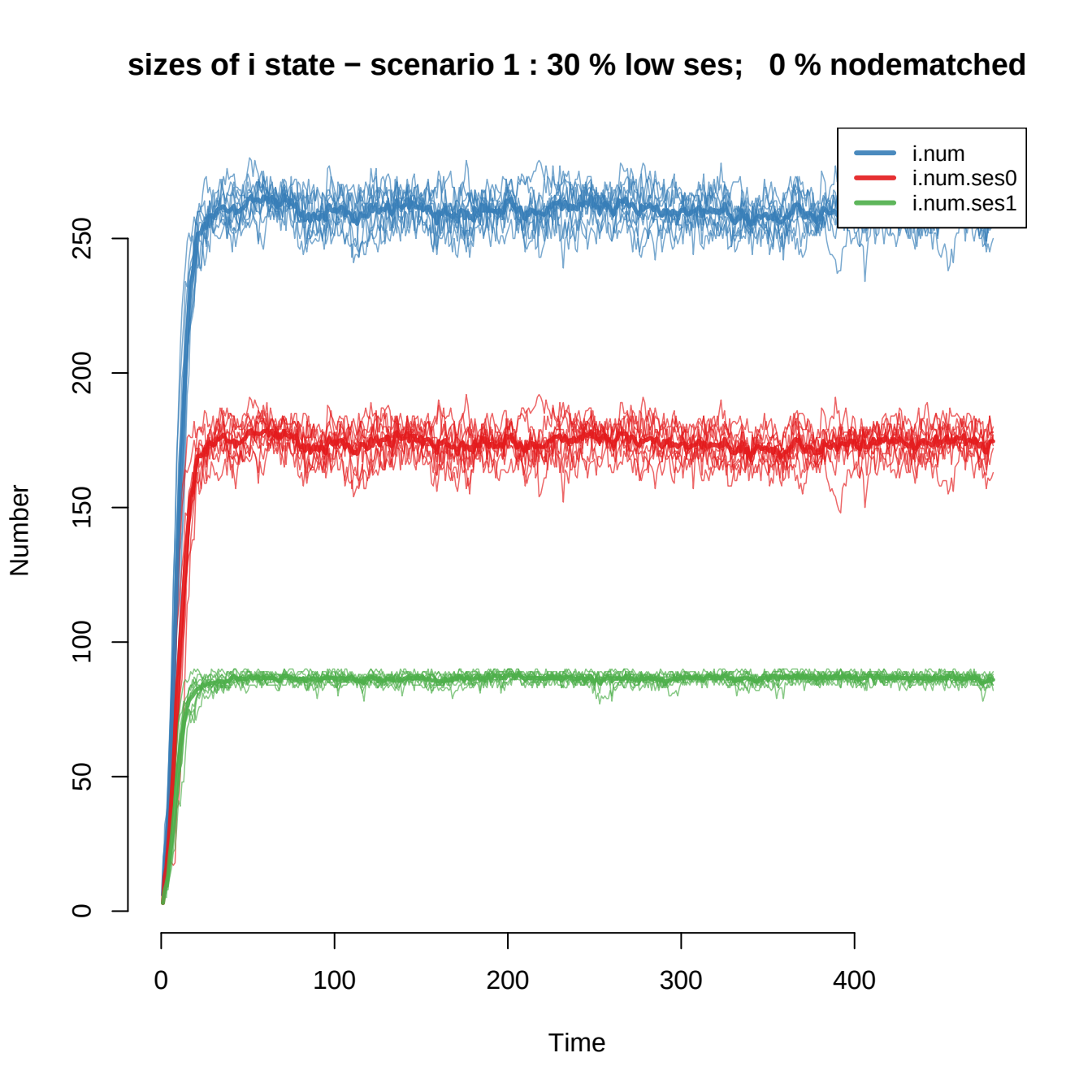
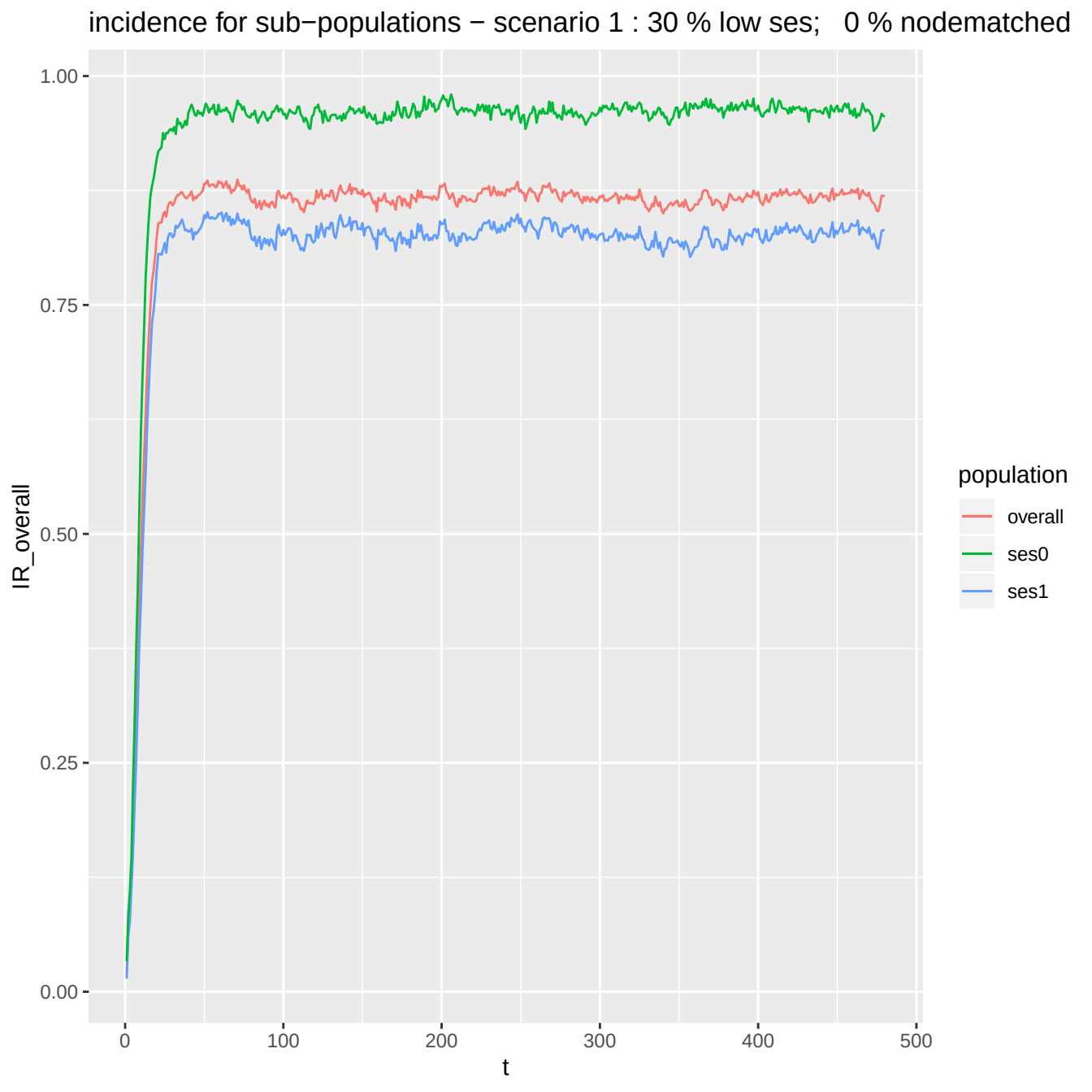
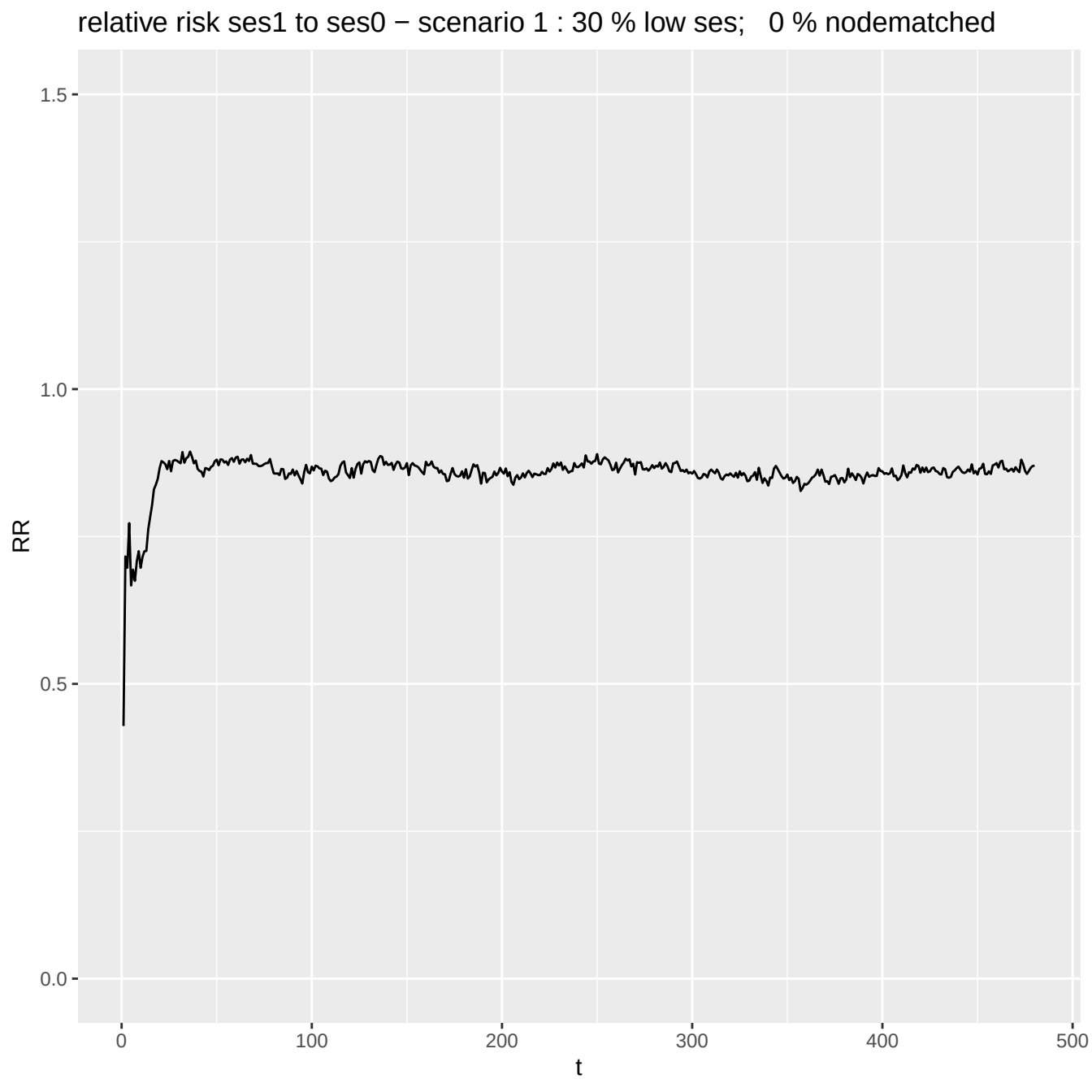
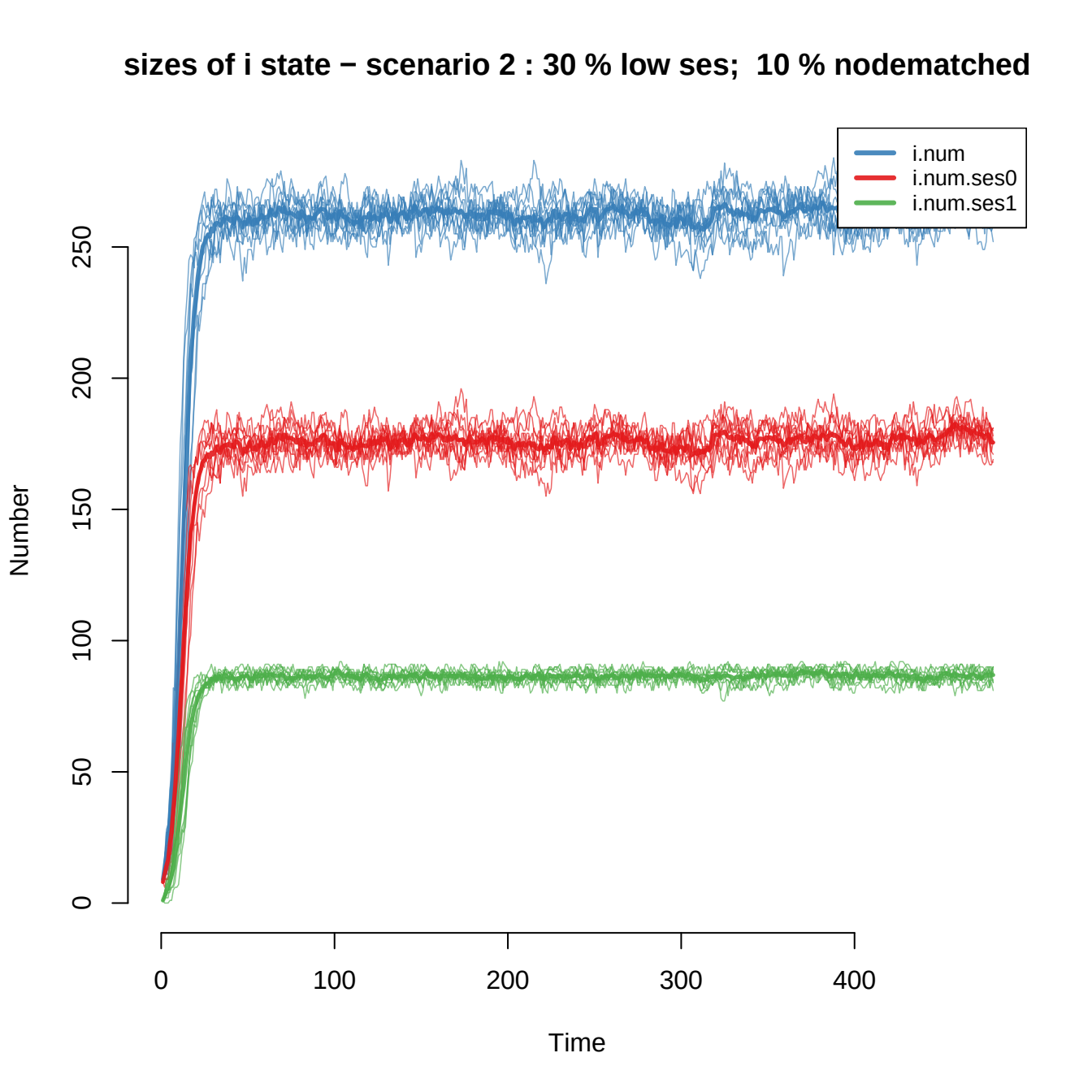
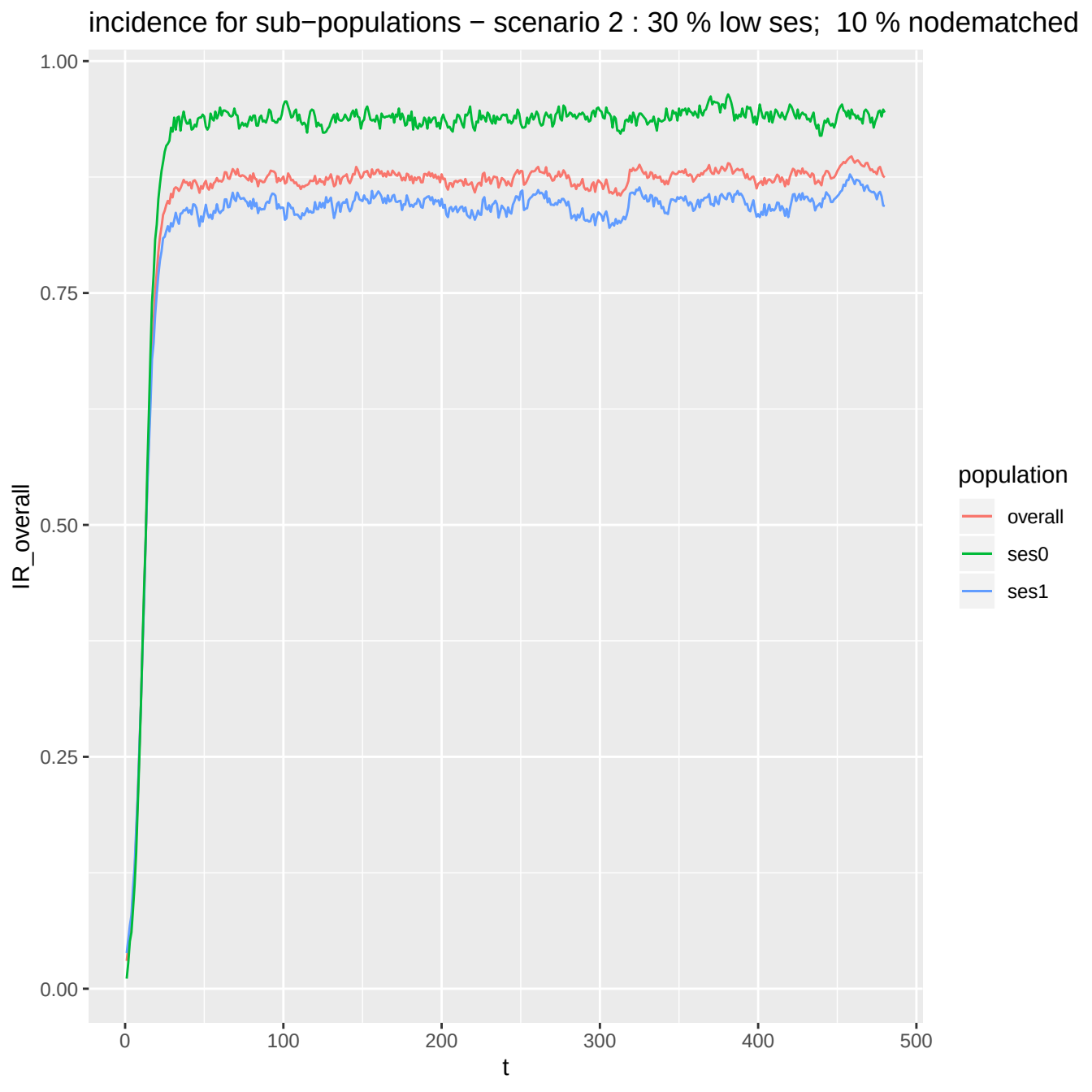
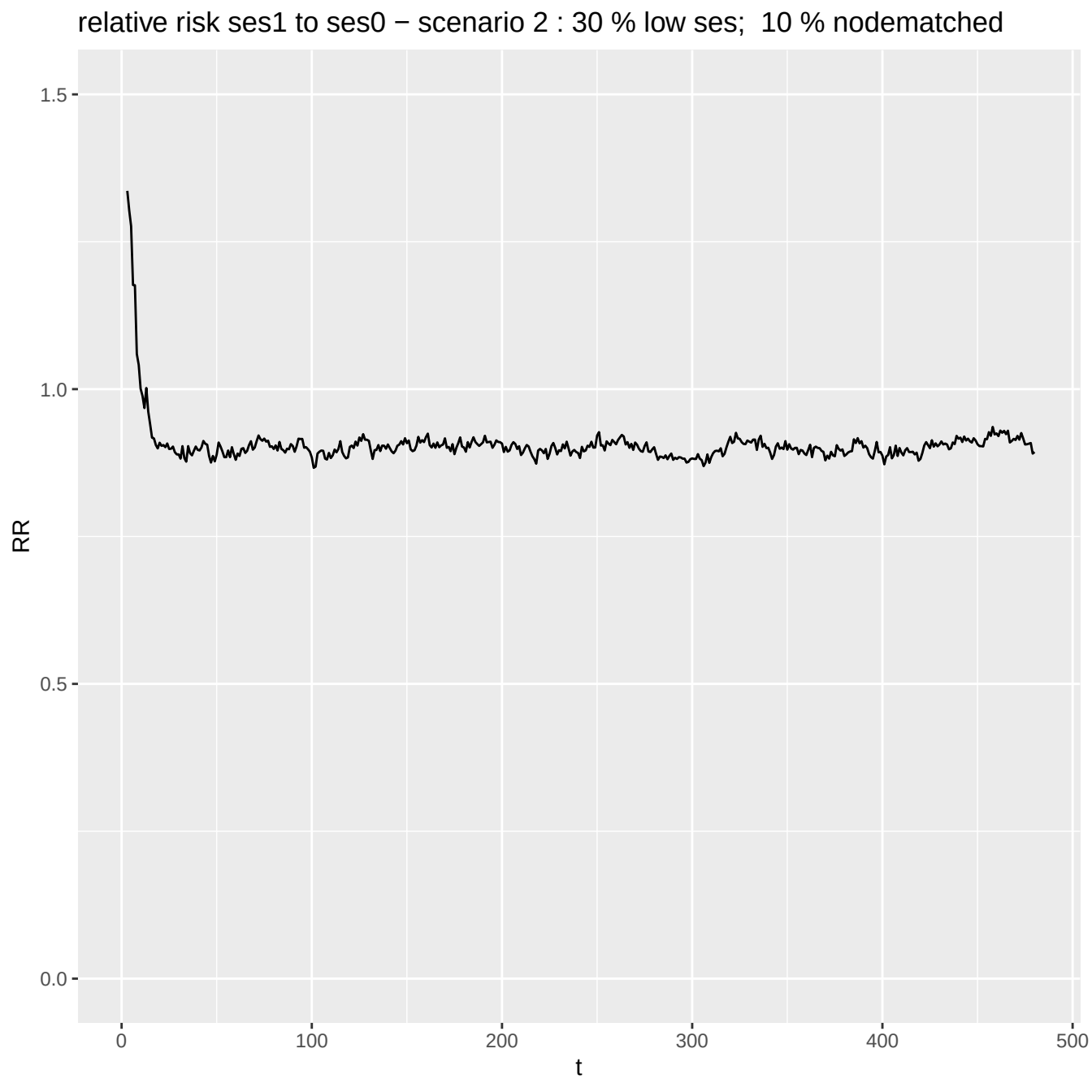

sizes of i state − scenario 1 : 30 % low ses; 0 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
50
100
150
200
250
0
100
200
300
400
Time

