

Sample Size for 2016 Chinook Tagging Study: Supplemental Release

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Summary of Recommendations

Assume

- Barrier is installed at head of Old River
- Approximately 25% of fish reaching the Turner Cut Junction enter Turner Cut
- The survival probability from Turner Cut to Chipps Island is 0.05 for fish entering Turner Cut
- A primary release at Durham Ferry is paired with a supplemental release in the San Joaquin River downstream of the Old River flow split
- Total sample size = 648

Recommend

- To estimate route selection probability at the head of Old River: Release scenario II
- To estimate route selection probability at Turner Cut: Release scenario I
- To estimate survival from through the Old River route to Chipps Island: Release scenario II
- To estimate survival from the lower San Joaquin River or Turner Cut to Chipps Island: Release scenario I
- To estimate survival from Mossdale to the Turner Cut junction: Release scenario II

Where

- Release scenario I = release 324 fish at Durham Ferry, 324 fish in supplemental release
- Release scenario II = release 500 fish at Durham Ferry, 148 fish in supplemental release
- “Low” or “Low Mixed” survival was assumed

Introduction

This analysis updates previous sample size analyses for Chinook salmon acoustic tagging studies in the South Delta. It uses data simulation to select between two scenarios using a primary release at Durham Ferry paired with a supplemental release in the San Joaquin River downstream of the Old River flow split. Scenario I uses equal release sizes at the two release locations: 324 fish released both at Durham Ferry and in the supplemental release. Scenario II uses 500 fish released at Durham Ferry and 148 fish released in the supplemental release. Both scenarios use a total release size of 648. Data were simulated under the “Low” and “Low Mixed” survival scenarios from a previous sample size analysis (Buchanan 2014). Primary focus is on estimating survival from various points to Chipps Island

Methods

Analysis methods were based on the methods described in Buchanan (2014). Detection data were simulated 10,000 times from a simplified survival model using 48 parameter sets and two candidate release scenarios, as described above. For each simulated data set, parameter estimates were computed using Method of Moments. For each parameter set and release size combination, the mean, maximum, and standard deviation of the sampling distribution of parameter estimates was computed, as well as the number of simulations in which each parameter was estimable. The preferred release scenario of the two scenarios considered was identified based on several criteria on the estimability of parameters and validity of estimates.

Parameters estimated in the survival model were (Figure 1):

- Overall survival from Durham Ferry to Chipps Island (s_R)
- Survival from Durham Ferry to the head of Old River (s_{R0})
- Survival from the head of Old River to Chipps Island in both the San Joaquin River route and the Old River route (s_A, s_B)
- Survival from the head of Old River to Turner Cut in the San Joaquin River (s_{A1})
- Survival from the Turner Cut junction to Chipps Island in both the San Joaquin River route and the Turner Cut route (s_{A2}, s_F)
- Route selection at the head of Old River and at Turner Cut (ψ_{A1}, ψ_{A2})
- Detection probabilities at the dual arrays at the 5 detection sites

The supplemental release located downstream of the Old River flow split contributed to estimation of parameters s_{A2} , s_F , ψ_{A2} , and detection probabilities at 3 detection sites, when combined with data from the primary release at Durham Ferry. The survival model is described in more detail in Buchanan (2014).

