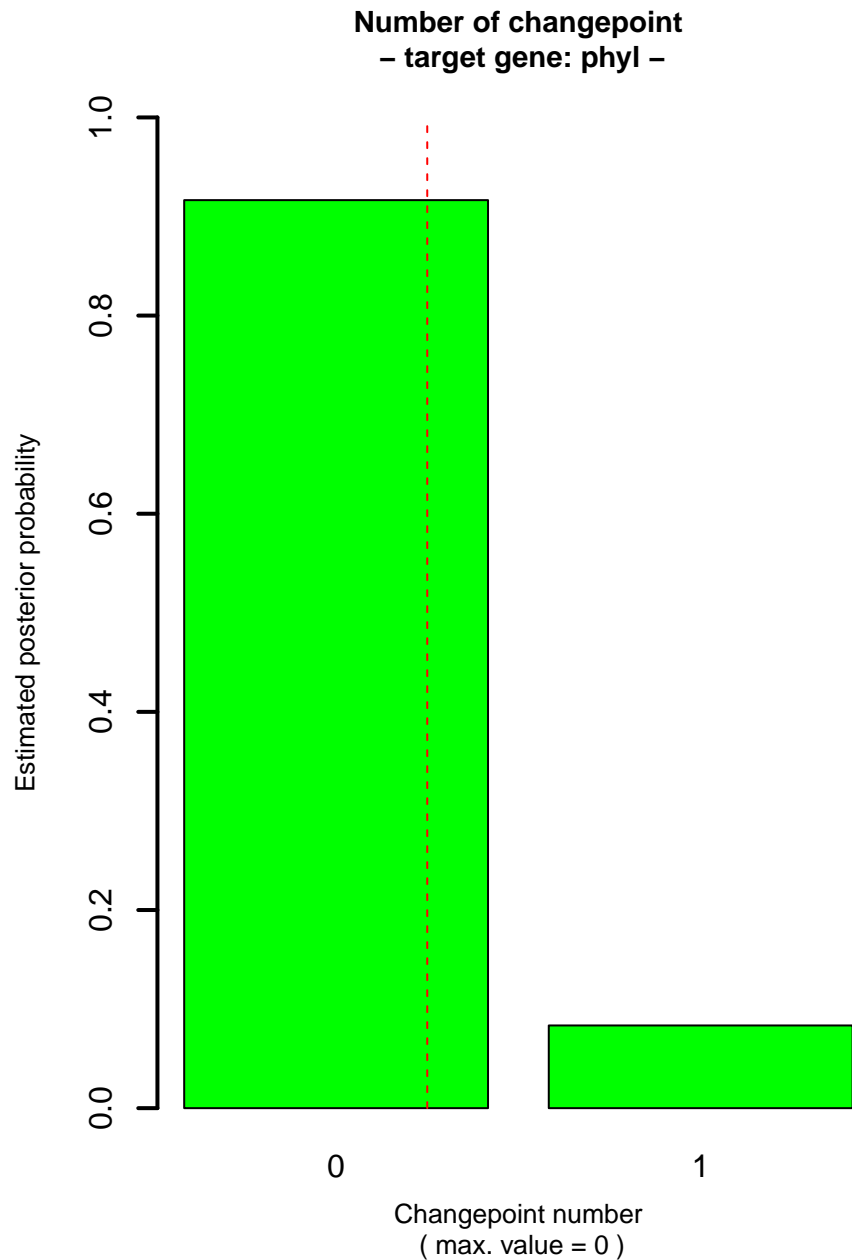
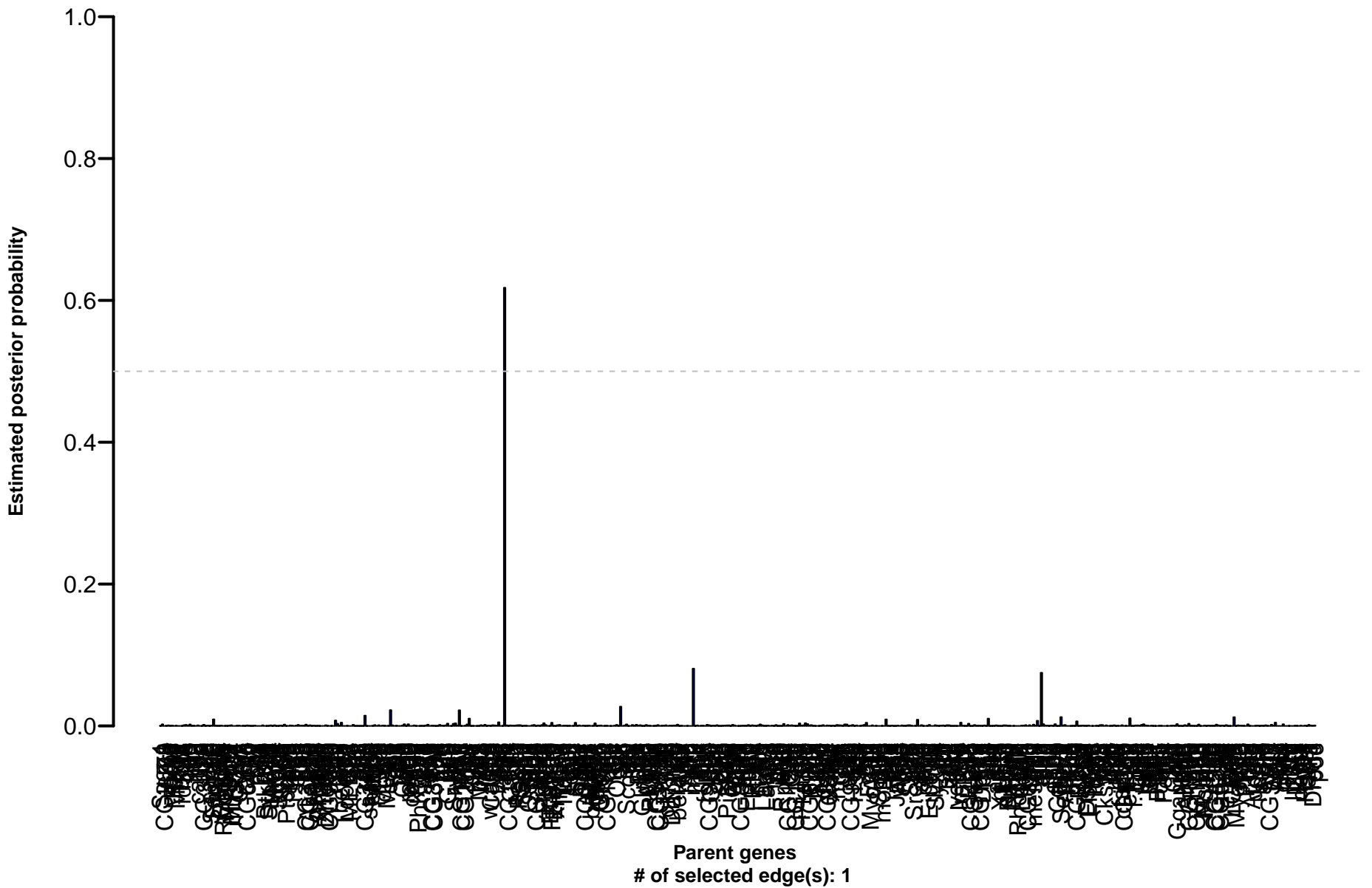