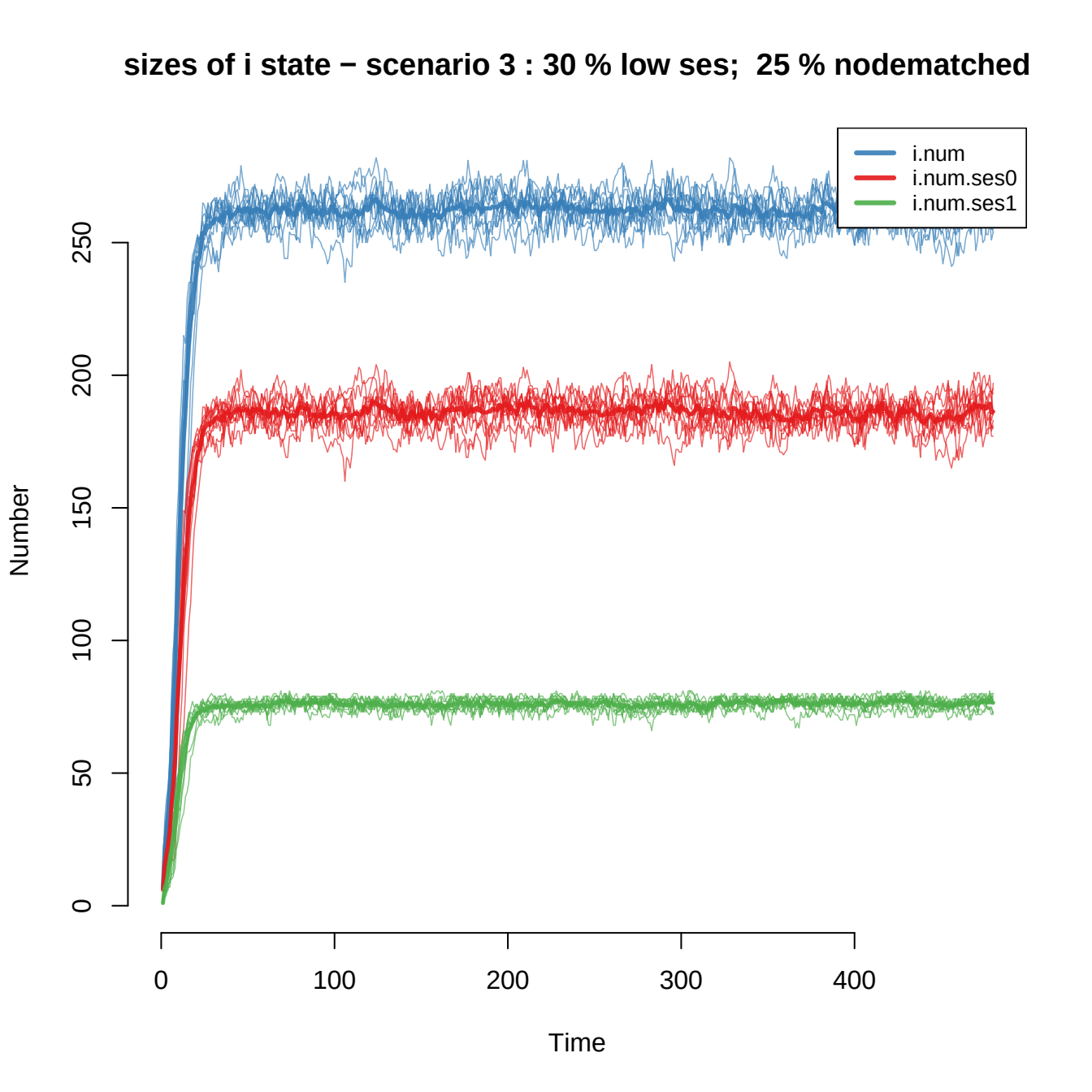

incidence for sub−populations − scenario 1 : 30 % low ses; 0 % nodematched
IR_overall
1.00
0.75
0.50
0.25
0.00
0
100
200
300
400
500
t
population
overall
ses0
ses1

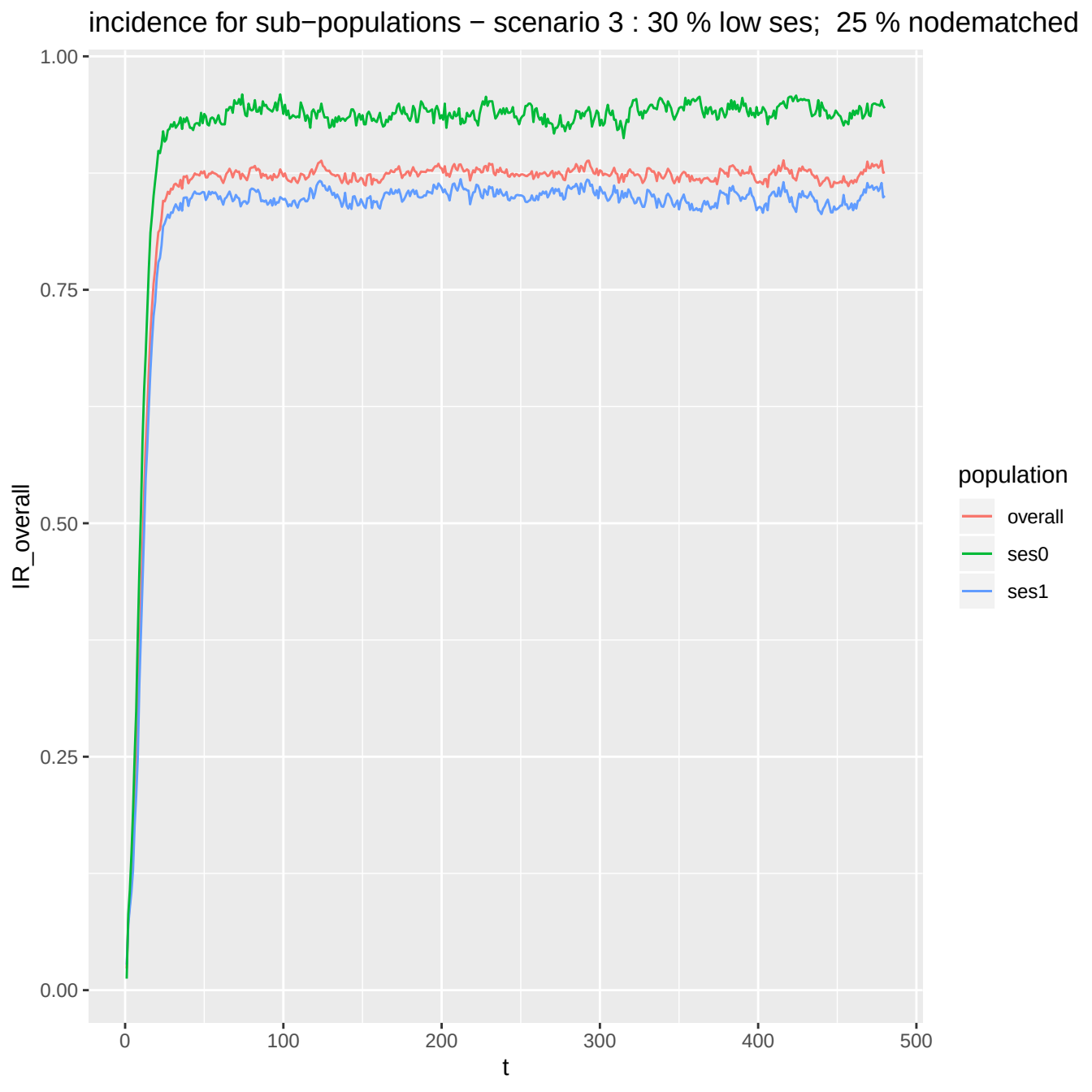

relative risk ses1 to ses0 − scenario 1 : 30 % low ses; 0 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

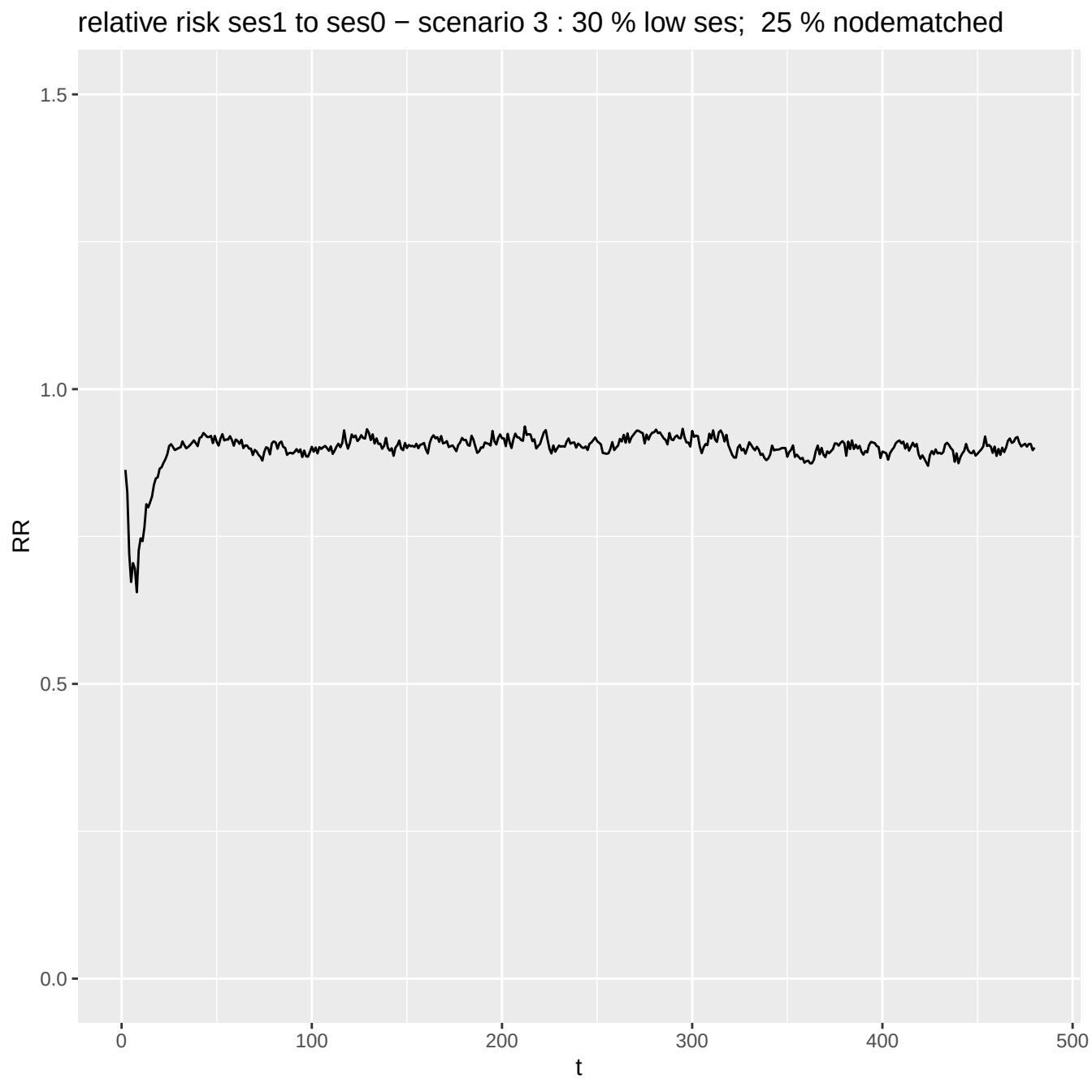

sizes of i state − scenario 2 : 30 % low ses; 10 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
50
100
150
200
250
0
100
200
300
400
Time

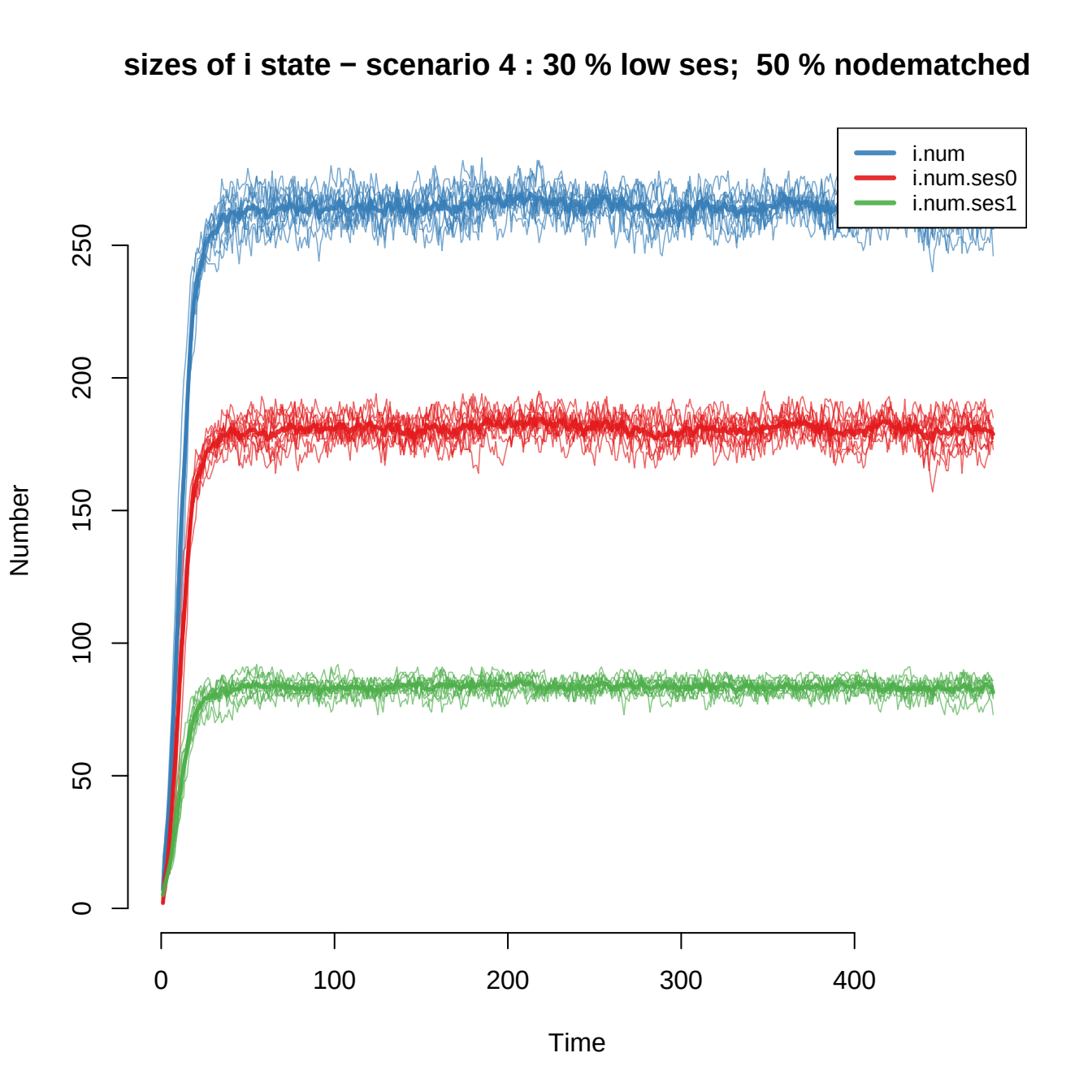

incidence for sub−populations − scenario 2 : 30 % low ses; 10 % nodematched
IR_overall
1.00
0.75
0.50
0.25
0.00
0
100
200
300
400
500
t
population
overall
ses0
ses1

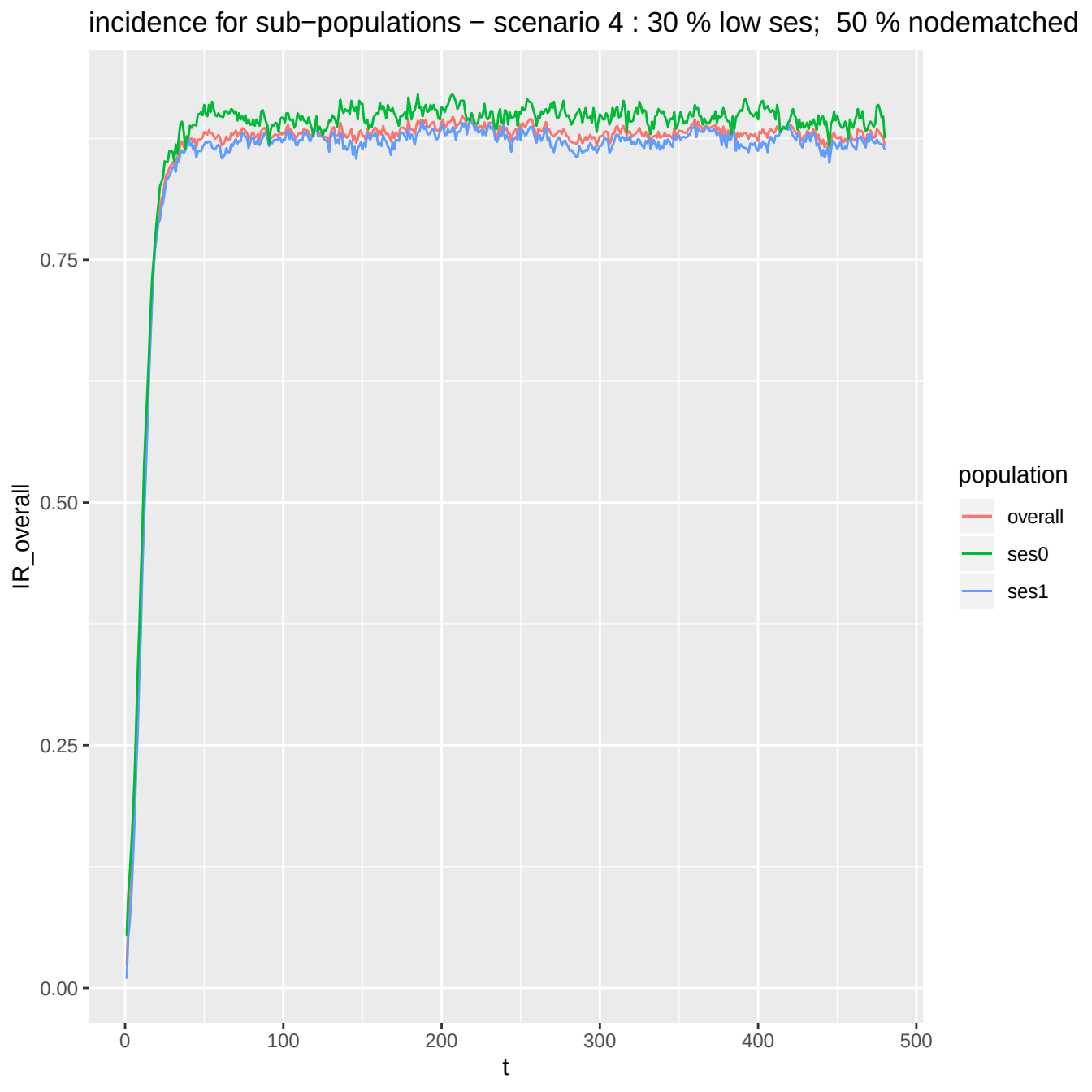

relative risk ses1 to ses0 − scenario 2 : 30 % low ses; 10 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