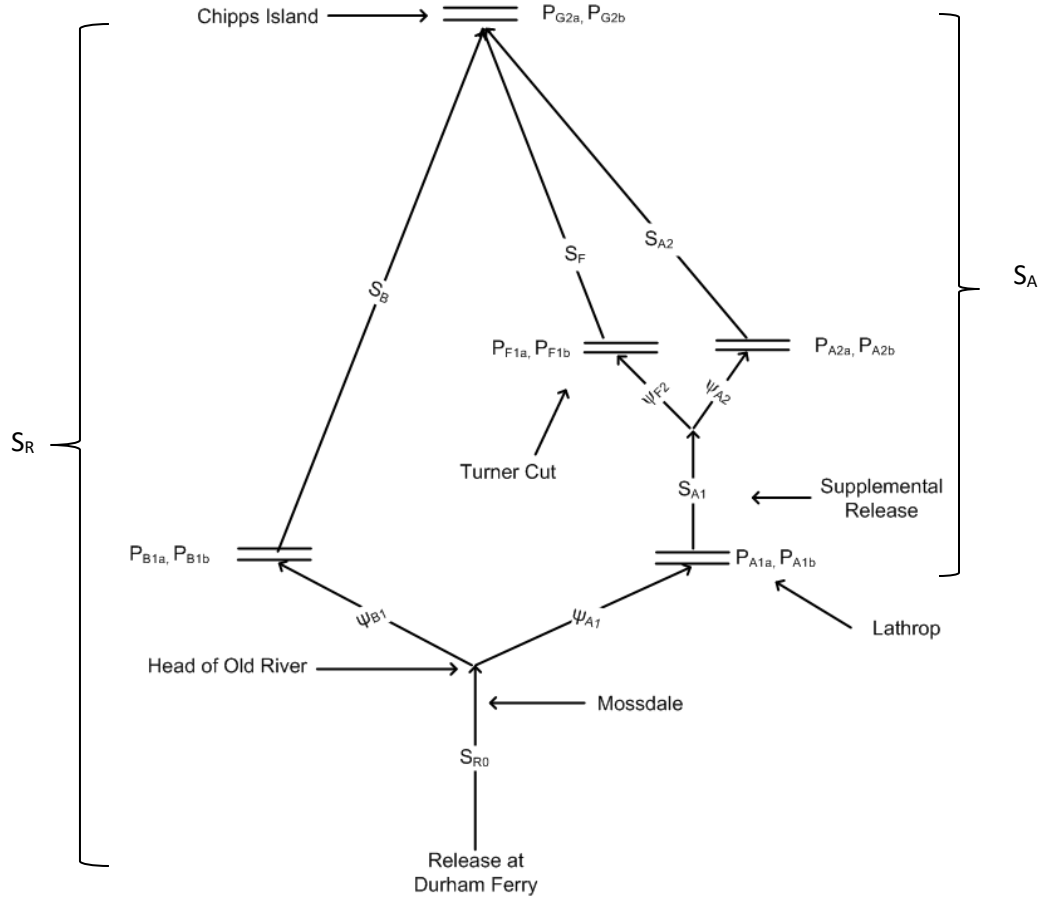


Figure 1. Schematic of model used in data simulations, with parameters: survival from Durham Ferry to Chipps Island (S_R), survival from head of Old River/Mosssdale in the San Joaquin River route (S_A) and Old River route (S_B), probability of survival from head of Old River/Mosssdale to Turner Cut (S_{A1}), survival from Turner Cut to Chipps Island in the San Joaquin River route (S_{A2}) and Turner Cut route (S_F), and the probabilities of remaining in the San Joaquin River at the head of Old River (ψ_{A1}) and at Turner Cut (ψ_{A2}). Other parameters are survival from Durham Ferry to the head of Old River/Mosssdale (S_{R0}), and detection probabilities on the upstream and downstream arrays at Lathrop (P_{A1a} , P_{A1b}), Old River (P_{B1a} , P_{B1b}), Turner Cut in the San Joaquin (P_{A2a} , P_{A2b}) and in Turner Cut (P_{F1a} , P_{F1b}), and at Chipps Island (P_{G2a} , P_{G2b}).

The total release size was assumed to be 648. Two candidate release scenarios were considered (Table 1), to determine the preferred distribution of the total release size over the two release locations (primary release at Durham Ferry, supplemental release downstream).

Table 1. Release scenarios considered, for a total release size of 648.

Scenario	Durham Ferry Release Size (R1)	Supplemental Release Size (R2)
I	324	324
II	500	148

Parameters Sets

Parameter values were selected from a subset of the parameters considered in a previous analysis (Buchanan 2014, Tables 2 and 3). The “Low” and “Low Mixed” parameter scenarios were considered for this analysis (scenarios 2 and 3 from Buchanan 2014), based on survival estimates from recent years combined with the possible expectation of higher water than in recent years. These two parameter scenarios both assume that survival from the Durham Ferry release site to Mossdale is $s_{R0}=0.4$, survival from Mossdale to the Turner Cut junction (sites A2 and F1) is $s_{A1}=0.3$, and 98% of the fish reaching the Old River junction remain in the San Joaquin, as expected if a barrier is installed in Old River (Table 2). Survival from site A2 to Chipps Island (site G2) was modeled as either 0.07 or 0.15, survival from Turner Cut to Chipps Island was assumed to be 0.05, and survival from the head of Old River to Chipps Island via the Old River route was assumed to be either 0.07 or 0.2 (Table 2). The probability of entering Turner Cut upon reaching the Turner Cut junction was assumed to be 0.25. Detection probabilities for each line of the dual arrays were assumed to be either 0.9 or 0.85 at the upriver sites (A1, B1), 0.85 at sites A2 and F1, and either 0.5 or 0.9 at site G2 (Chipps Island). Results for alternative values of the route selection parameters (ψ_{A1} and ψ_{A2}) and S_F are provided in Appendix A.

