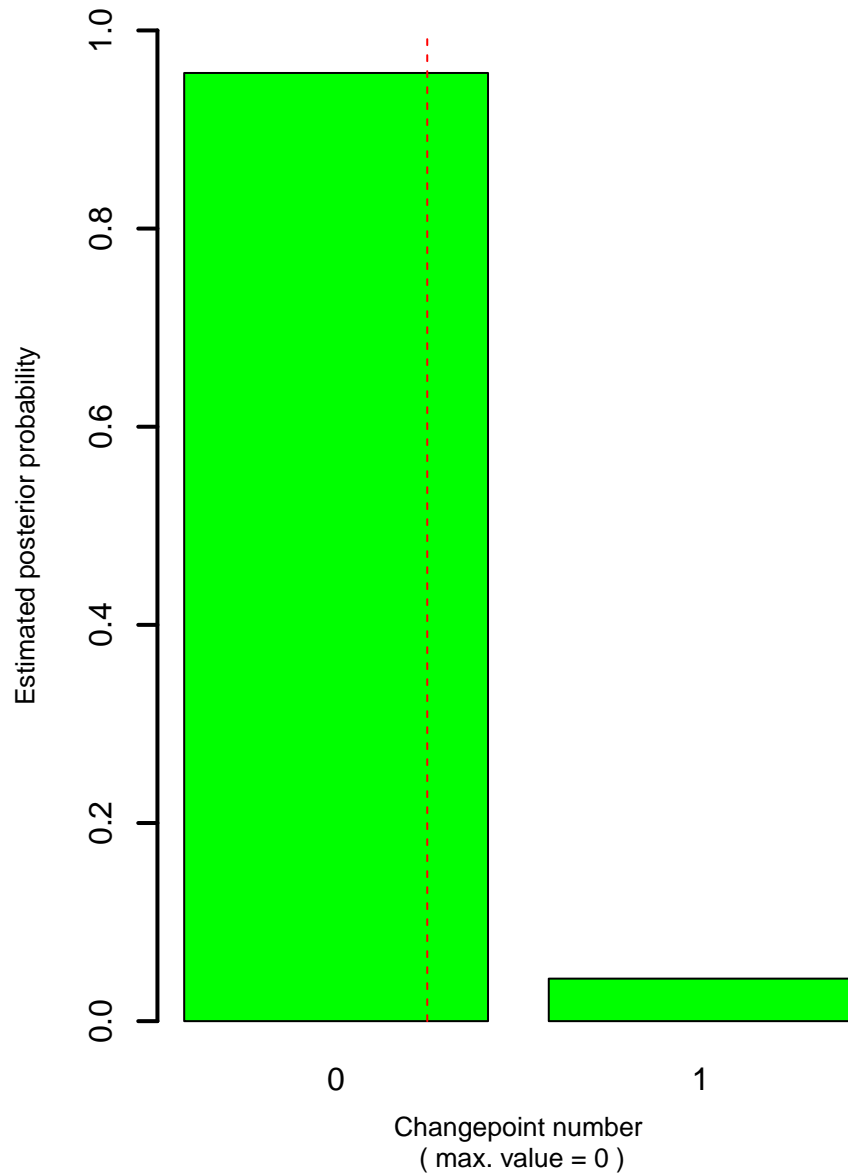


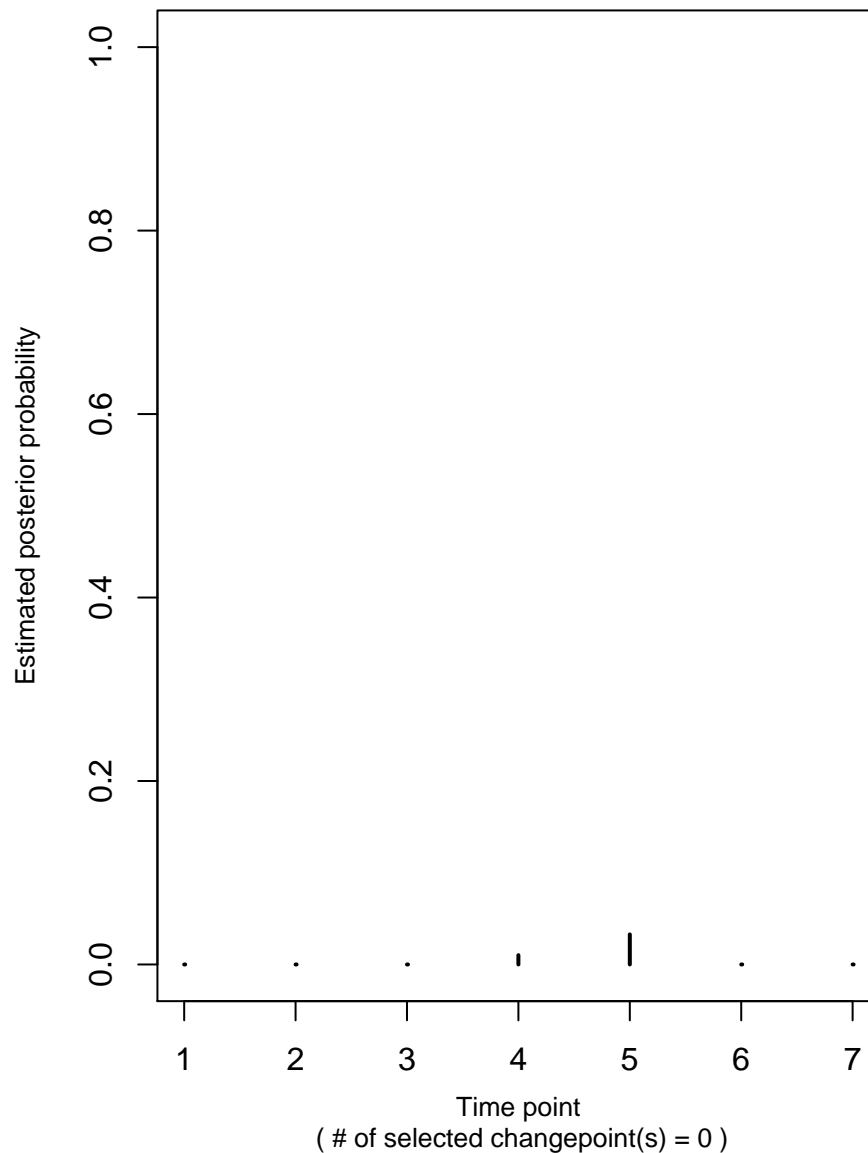
The graph displays the normalized concentration of the active form of the inhibitor over 30 time points. The concentration starts at 2.0, drops to 1.0 at time point 13, peaks at 2.0 at time point 14, drops to 1.0 at time point 15, rises to 2.0 at time point 19, drops to 1.0 at time point 