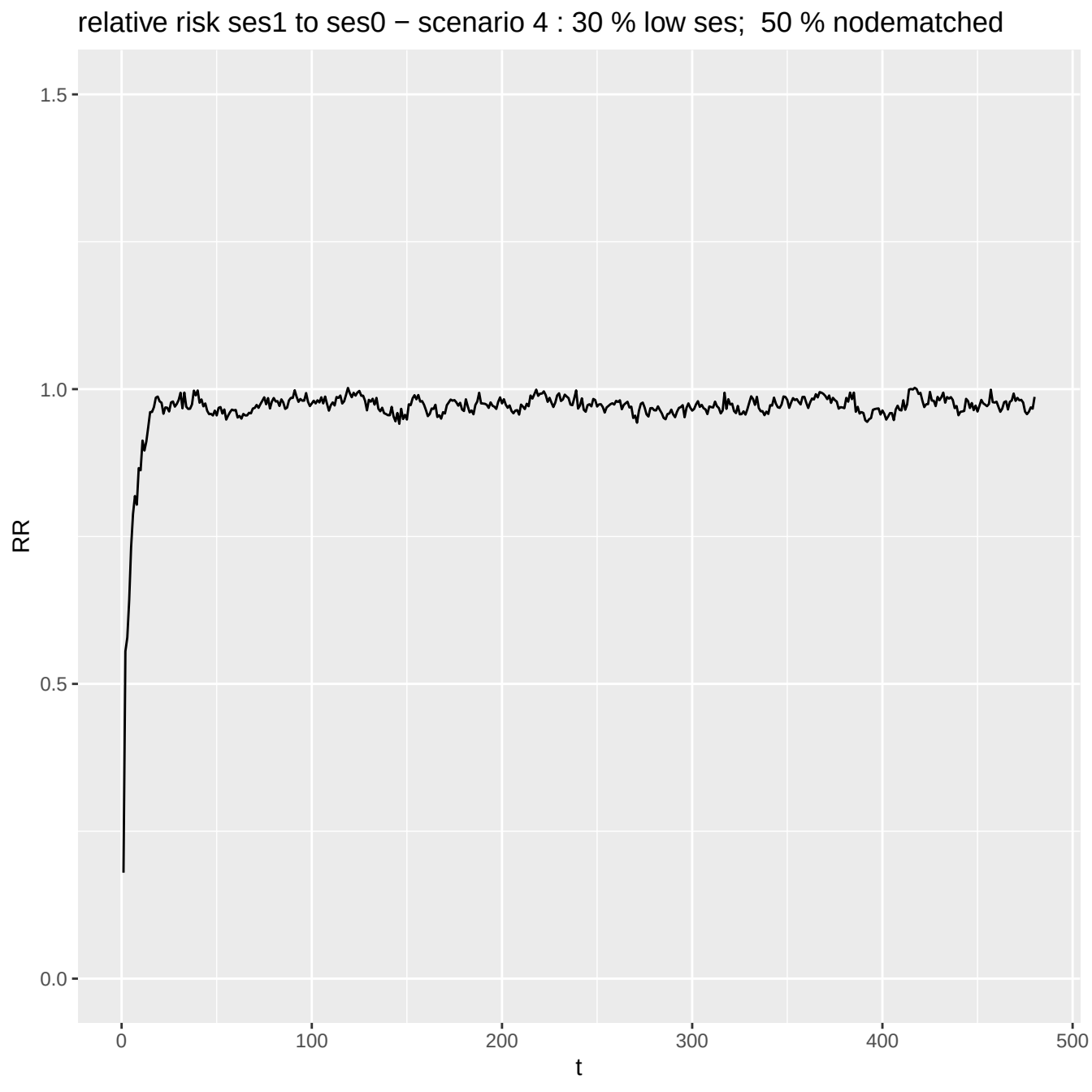

sizes of i state − scenario 3 : 30 % low ses; 25 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
50
100
150
200
250
0
100
200
300
400
Time

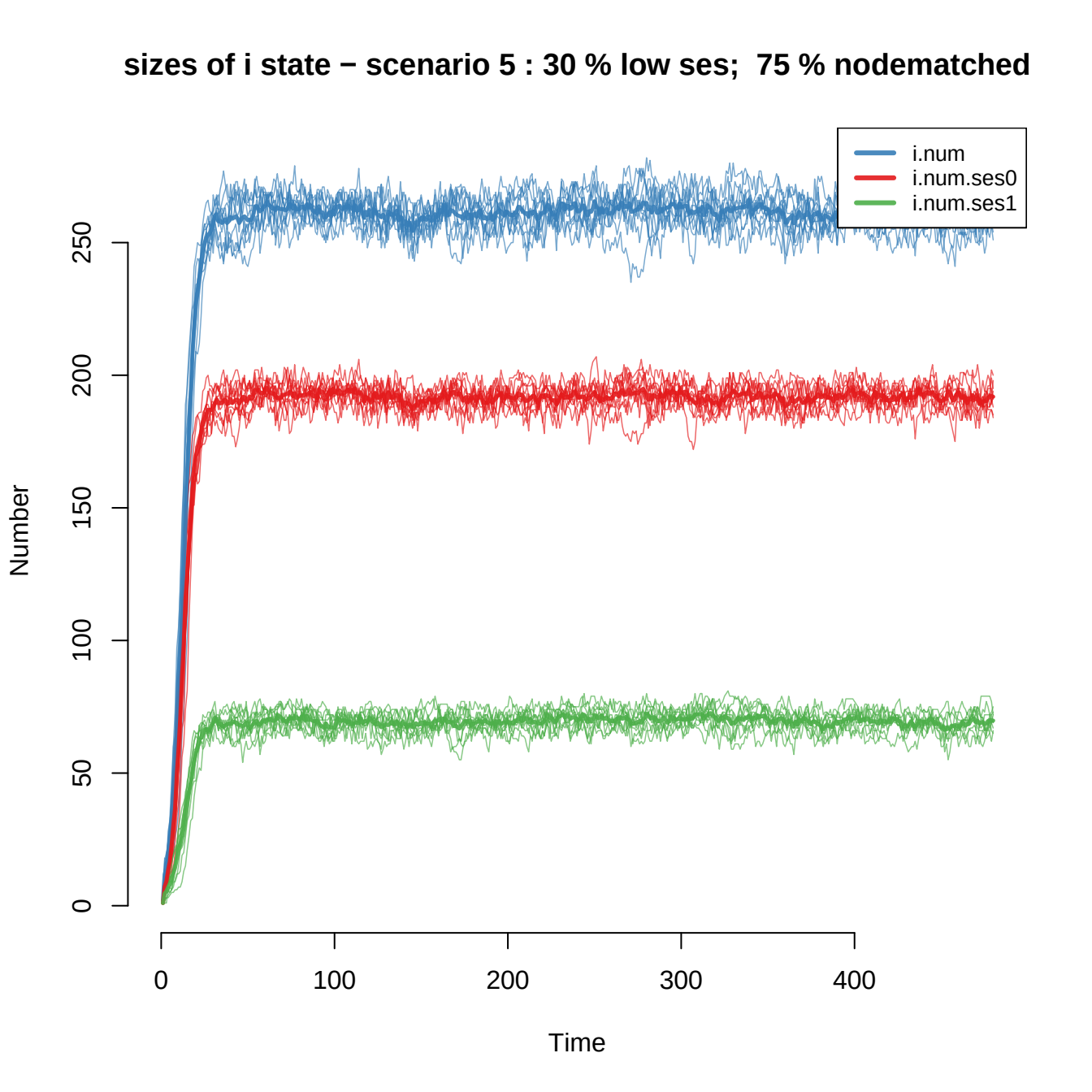

incidence for sub−populations − scenario 3 : 30 % low ses; 25 % nodematched
IR_overall
1.00
0.75
0.50
0.25
0.00
0
100
200
300
400
500
t
population
overall
ses0
ses1

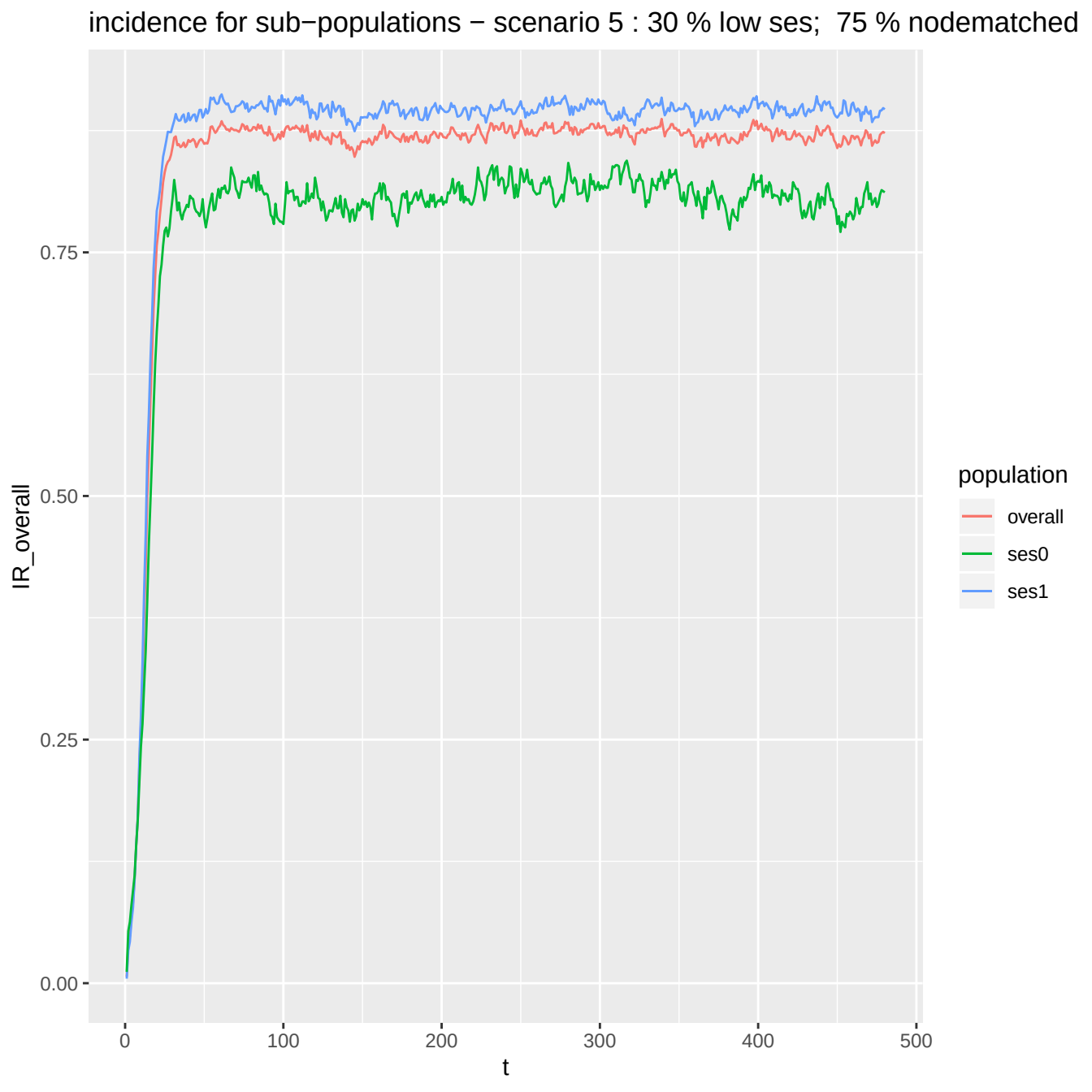

relative risk ses1 to ses0 − scenario 3 : 30 % low ses; 25 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

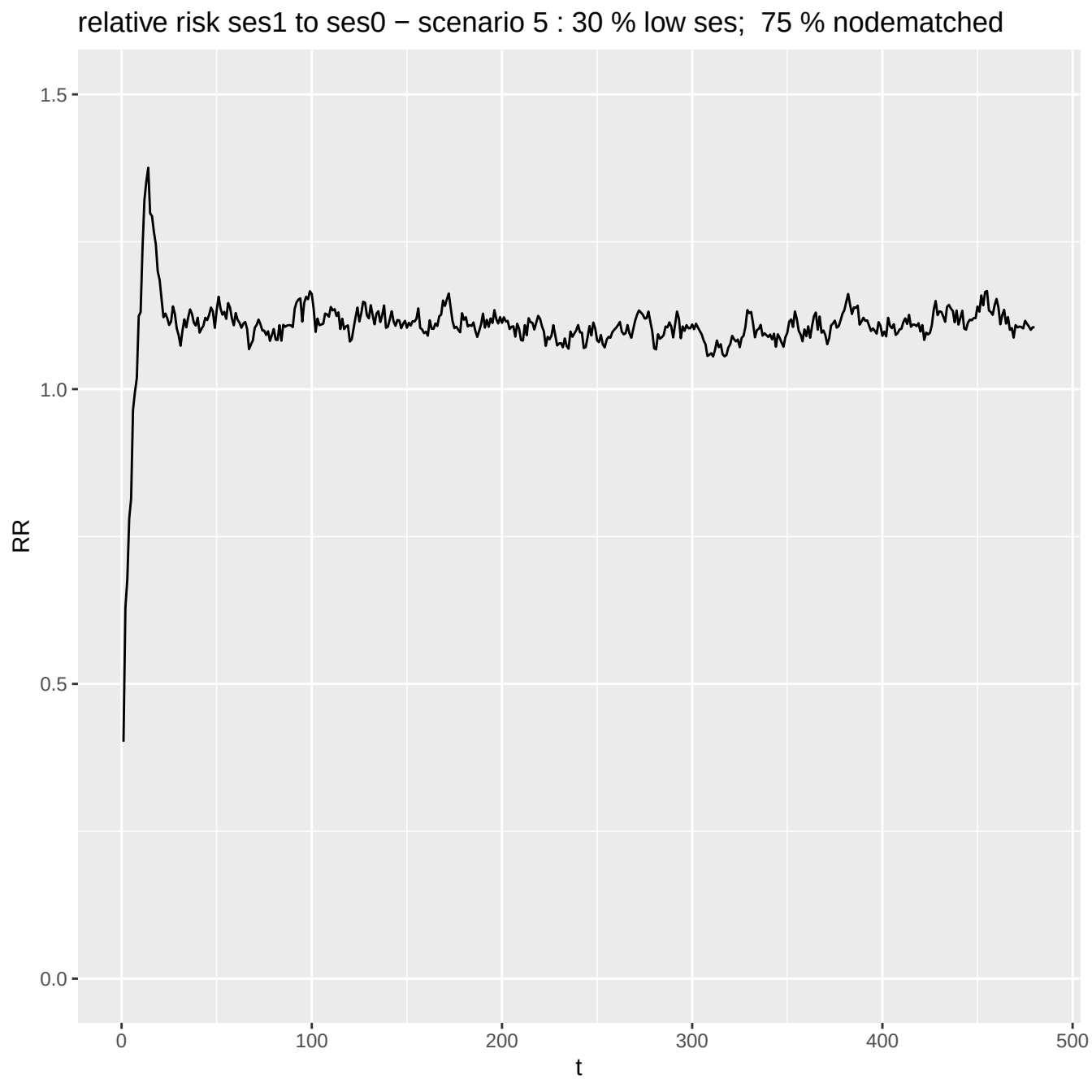

sizes of i state − scenario 4 : 30 % low ses; 50 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
50
100
150
200
250
0
100
200
300
400
Time

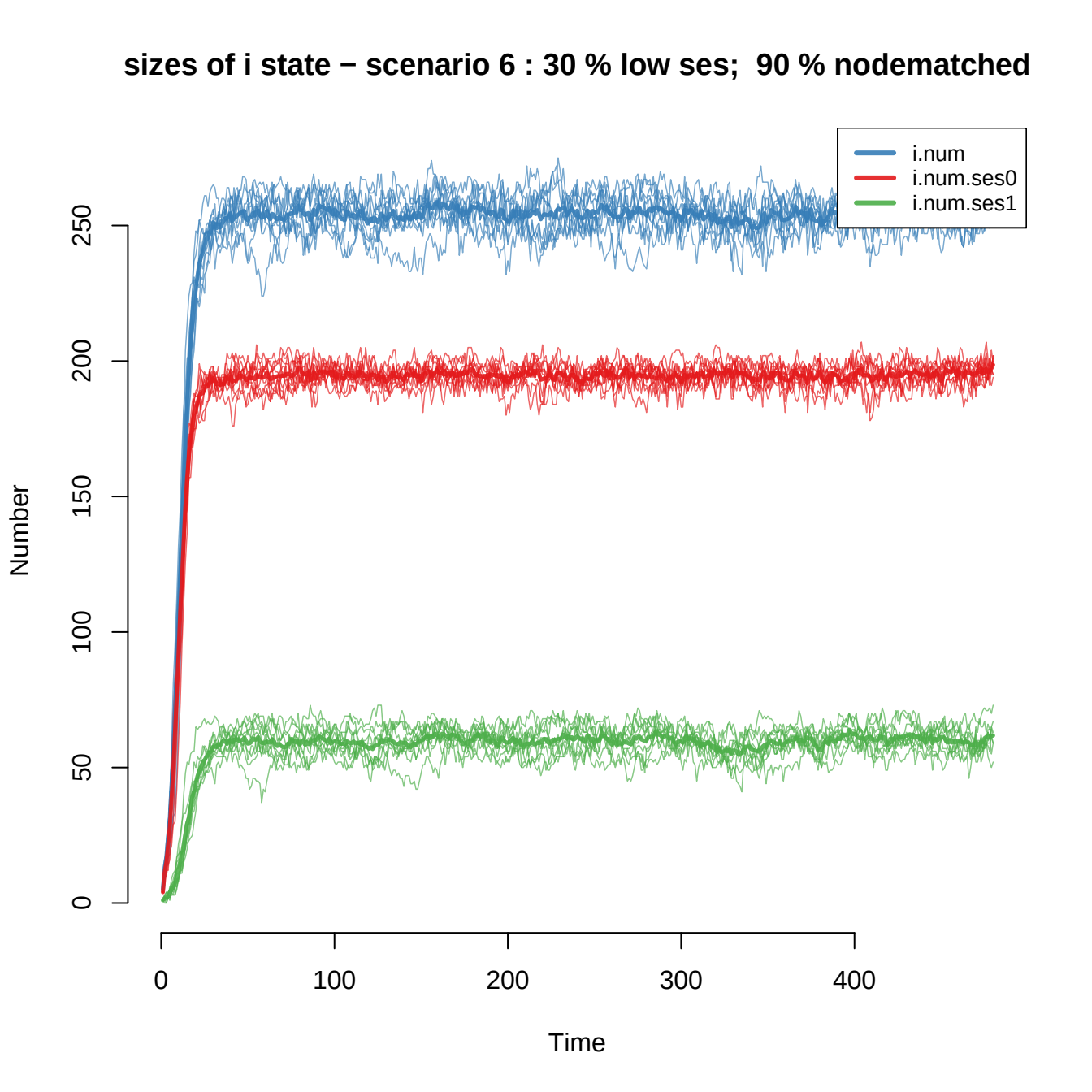

incidence for sub−populations − scenario 4 : 30 % low ses; 50 % nodematched
IR_overall
0.00
0.25
0.50
0.75
0
100
200
300
400
500
t
population
overall
ses0
ses1