The graph displays the normalized concentration of the active form of the inhibitor, I_a , over a period of 30 time points. The concentration is represented by a blue line. It starts at a baseline of 1.0. At time point 3, it rises to 2.0 and remains constant until time point 12. It then drops back to 1.0 and remains constant until time point 17. At time point 18, it spikes to 2.2. It then drops back to 1.0 and remains constant until time point 23. Finally, it rises to 2.0 and remains constant until time point 30.

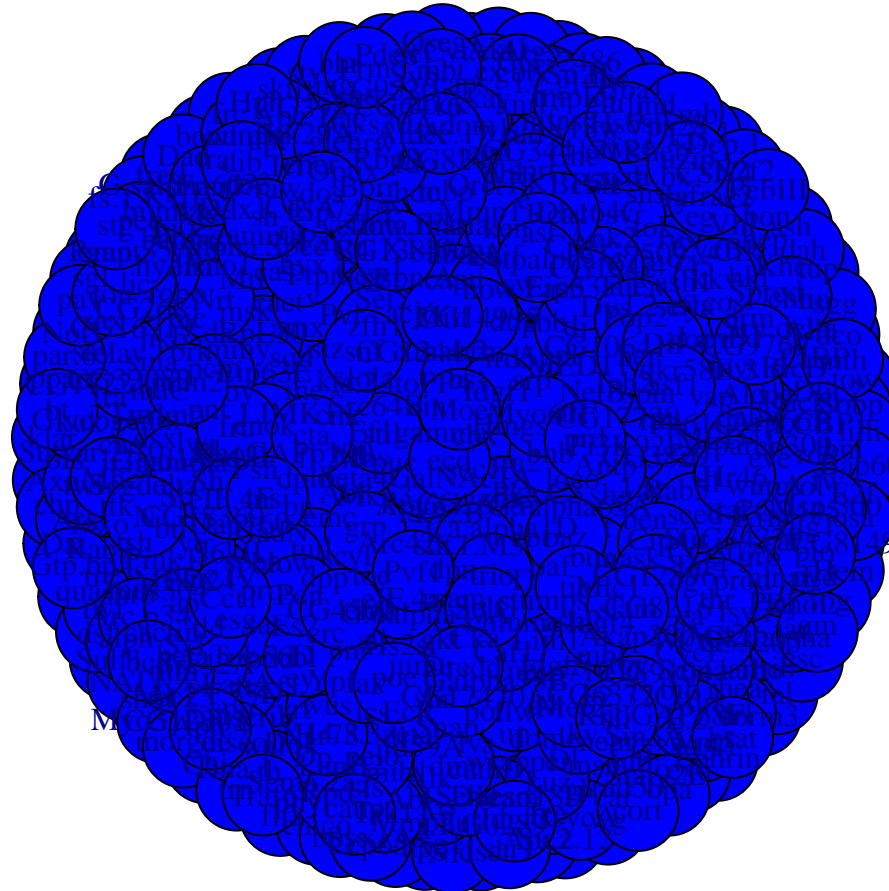
Time point	Normalized concentration of I_a
0	1.0
1	1.0
2	1.0
3	2.0
4	2.0
5	2.0
6	2.0
7	2.0
8	2.0
9	2.0
10	2.0
11	2.0
12	2.0
13	1.0
14	1.0
15	1.0
16	1.0
17	1.0
18	2.2
19	1.0
20	1.0
21	1.0
22	1.0
23	1.0
24	2.0
25	2.0
26	2.0
27	2.0
28	2.0
29	2.0
30	2.0



Regulatory model for target gene: phyl
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
550	1	0.6179	2	6	+