Table 2. Parameter sets used in data simulations to estimate survival to Turner Cut and Chipps Island and route selection at Old River and Turner Cut. Values of S_R and S_A were computed from values of other parameters. For each scenario, $s_F=0.05$, $\psi_{A2}=0.75$, $p_{G2a}=p_{G2b}=0.5$ or 0.9 (equivalent to $p_{G2}=0.75$ or 0.99), and $p_{A2a} = p_{A2b} = p_{F1a} = p_{F1b} = 0.85$.

Scenario	S_{R0}	S_{A1}	S_{A2}	S_B	ψ_{A1}	S_R	S_A	$p_{A1a}, p_{A1b}, p_{B1a}, p_{B1b}$
2: Low	0.4	0.3	0.07	0.07	0.98	0.008	0.020	0.9
3: Low Mixed	0.4	0.3	0.15	0.2	0.98	0.016	0.038	0.85

Criteria

Four criteria were used to assess the candidate release scenarios (Table 3). These criteria are the same as those used in previous analyses (e.g., Buchanan 2014), except that the maximum standard error considered for criterion C3 was reduced from 0.10 to 0.05.

Table 3. Criteria used for identification of minimum sample size necessary to estimate a model parameter, assuming 10,000 simulations. The standard error in Criterion C3 is calculated as the standard deviation of the observed parameter estimates over all simulations.

Criterion	Definition
C1	Parameter is estimable in at least 95% of simulations (9500 or more)
C2	Probability estimate is not greater than 1.1 in 95% of simulations (9500 or more)
C3	Standard error on parameter estimate is not greater than 0.05
C4	Difference between average of parameter estimates and true parameter value is not greater than 0.05

Results

Results for the four criteria from simulations using the “Low” and “Low Mixed” survival scenarios, and assuming that the probability of remaining in the San Joaquin River at the head of Old River is 0.98, are shown in Table 4. Simulation results indicate that many parameters can be estimated with confidence using either release scenario I (equal split of 324 at both release locations) or release scenario II (500 released at Durham Ferry, 148 released downstream). Release scenario I favors estimation of downstream parameters such as S_{A2} , S_F , and ψ_{A2} , whereas release scenario II favors estimation of upstream parameters such as S_{R0} , as well as S_B . Giving priority to estimation of downstream parameters such as S_{A2} , S_F , and ψ_{A2} , release scenario I is recommended. This recommendation is based on the assumption that a physical barrier will block the majority of access to Old River.

Table 5 presents results for the same survival scenarios but assuming that the probability of remaining in the San Joaquin River at the head of Old River is 0.4 (e.g., if the barrier is either absent or ineffective). Under this assumption, all parameters are expected to be estimable (Criterion C1) using either release scenario (I or II). The only criterion that is sometimes unmet is C3, which requires a standard error < 0.5. Release scenario I is recommended to meet this criterion for downstream San Joaquin River and Turner Cut parameters (S_{A2} , S_F , ψ_{A2}), whereas release scenario II is recommended to meet this criterion for upstream San Joaquin River and Old River parameters (S_{A1} , S_B , ψ_{A1}).

Simulation results for all parameter sets are presented in Appendix A.

Summary

Using a primary release at Durham Ferry and a supplemental release in the San Joaquin River downstream of the Old River flow split, and a total release size of 648, the following recommendations are made:

- To estimate route selection probability at the head of Old River: Release scenario II
- To estimate route selection probability at Turner Cut: Release scenario I
- To estimate survival from through the Old River route to Chipps Island: Release scenario II
- To estimate survival from the lower San Joaquin River or Turner Cut to Chipps Island: Release scenario I
- To estimate survival from Mossdale to the Turner Cut junction: Release scenario II

Where

- Release scenario I = release 324 fish at Durham Ferry, 324 fish in supplemental release
- Release scenario II = release 500 fish at Durham Ferry, 148 fish in supplemental release
- “Low” or “Low Mixed” survival was assumed

Table 4. Release scenario selected for estimation of each parameter based on the four criteria defined in Table 3, assuming that a 98% probability of remaining in the San Joaquin River at the head of Old River. Release scenarios are: I = (R1 = R2 = 324), and II = (R1 = 500, R2 = 148), where R1 = release size at Durham Ferry, and R2 = size of supplemental release. Release scenarios identified in [brackets] indicate the preferred release scenario in the case when neither release scenario satisfied the criterion. Parameter set numbers refer to Table 6.