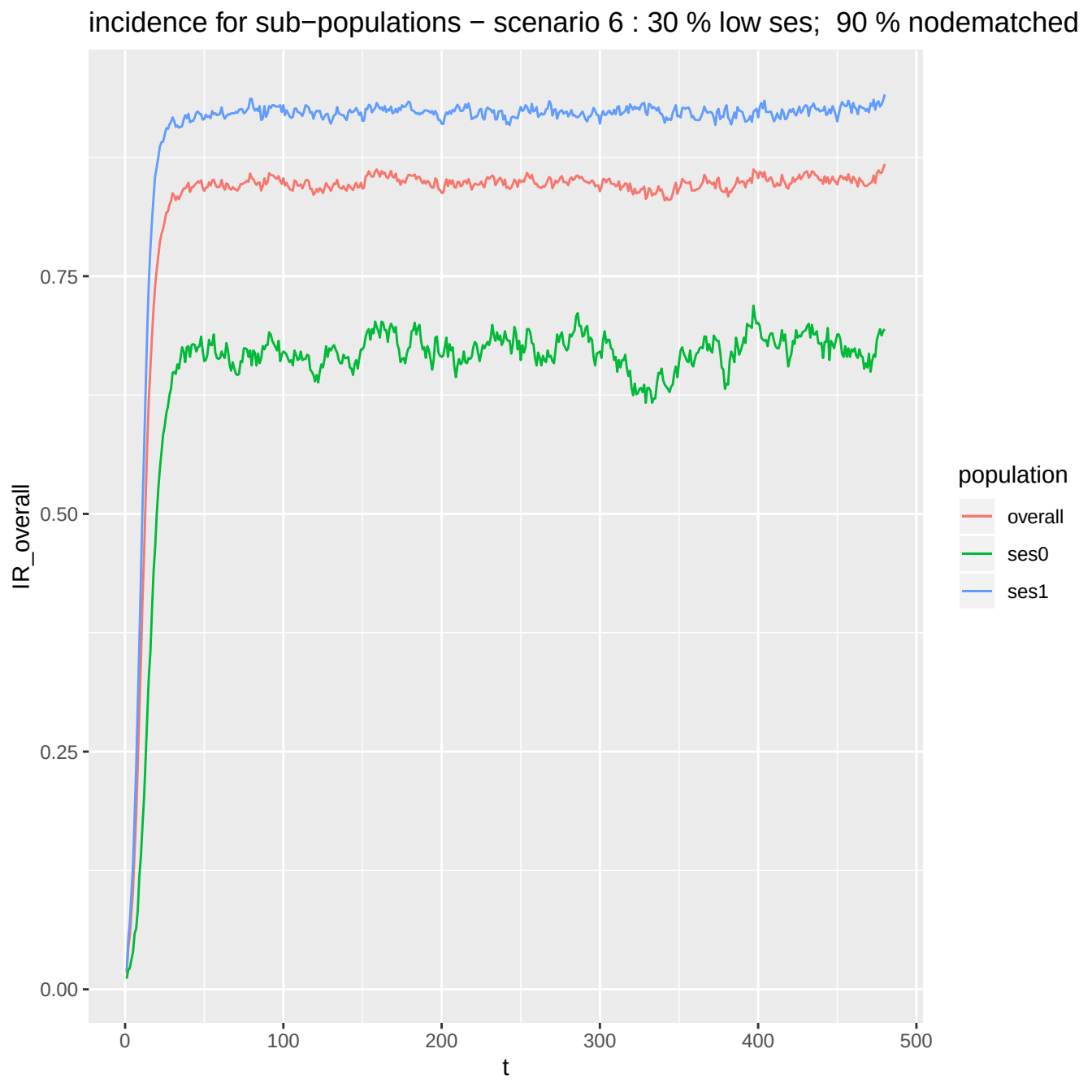

relative risk ses1 to ses0 − scenario 4 : 30 % low ses; 50 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

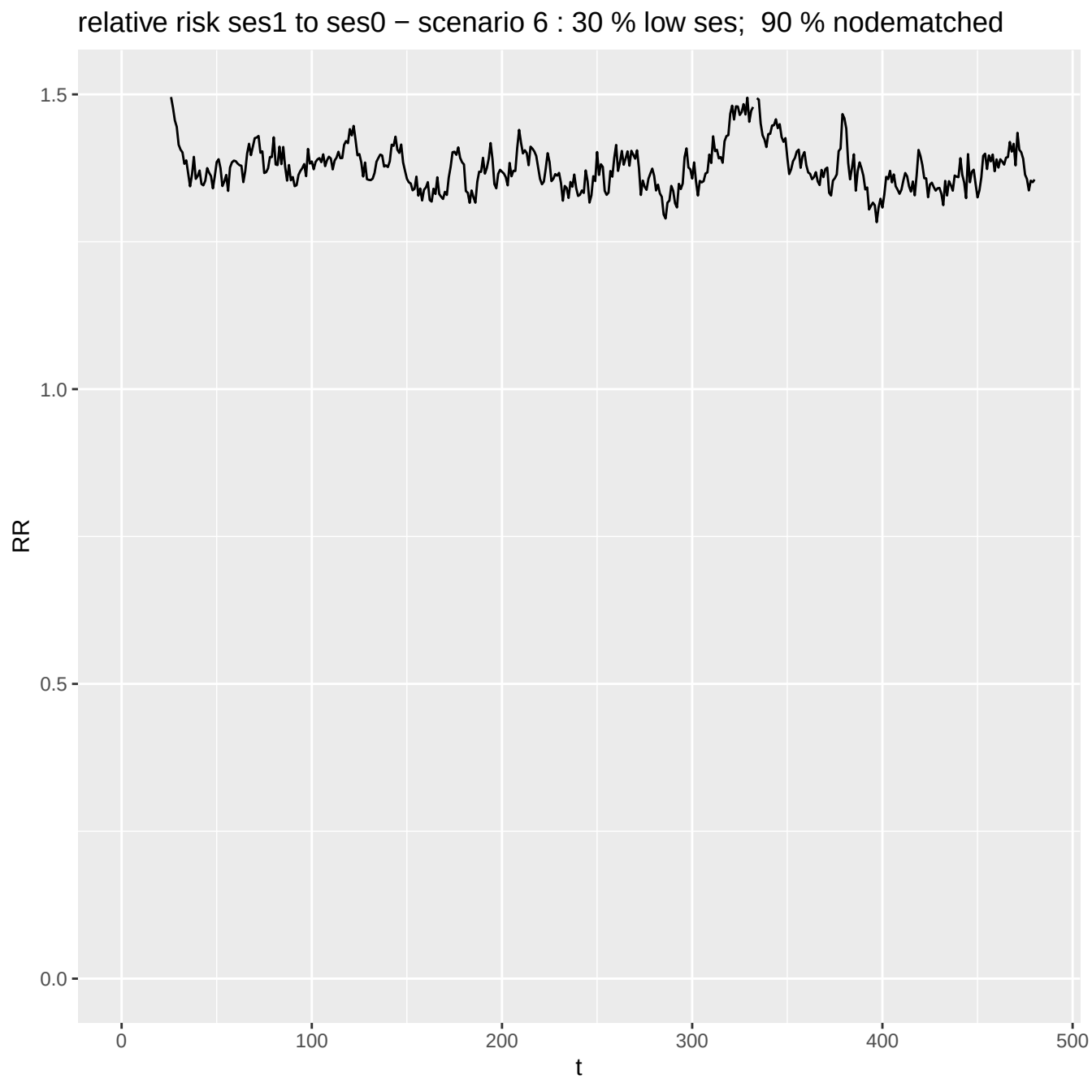

sizes of i state − scenario 5 : 30 % low ses; 75 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
50
100
150
200
250
0
100
200
300
400
Time

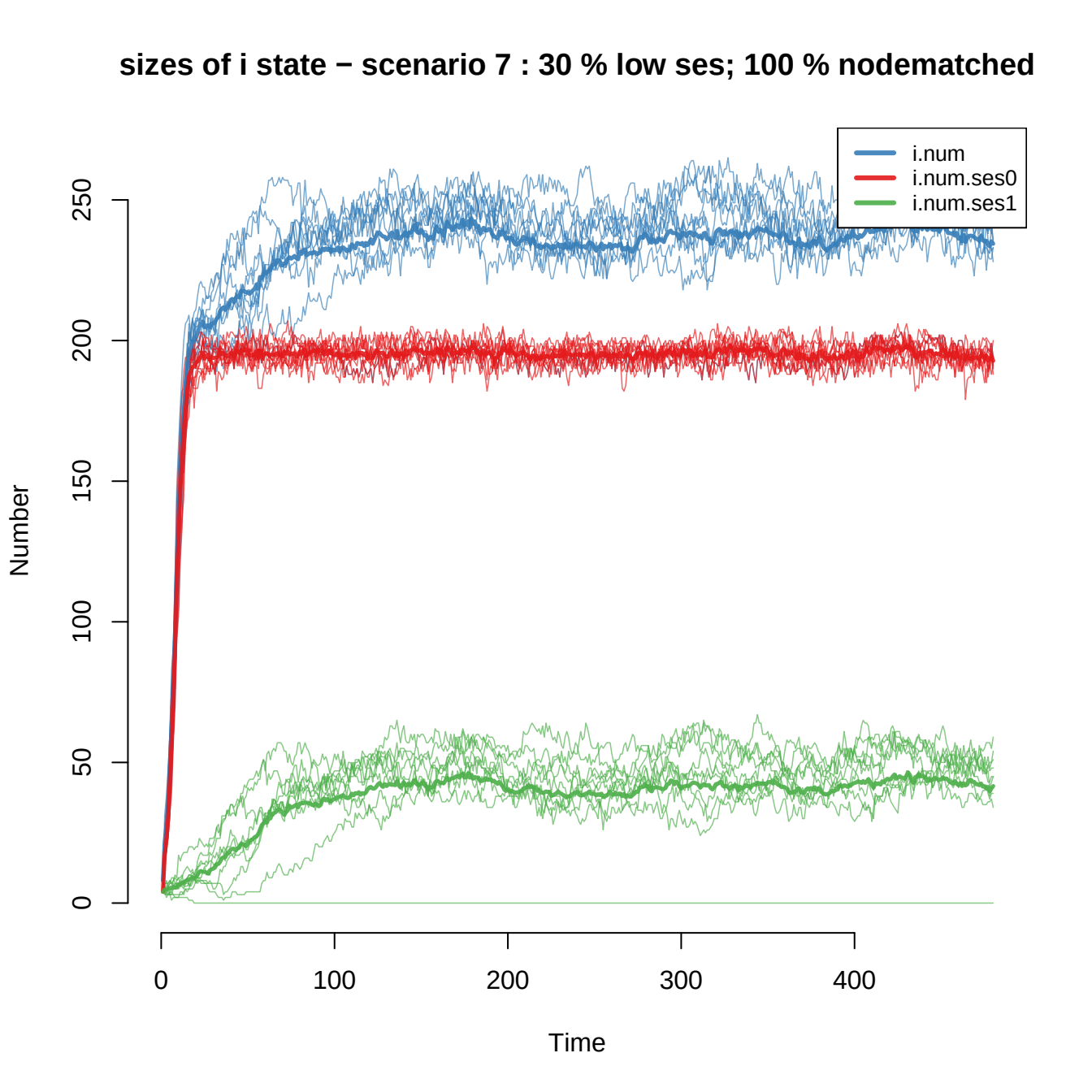

incidence for sub−populations − scenario 5 : 30 % low ses; 75 % nodematched
IR_overall
0.00
0.25
0.50
0.75
0
100
200
300
400
500
t
population
overall
ses0
ses1

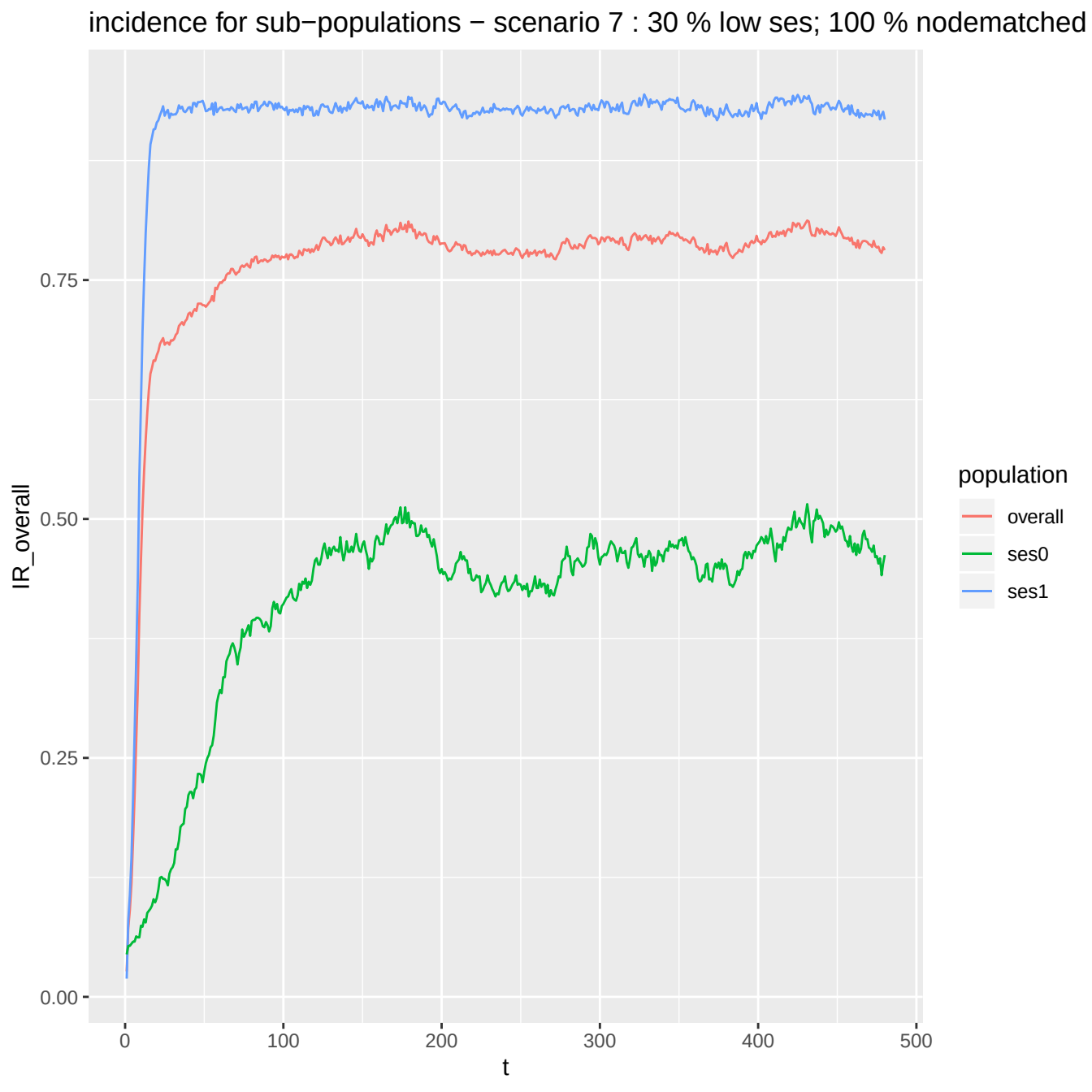

relative risk ses1 to ses0 − scenario 5 : 30 % low ses; 75 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

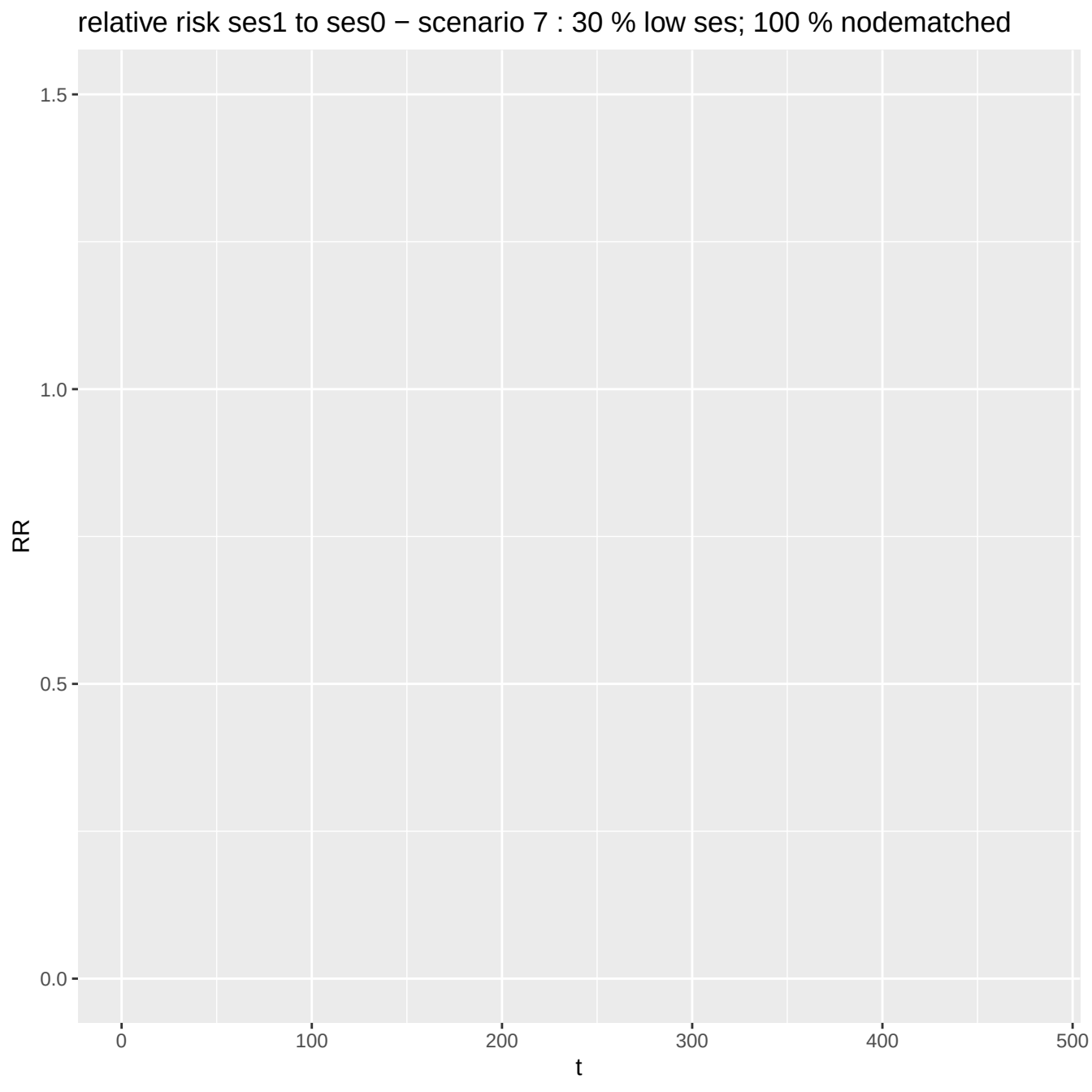

sizes of i state − scenario 6 : 30 % low ses; 90 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
50
100
150
200
250
0
100
200
300
400
Time