21, and returns to 2.0 at time point 23, remaining constant thereafter.

Time point	Normalized concentration
1	2.0
2	2.0
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	2.0
15	1.0
16	1.0
17	1.0
18	1.0
19	2.0
20	2.0
21	1.0
22	1.0
23	2.0
24	2.0
25	2.0
26	2.0
27	2.0
28	2.0
29	2.0
30	2.0

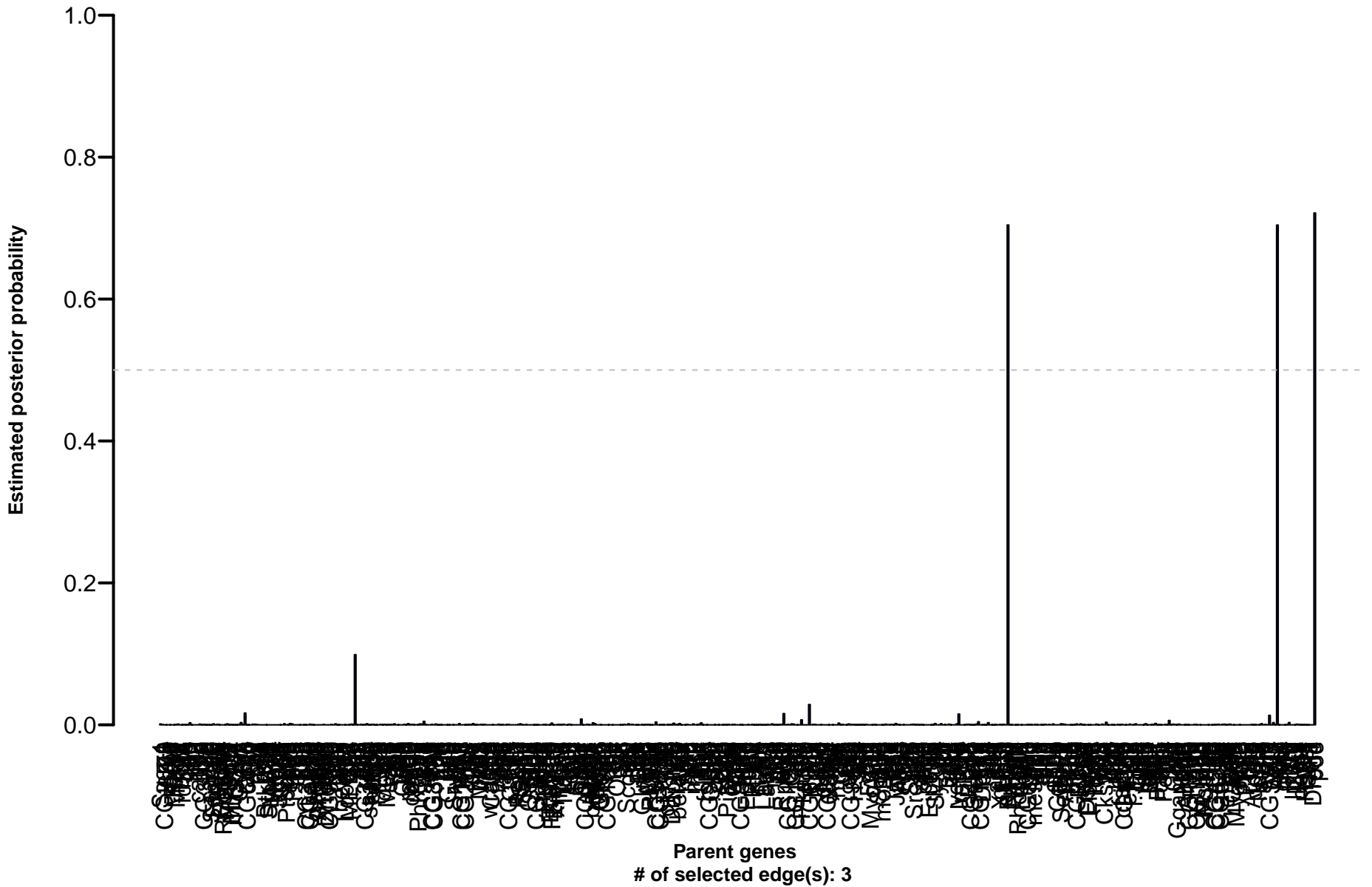
Number of changepoint
– target gene: sgl –