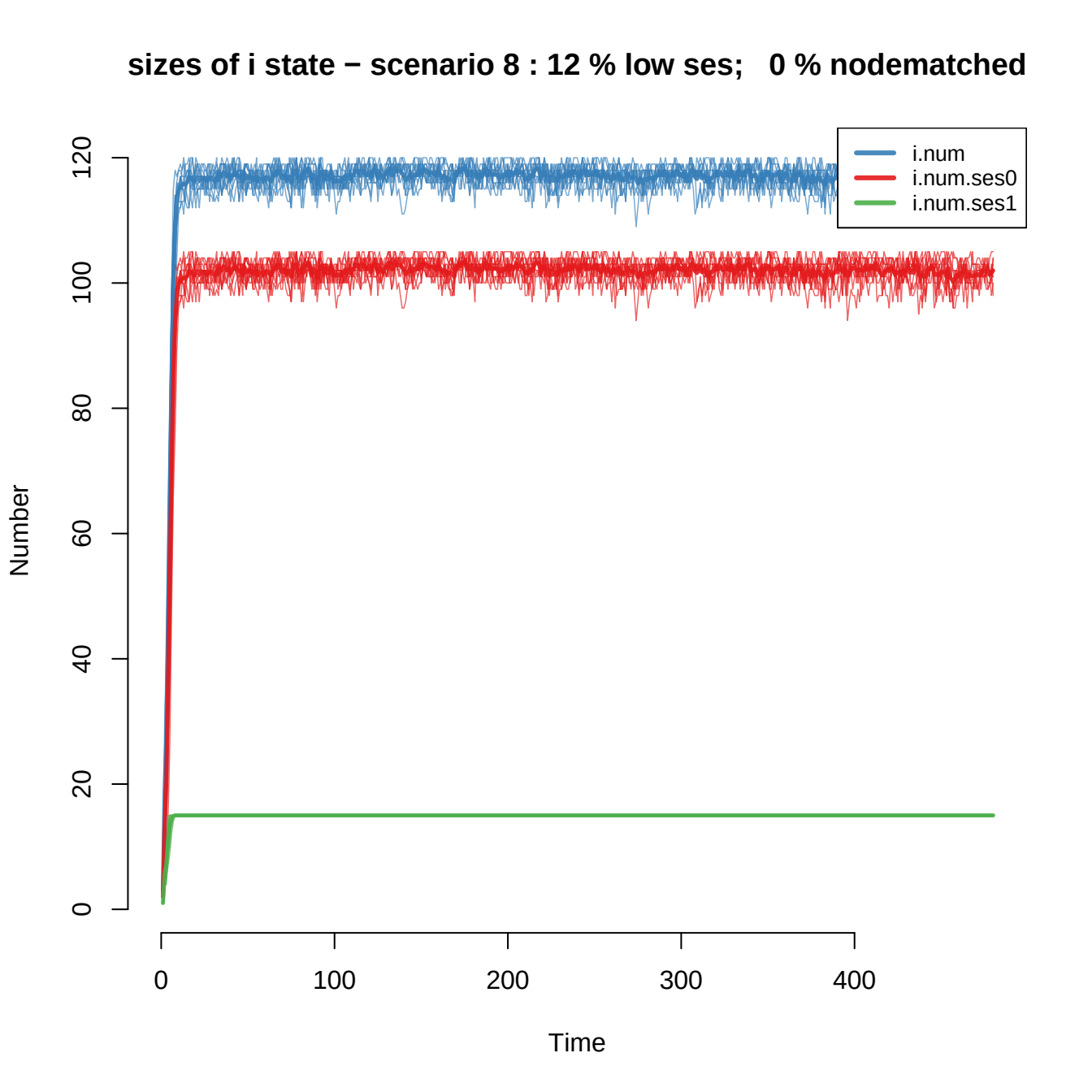

incidence for sub−populations − scenario 6 : 30 % low ses; 90 % nodematched
IR_overall
0.00
0.25
0.50
0.75
0
100
200
300
400
500
t
population
overall
ses0
ses1

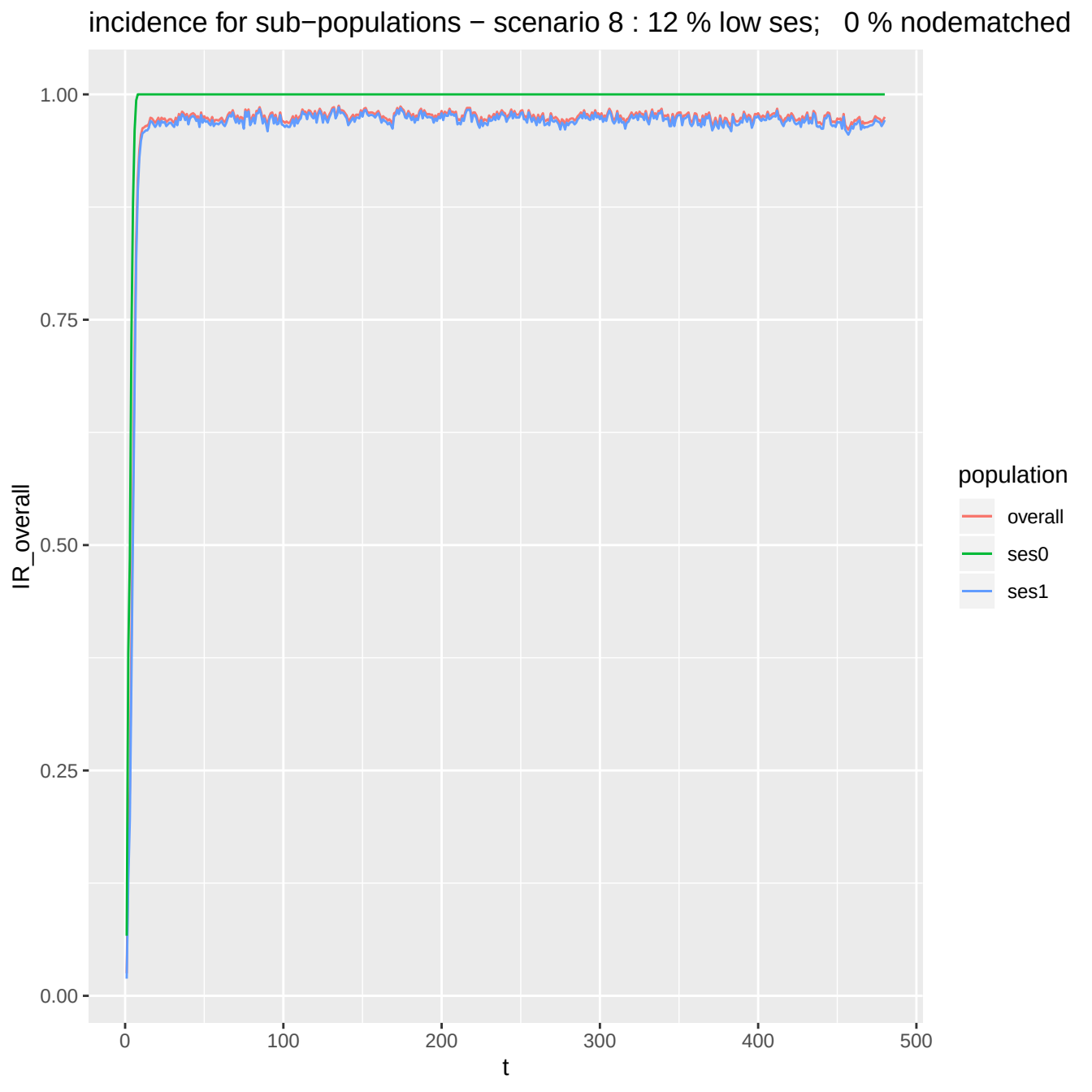

relative risk ses1 to ses0 − scenario 6 : 30 % low ses; 90 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

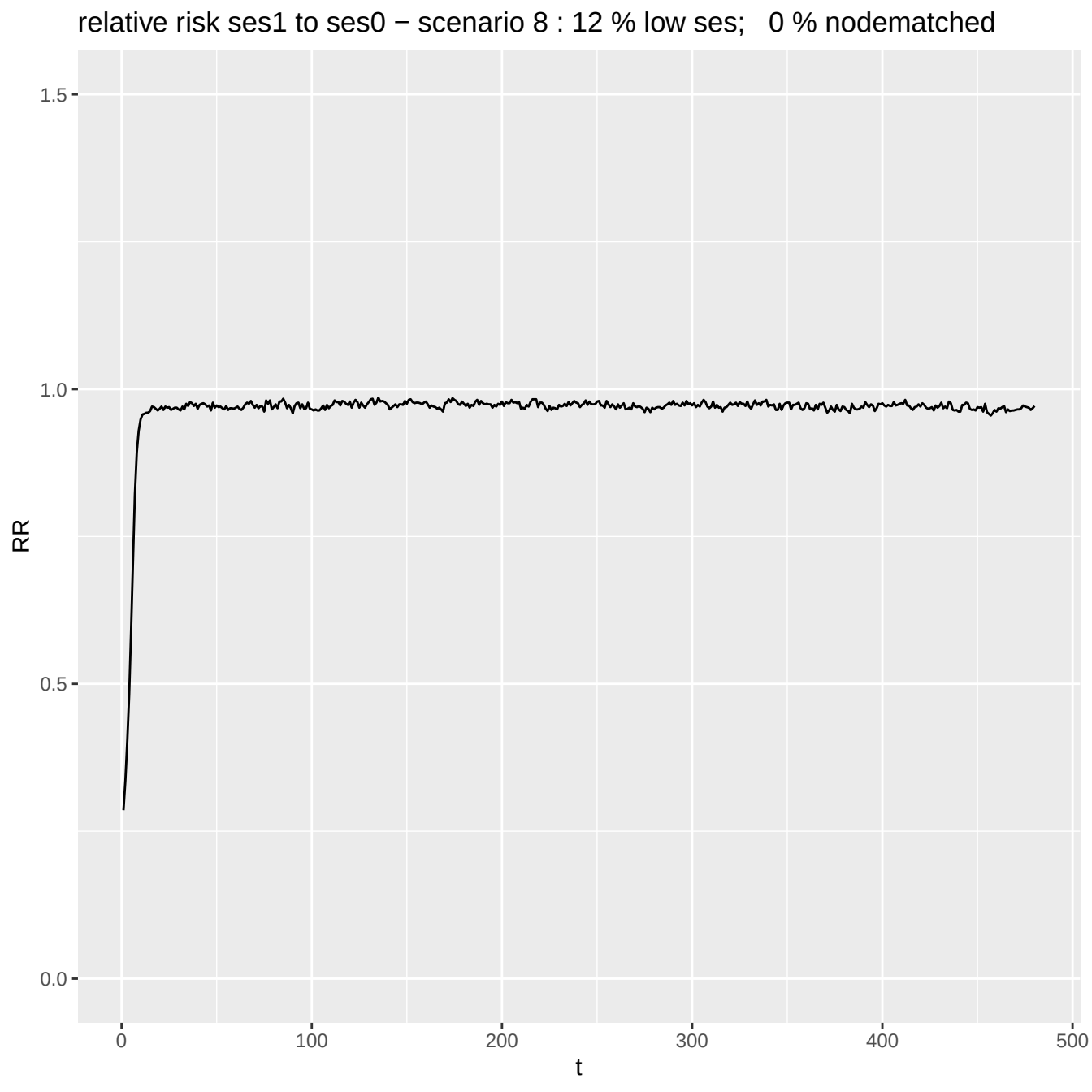

sizes of i state − scenario 7 : 30 % low ses; 100 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
50
100
150
200
250
0
100
200
300
400
Time

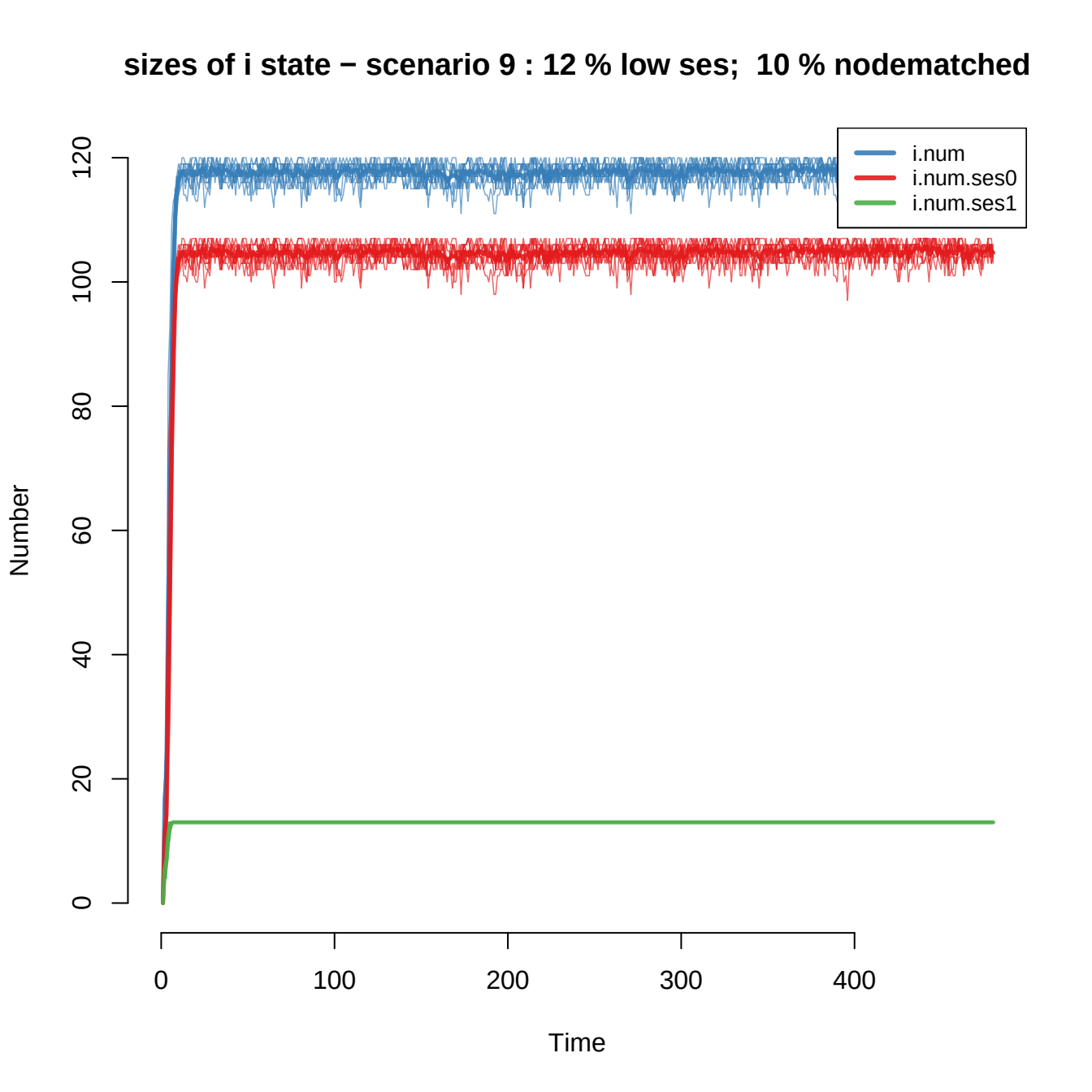

incidence for sub−populations − scenario 7 : 30 % low ses; 100 % nodematched
IR_overall
0.00
0.25
0.50
0.75
0
100
200
300
400
500
t
population
overall
ses0
ses1

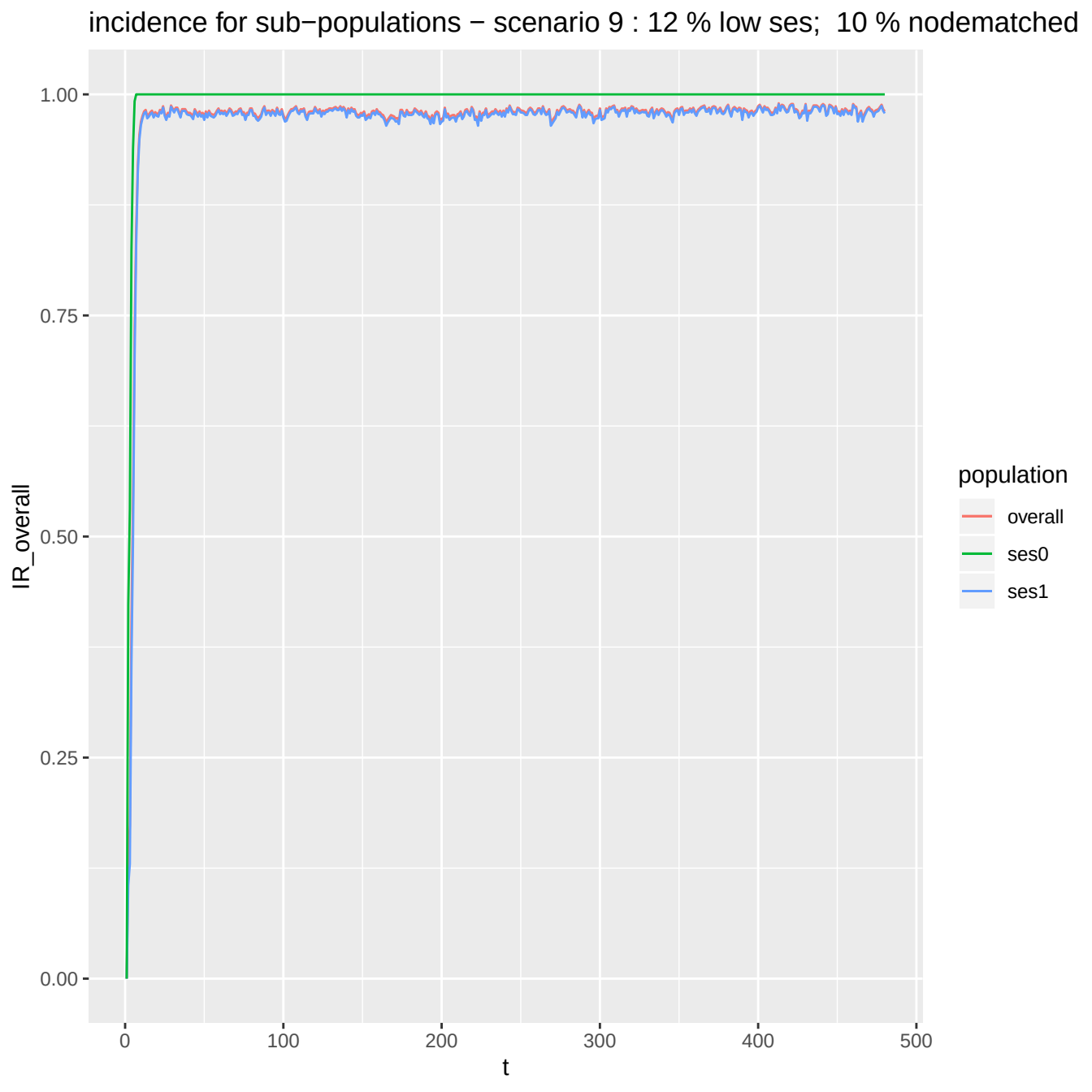

relative risk ses1 to ses0 − scenario 7 : 30 % low ses; 100 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

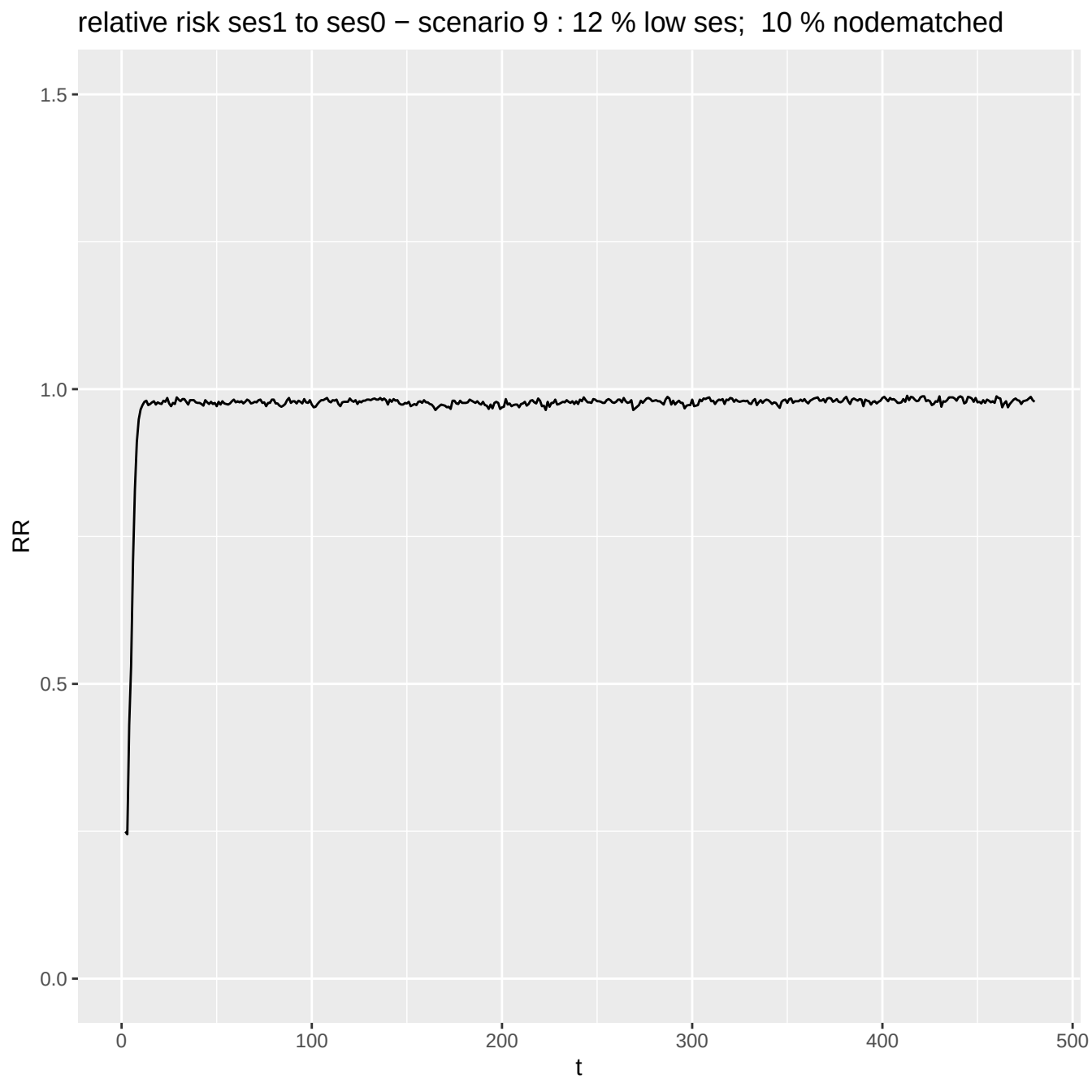

sizes of i state − scenario 8 : 12 % low ses; 0 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
20
40
60
80
100
120
0
100
200
300
400
Time

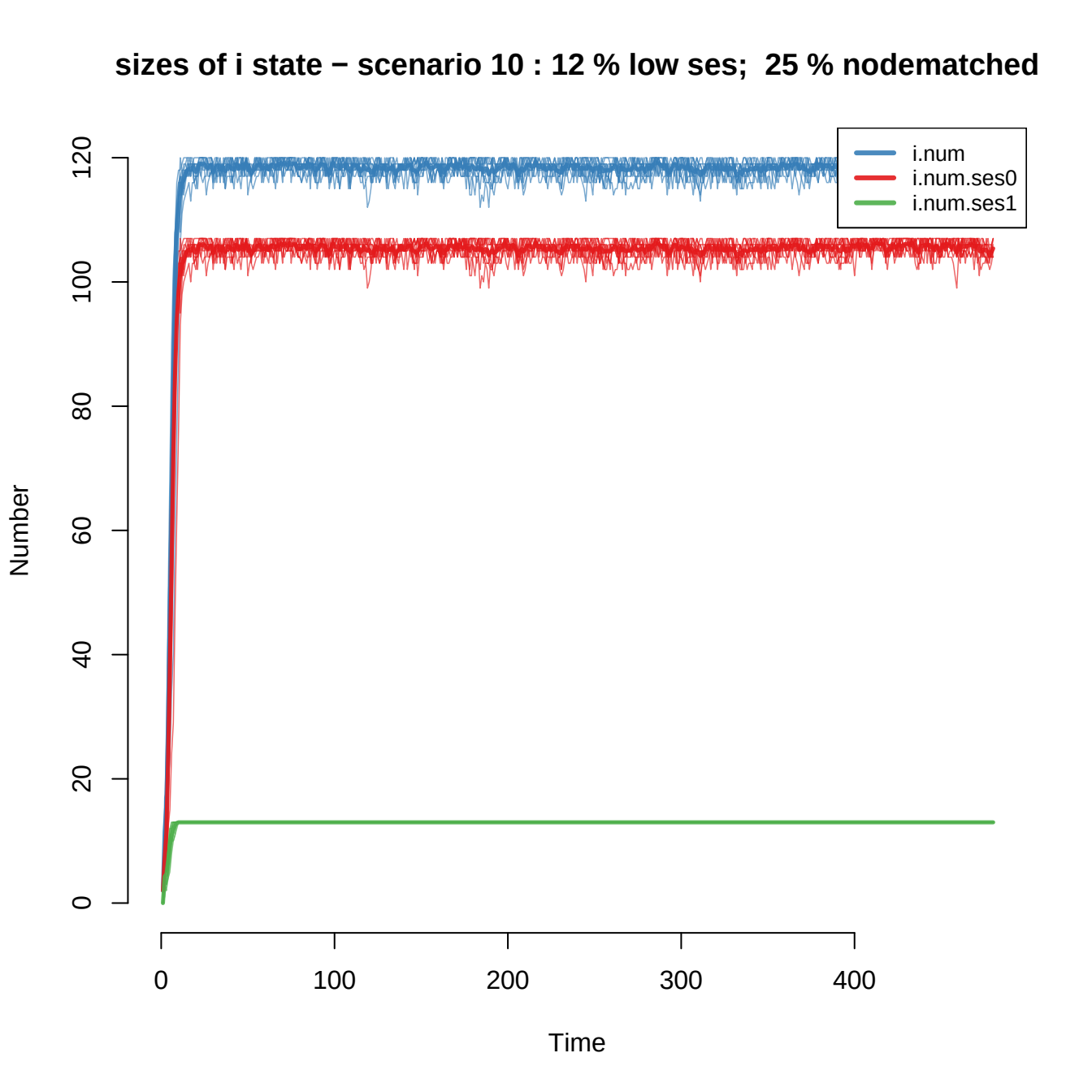

incidence for sub−populations − scenario 8 : 12 % low ses; 0 % nodematched
IR_overall
1.00
0.75
0.50
0.25
0.00
0
100
200
300
400
500
t
population
overall
ses0
ses1

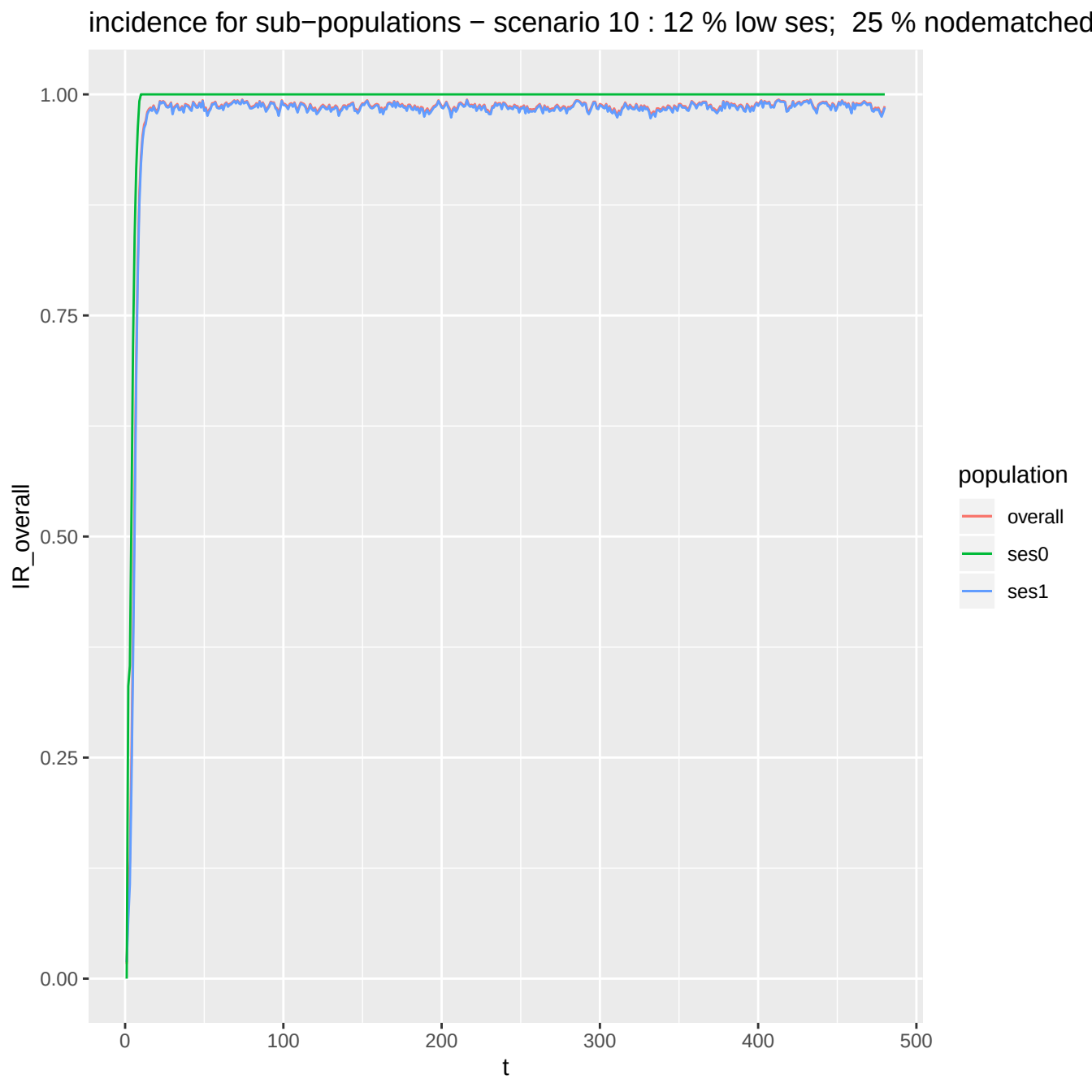

relative risk ses1 to ses0 − scenario 8 : 12 % low ses; 0 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

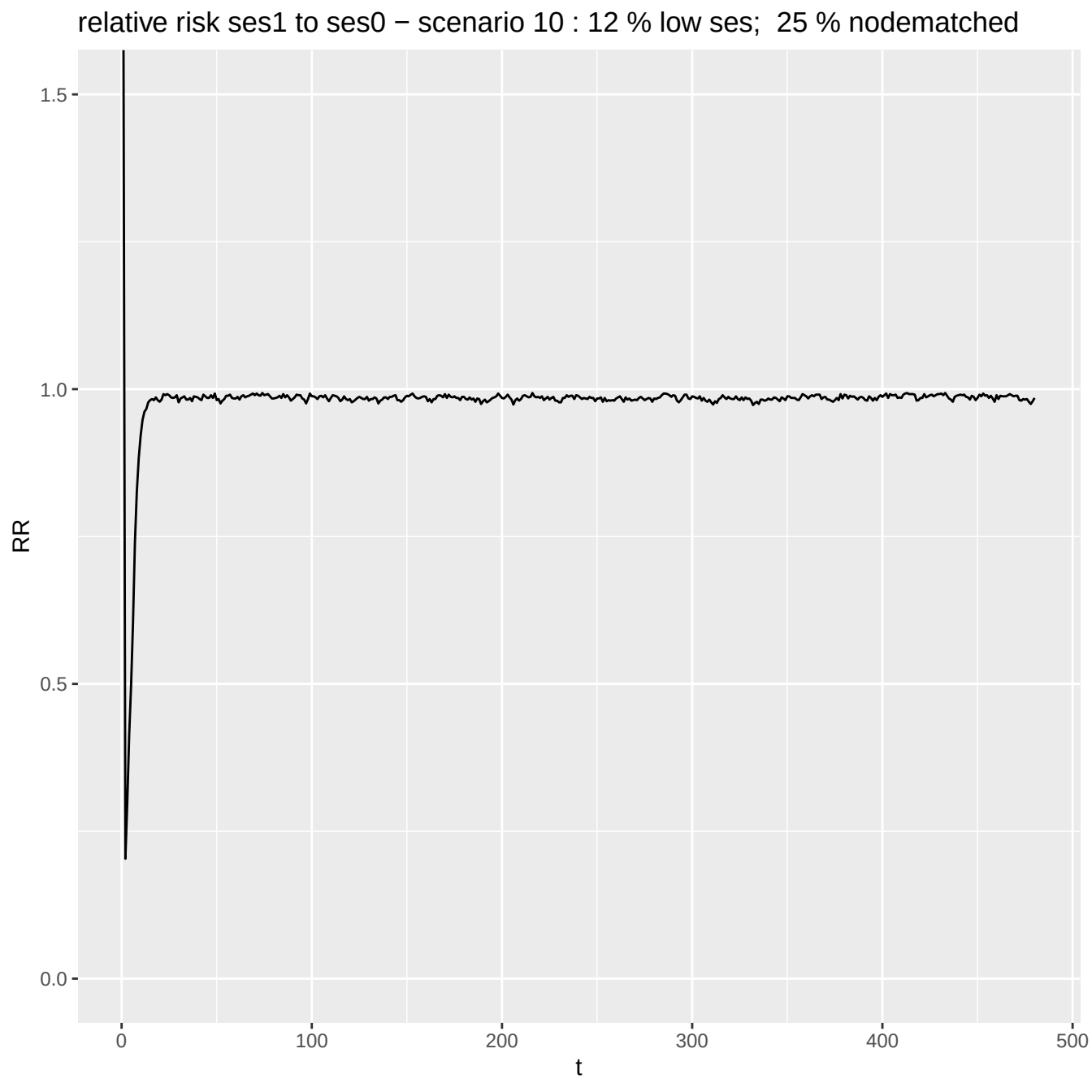

sizes of i state − scenario 9 : 12 % low ses; 10 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
20
40
60
80
100
120
0
100
200
300
400
Time

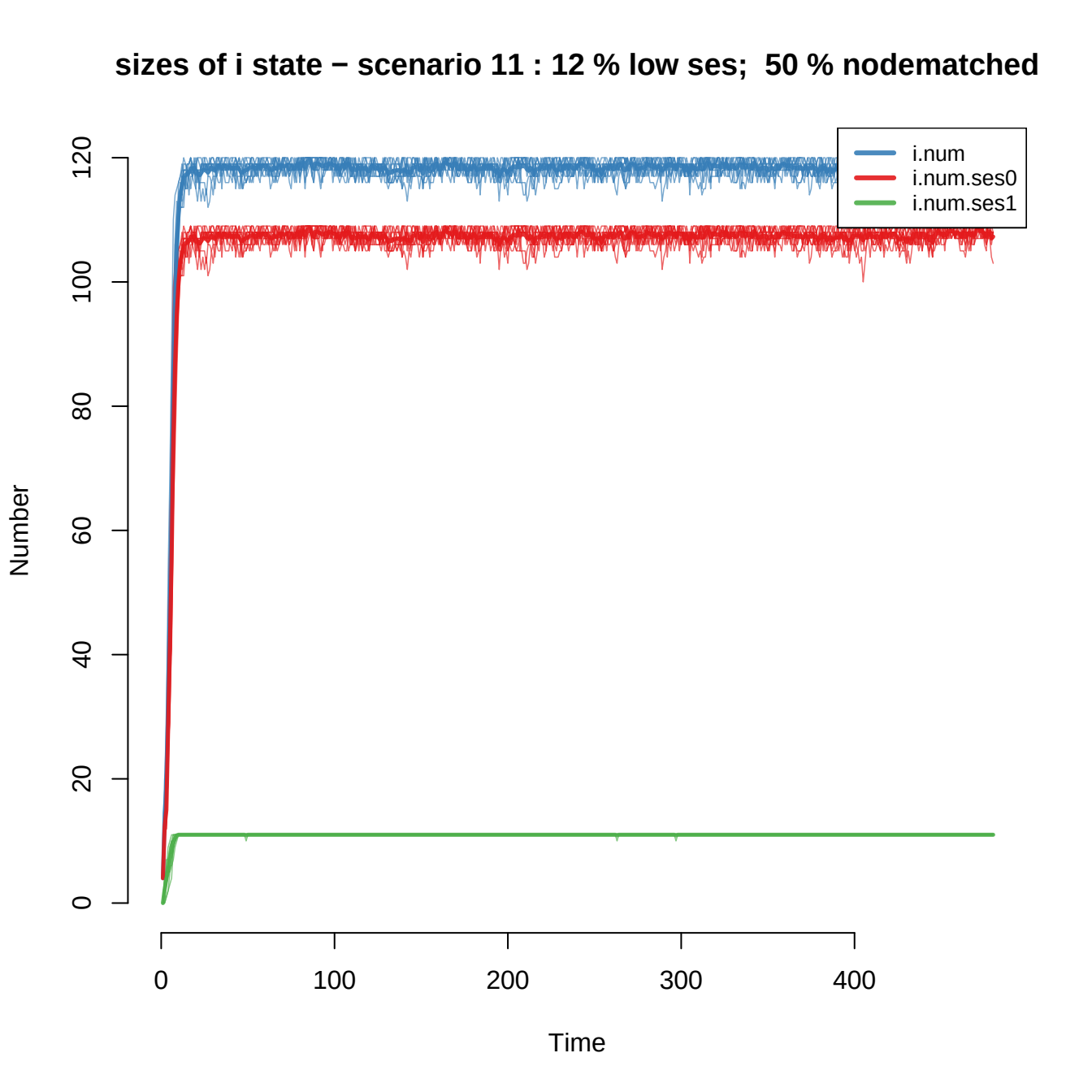

incidence for sub−populations − scenario 9 : 12 % low ses; 10 % nodematched
IR_overall
1.00
0.75
0.50
0.25
0.00
0
100
200
300
400
500
t
population
overall
ses0
ses1

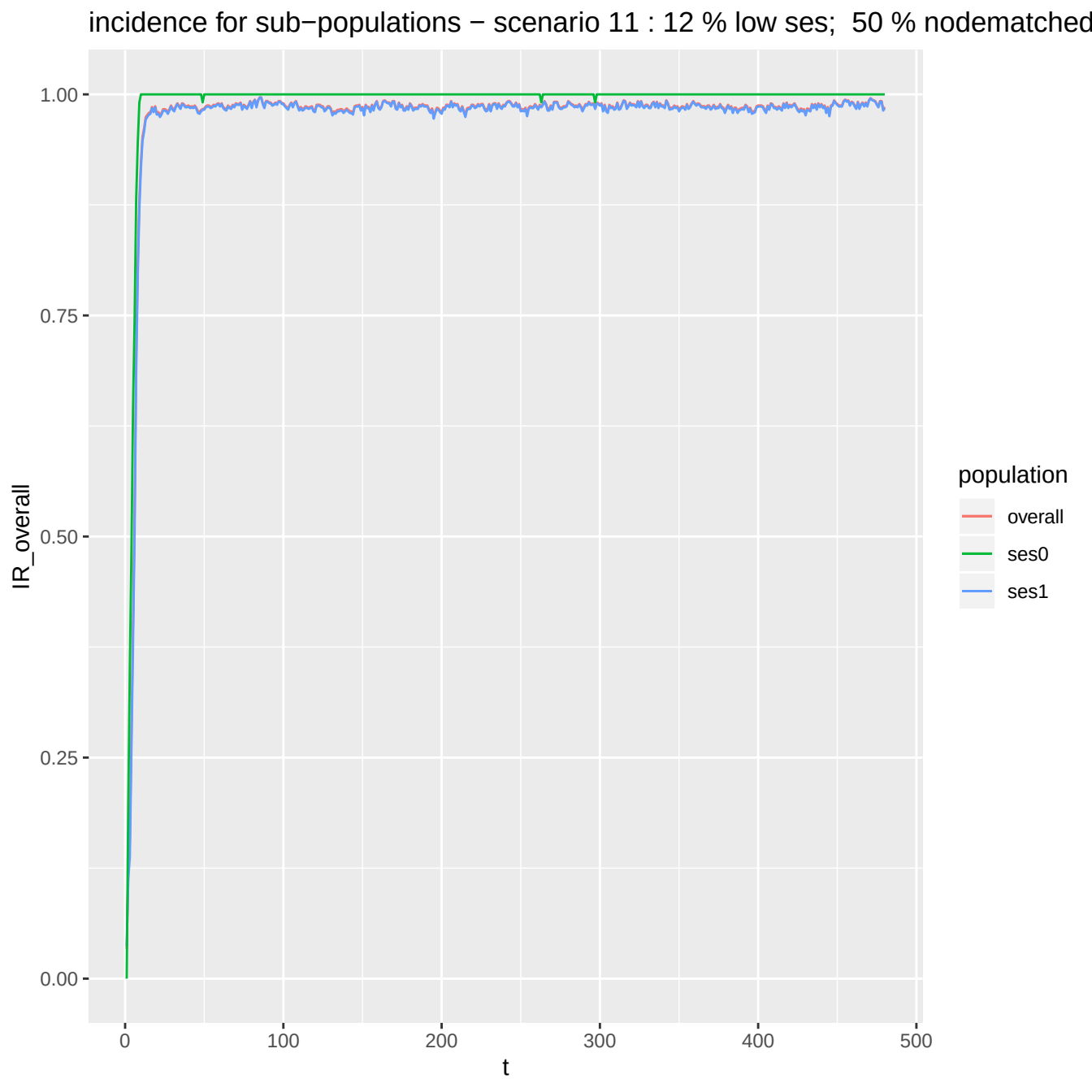

relative risk ses1 to ses0 − scenario 9 : 12 % low ses; 10 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