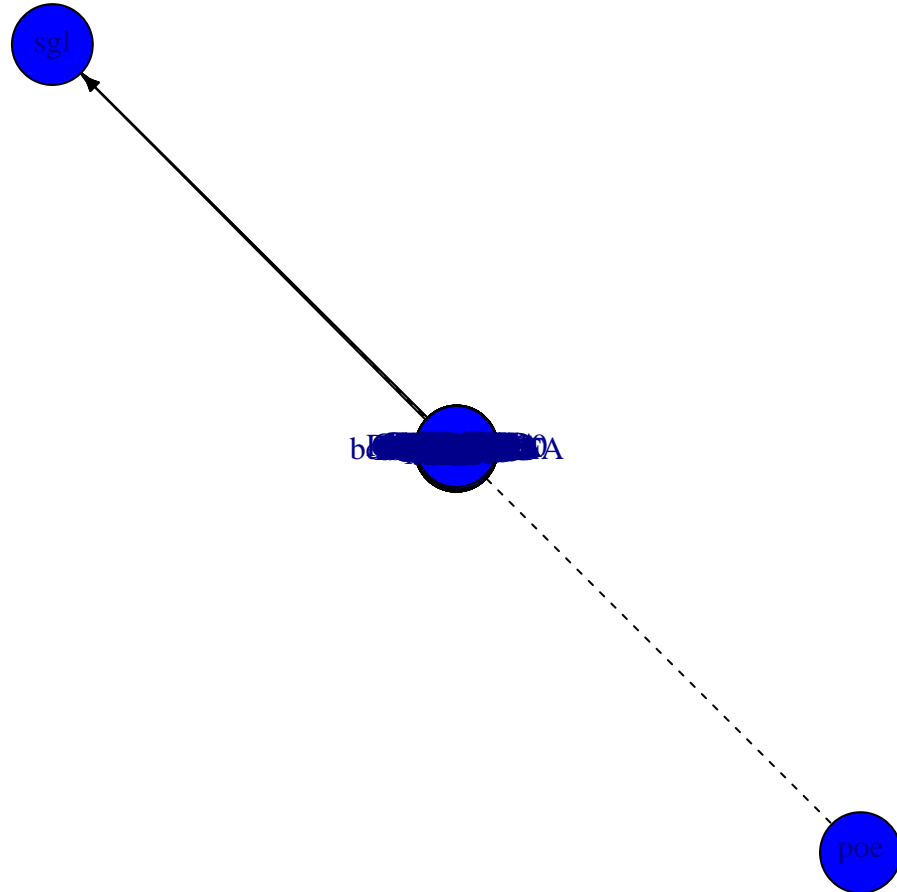
Changepoint position
– target gene: sgl –



Regulatory model for target gene: *sgl*
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
415	1	0.7213	2	6	-
40	1	0.7044	2	6	+
264	1	0.7043	2	6	-