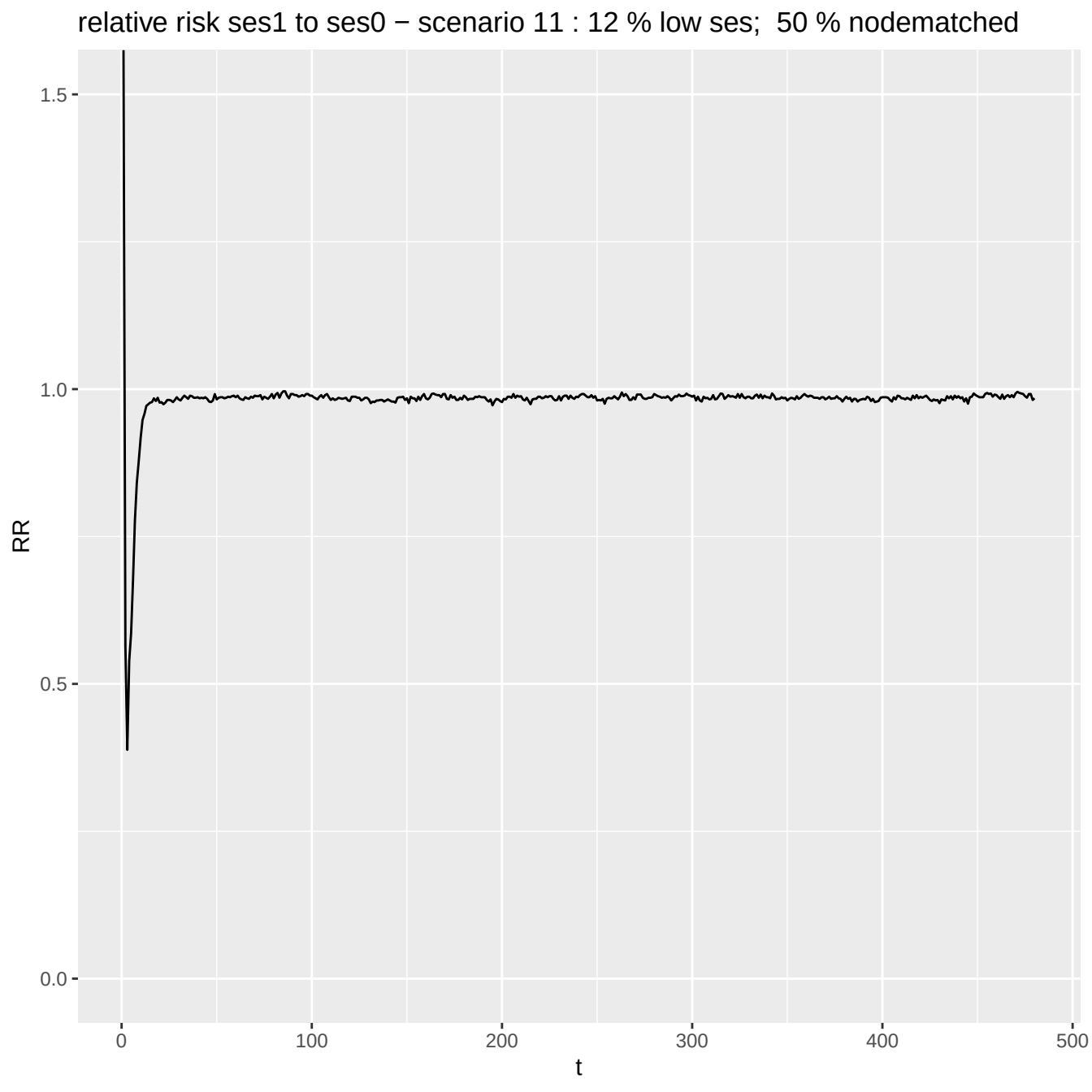

sizes of i state − scenario 10 : 12 % low ses; 25 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
20
40
60
80
100
120
0
100
200
300
400
Time

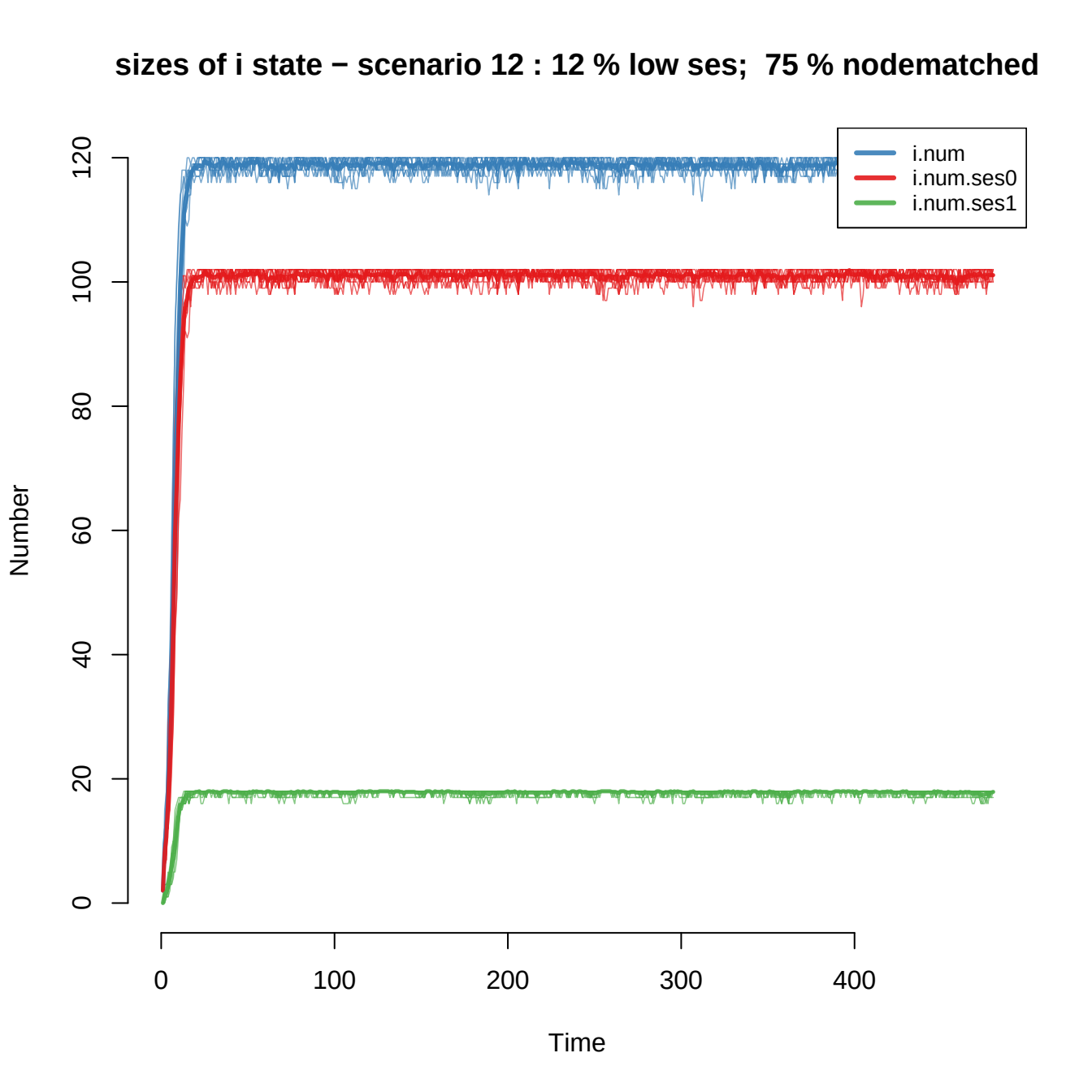

incidence for sub−populations − scenario 10 : 12 % low ses; 25 % nodematched
IR_overall
1.00
0.75
0.50
0.25
0.00
0
100
200
300
400
500
t
population
overall
ses0
ses1

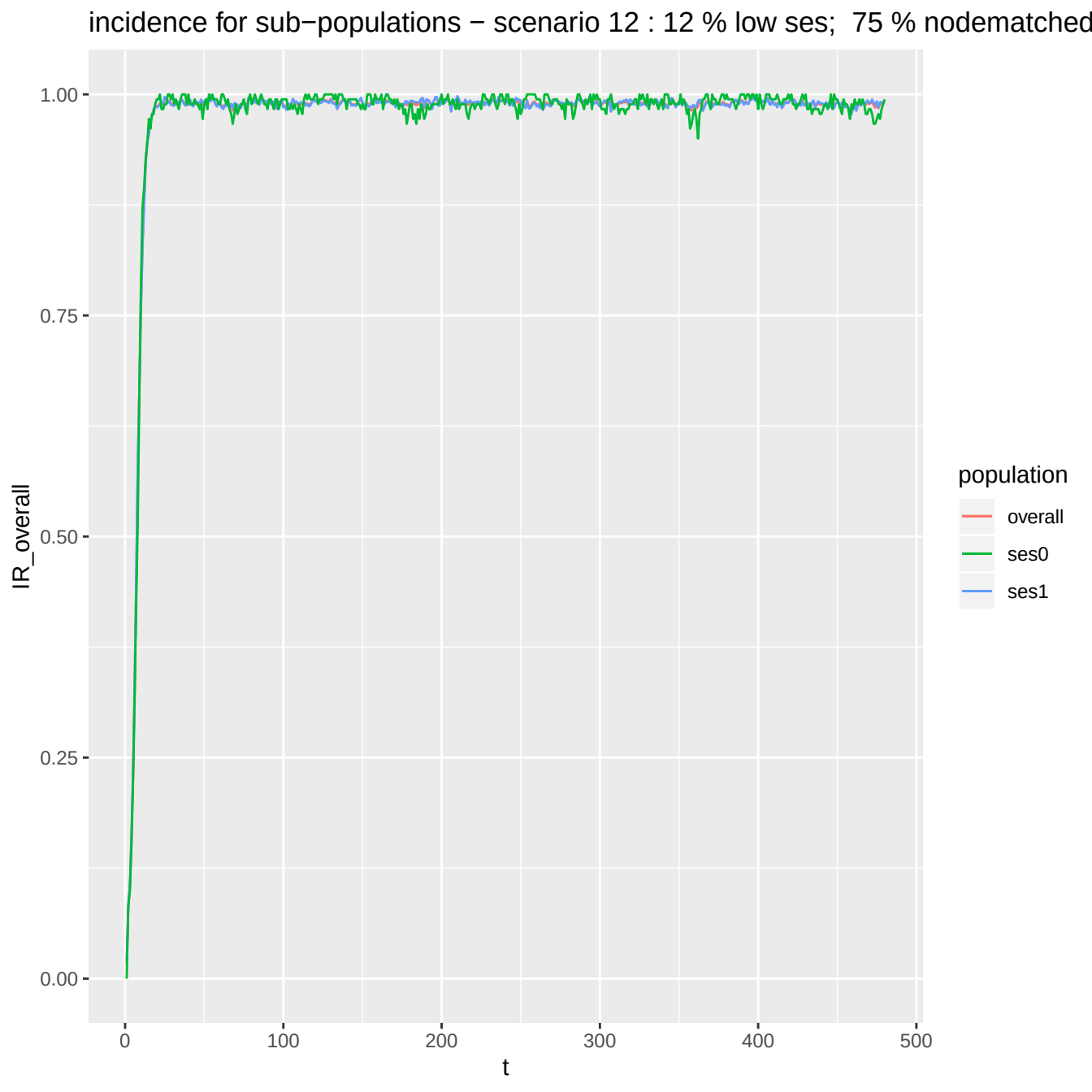

relative risk ses1 to ses0 − scenario 10 : 12 % low ses; 25 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

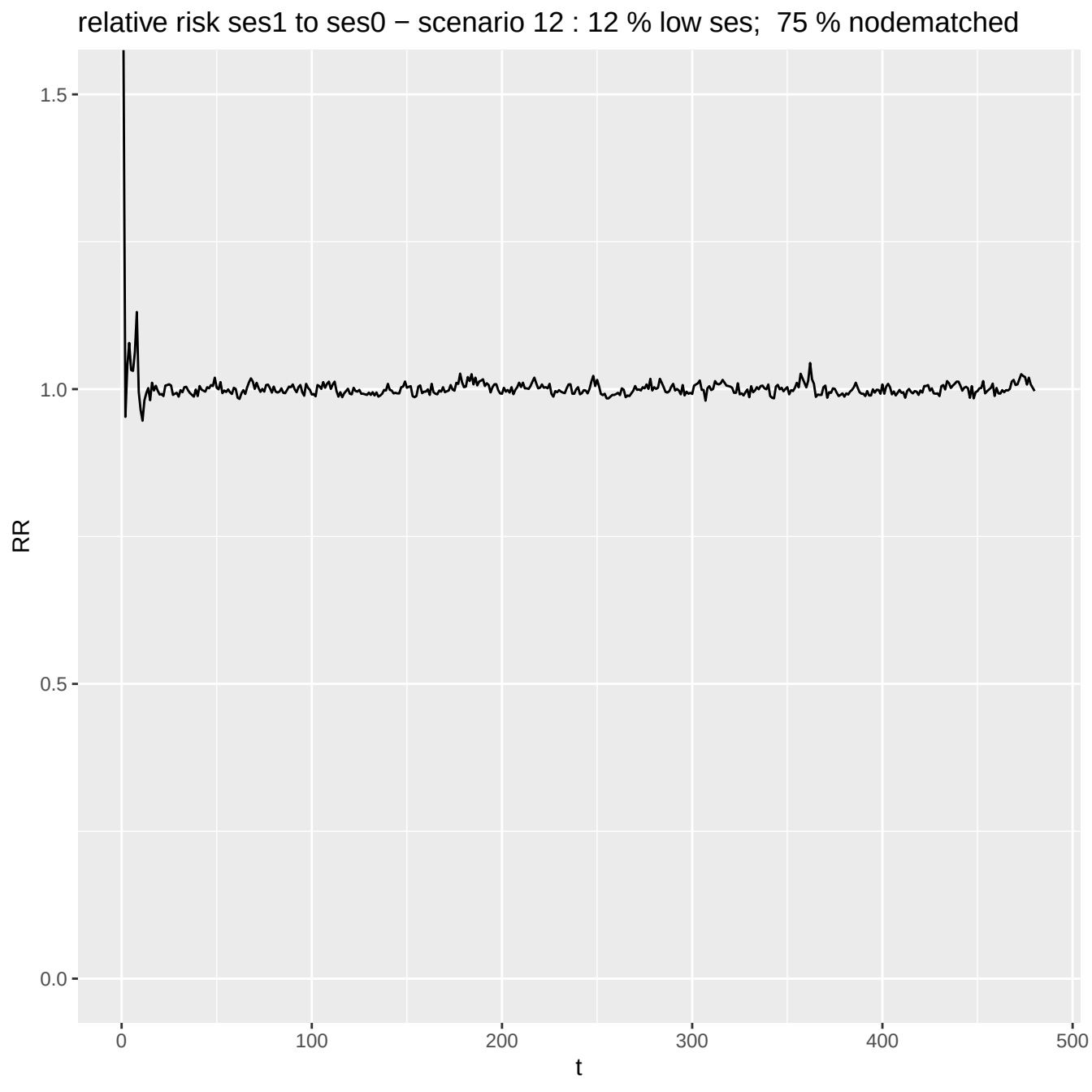

sizes of i state − scenario 11 : 12 % low ses; 50 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
20
40
60
80
100
120
0
100
200
300
400
Time

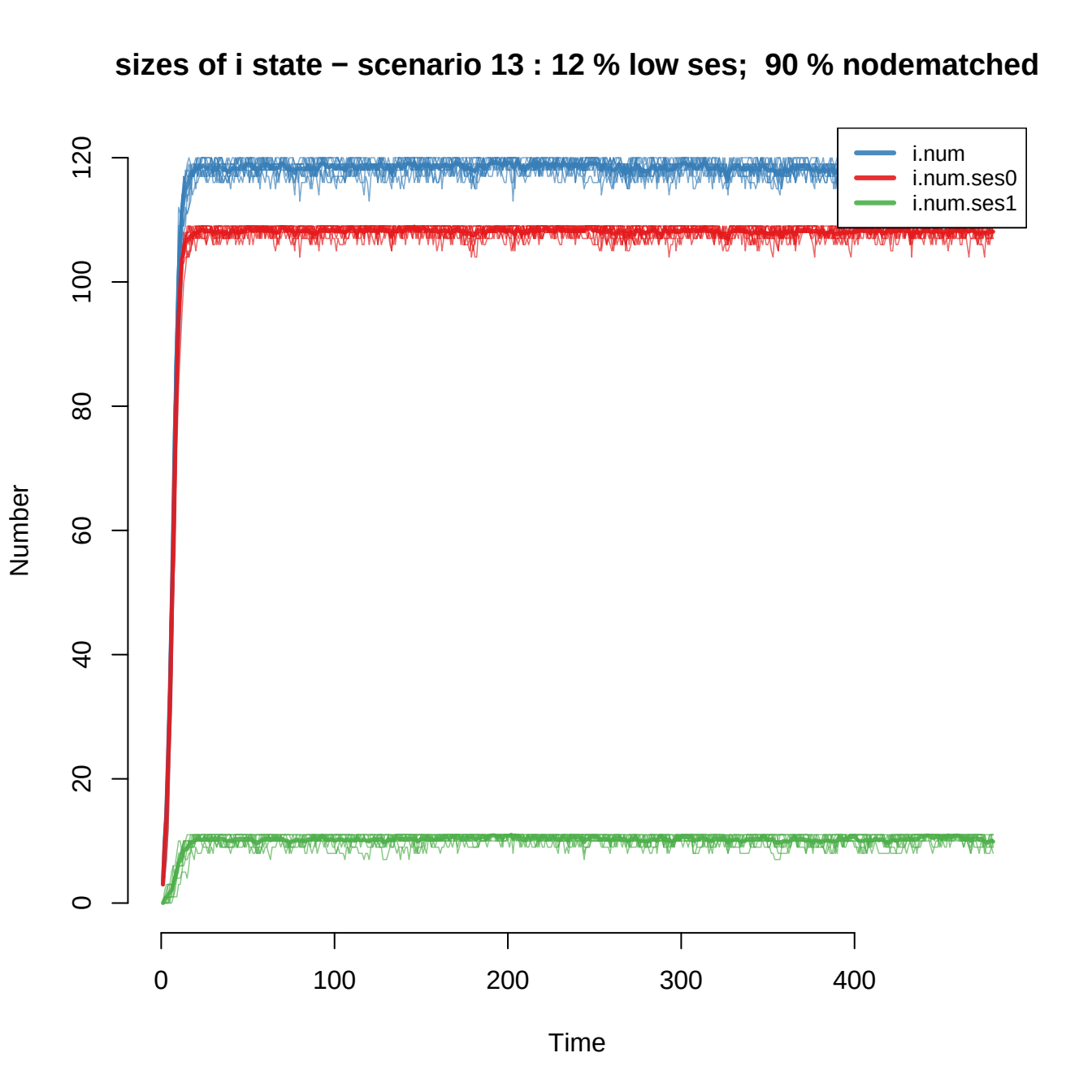

incidence for sub−populations − scenario 11 : 12 % low ses; 50 % nodematched
IR_overall
1.00
0.75
0.50
0.25
0.00
0
100
200
300
400
500
t
population
overall
ses0
ses1

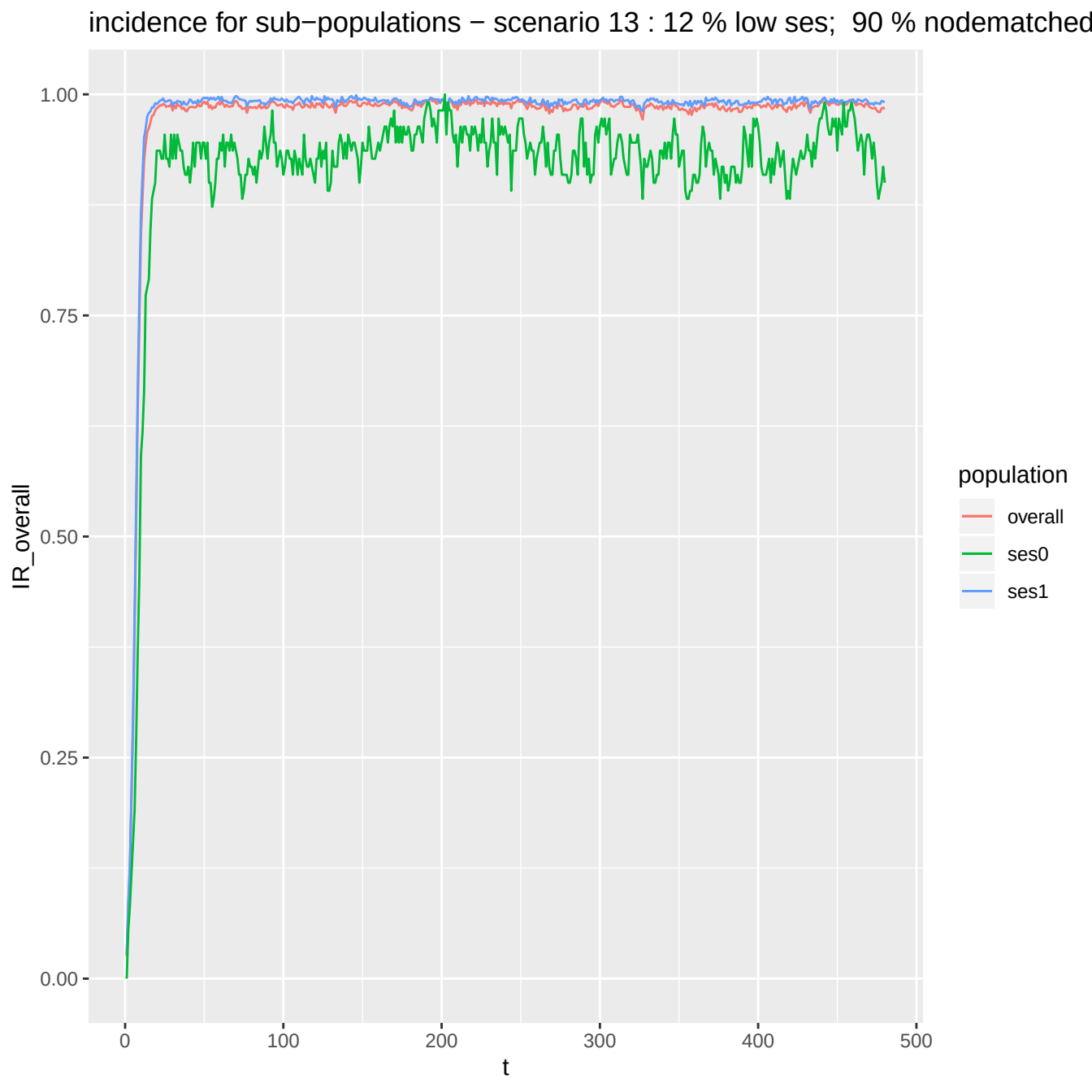

relative risk ses1 to ses0 − scenario 11 : 12 % low ses; 50 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

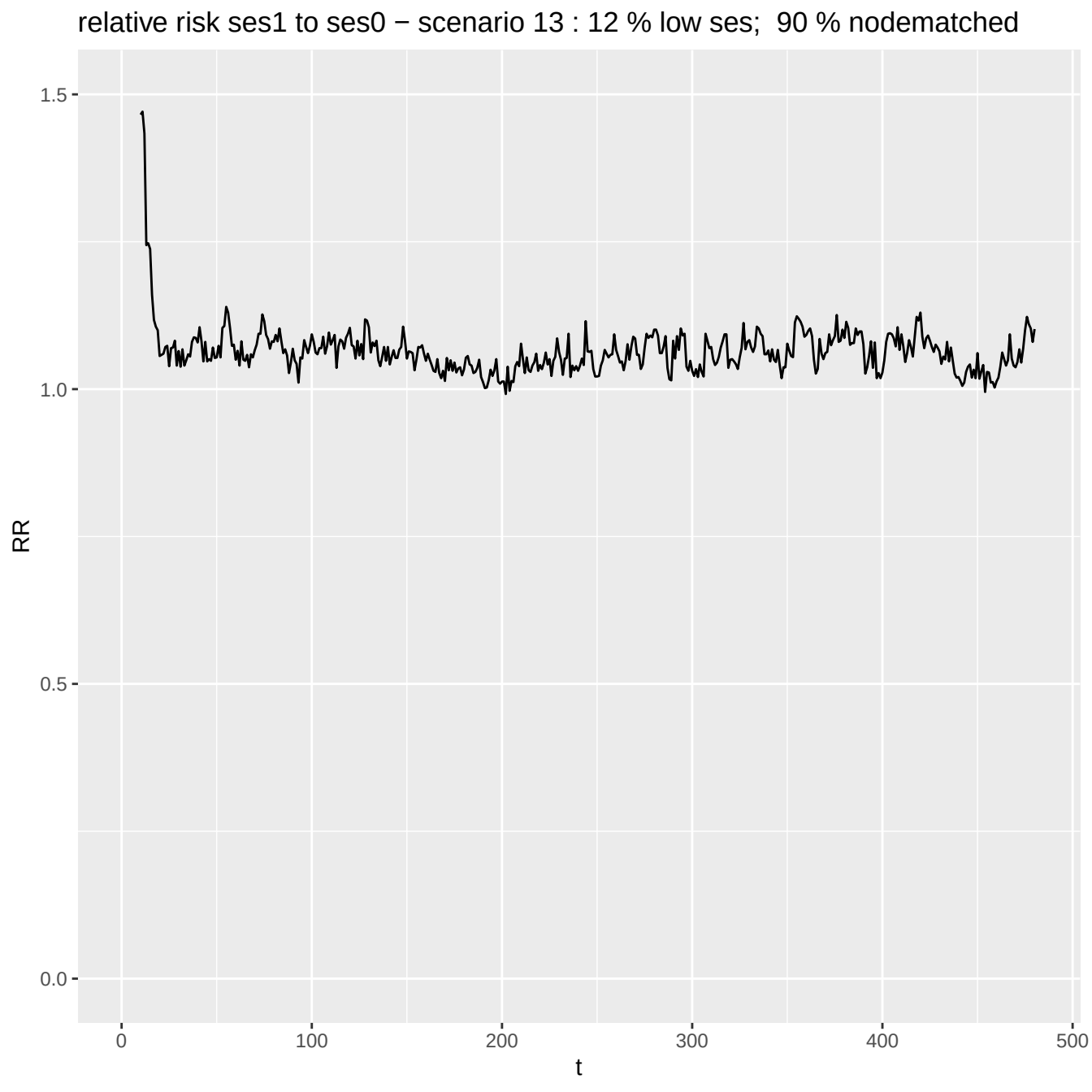

sizes of i state − scenario 12 : 12 % low ses; 75 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
20
40
60
80
100
120
0
100
200
300
400
Time

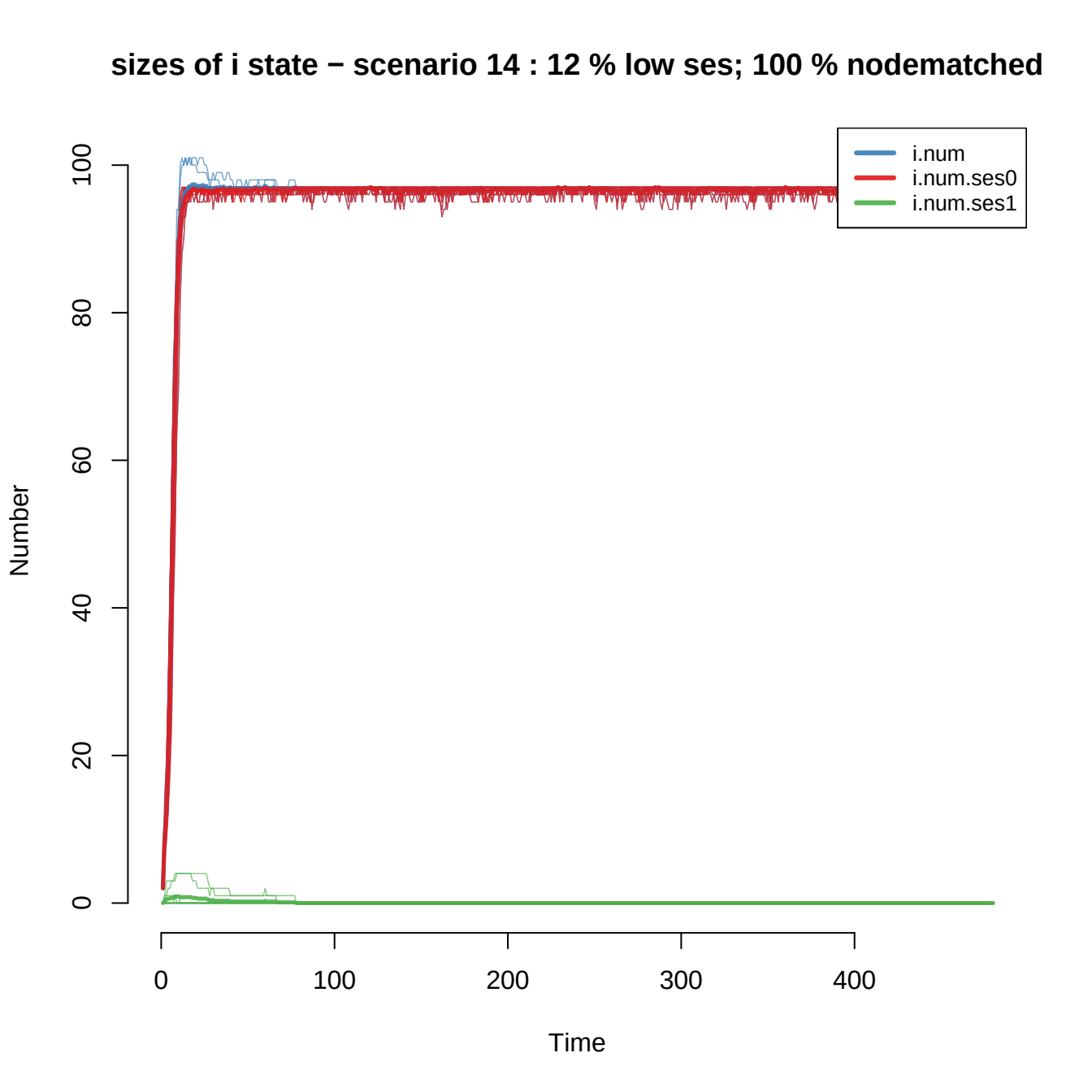

incidence for sub−populations − scenario 12 : 12 % low ses; 75 % nodematched
IR_overall
1.00
0.75
0.50
0.25
0.00
0
100
200
300
400
500
t
population
overall
ses0
ses1

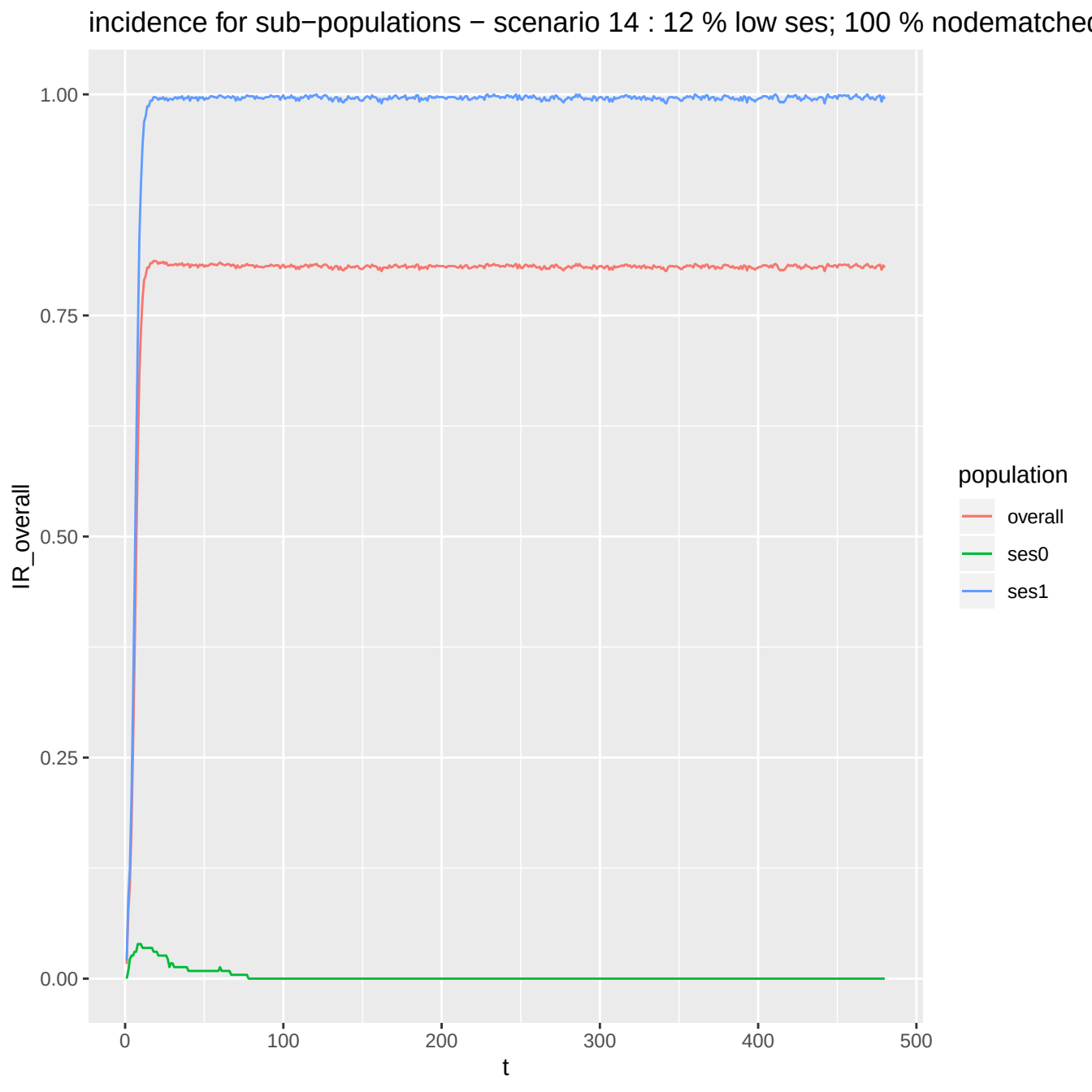

relative risk ses1 to ses0 − scenario 12 : 12 % low ses; 75 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

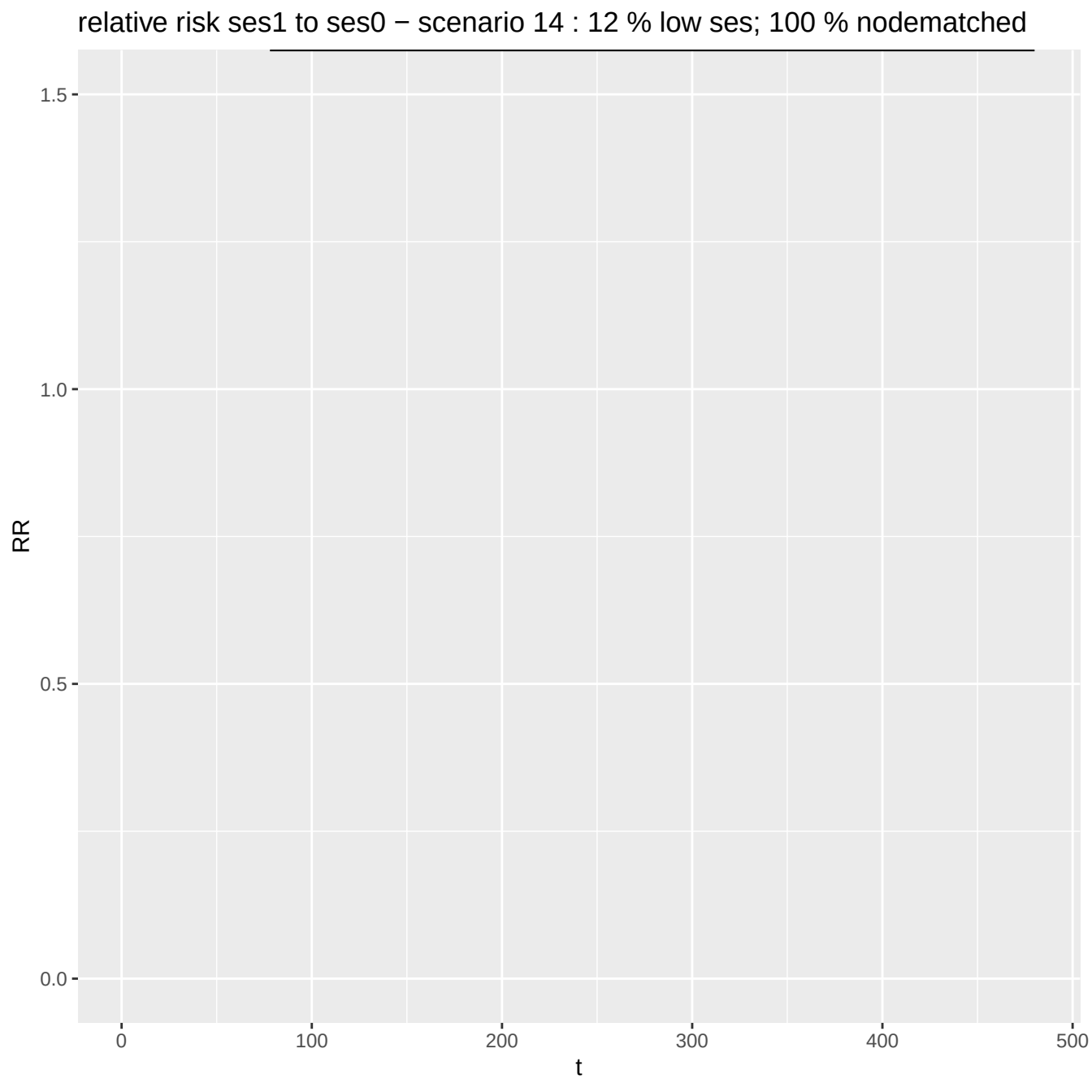

sizes of i state − scenario 13 : 12 % low ses; 90 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
20
40
60
80
100
120
0
100
200
300
400
Time


incidence for sub−populations − scenario 13 : 12 % low ses; 90 % nodematched
IR_overall
1.00
0.75
0.50
0.25
0.00
0
100
200
300
400
500
t
population
overall
ses0
ses1


relative risk ses1 to ses0 − scenario 13 : 12 % low ses; 90 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t


sizes of i state − scenario 14 : 12 % low ses; 100 % nodematched
i.num
i.num.ses0
i.num.ses1
Number
0
20
40
60
80
100
0
100
200
300
400
Time


incidence for sub−populations − scenario 14 : 12 % low ses; 100 % nodematched
IR_overall
1.00
0.75
0.50
0.25
0.00
0
100
200
300
400
500
t
population
overall
ses0
ses1


relative risk ses1 to ses0 − scenario 14 : 12 % low ses; 100 % nodematched
1.5
1.0
0.5
0.0
RR
0
100
200
300
400
500
t

	scenario	overall_mean	overall_sd	ses1_mean	ses1_sd	ses0_mean	ses0_sd
1	1	0.868	0.006	0.827	0.009	0.962	0.006
2	2	0.875	0.008	0.847	0.011	0.94	0.008
3	3	0.874	0.006	0.85	0.008	0.94	0.009
4	4	0.881	0.007	0.874	0.008	0.898	0.009
5	5	0.872	0.006	0.897	0.006	0.812	0.014
6	6	0.848	0.006	0.923	0.006	0.671	0.019
7	7	0.789	0.009	0.93	0.006	0.459	0.022
8	8	0.975	0.005	0.971	0.005	1	0
9	9	0.982	0.004	0.979	0.005	1	0
10	10	0.987	0.004	0.985	0.004	1	0
11	11	0.987	0.003	0.986	0.004	1	0.001
12	12	0.99	0.003	0.99	0.003	0.99	0.008
13	13	0.987	0.004	0.992	0.003	0.937	0.026
14	14	0.805	0.002	0.996	0.002	0	0