Survival Scenario	Parameter	Value	Criterion C1	Criterion C2	Criterion C3	Criterion C4
Low survival, low detection probability at Chipps Island – Parameter Set 6	S _{R0}	0.4	II	I or II	I or II	I or II
	S _{A1}	0.3	I or II	I or II	I or II	I or II
	S _{A2}	0.07	[I]	I or II	I or II	I or II
	S _B	0.07	[I]	I or II	[II]	I or II
	S _F	0.05	[I]	I or II	[I]	I or II
	Ψ _{A1}	0.98	II	I or II	[II]	I or II
	Ψ _{A2}	0.75	I or II	I or II	I or II	I or II
	P _{G1a}	0.5	I or II	I or II	[I]	I or II
	P _{G2b}	0.5	I or II	I or II	[I]	I or II
	S _A	0.020	[I]	I or II	I or II	I or II
	S _R	0.008	[II]	I or II	I or II	I or II
Low survival, high detection probability at Chipps Island – Parameter Set 18	S _{R0}	0.4	II	I or II	I or II	I or II
	S _{A1}	0.3	I or II	I or II	I or II	I or II
	S _{A2}	0.07	I or II	I or II	I or II	I or II
	S _B	0.07	II	I or II	[II]	I or II
	S _F	0.05	I or II	I or II	I or II	I or II
	Ψ _{A1}	0.98	II	I or II	[II]	I or II
	Ψ _{A2}	0.75	I or II	I or II	I or II	I or II
	P _{G1a}	0.9	I or II	I or II	[I]	I or II
	P _{G2b}	0.9	I or II	I or II	[I]	I or II
	S _A	0.020	I or II	I or II	I or II	I or II
	S _R	0.008	II	I or II	I or II	I or II
Low Mixed survival, low detection probability at Chipps Island – Parameter Set 30	S _{R0}	0.4	II	I or II	I or II	I or II
	S _{A1}	0.3	I or II	I or II	I or II	I or II
	S _{A2}	0.15	I or II	I or II	[I]	I or II
	S _B	0.2	II	I or II	[II]	I or II
	S _F	0.05	I or II	I or II	[I]	I or II
	Ψ _{A1}	0.98	II	I or II	[II]	II
	Ψ _{A2}	0.75	I or II	I or II	I or II	I or II
	P _{G1a}	0.5	I or II	I or II	[I]	I or II
	P _{G2b}	0.5	I or II	I or II	[I]	I or II
	S _A	0.038	I or II	I or II	I or II	I or II
	S _R	0.016	[II]	I or II	I or II	I or II
Low Mixed survival, high detection probability at Chipps Island – Parameter Set 42	S _{R0}	0.4	[II]	I or II	I or II	I or II
	S _{A1}	0.3	I or II	I or II	I or II	I or II
	S _{A2}	0.15	I or II	I or II	I or II	I or II
	S _B	0.2	II	I or II	[II]	I or II
	S _F	0.05	I or II	I or II	I or II	I or II
	Ψ _{A1}	0.98	II	I or II	[II]	I or II
	Ψ _{A2}	0.75	I or II	I or II	I or II	I or II
	P _{G1a}	0.9	I or II	I or II	[I]	I or II
	P _{G2b}	0.9	I or II	I or II	[I]	I or II
	S _A	0.038	I or II	I or II	I or II	I or II
	S _R	0.016	[II]	I or II	I or II	I or II

Table 5. Release scenario selected for estimation of each parameter based on the four criteria defined in Table 3, assuming that a 40% probability of remaining in the San Joaquin River at the head of Old River. Release scenarios are: I = (R1 = R2 = 324), and II = (R1 = 500, R2 = 148), where R1 = release size at Durham Ferry, and R2 = size of supplemental release. Release scenarios identified in [brackets] indicate the preferred release scenario in the case when neither release scenario satisfied the criterion. Parameter set numbers refer to Table 6.

Survival Scenario	Parameter	Value	Criterion C1	Criterion C2	Criterion C3	Criterion C4
Low survival, low detection probability at Chipps Island – Parameter Set 2	S _{R0}	0.4	I or II	I or II	I or II	I or II
	S _{A1}	0.3	I or II	I or II	[II]	I or II
	S _{A2}	0.07	I or II	I or II	I	I or II
	S _B	0.07	I or II	I or II	I or II	I or II
	S _F	0.05	I or II	I or II	[I]	I or II
	Ψ _{A1}	0.4	I or II	I or II	I or II	I or II
	Ψ _{A2}	0.75	I or II	I or II	I	I or II
	P _{G1a}	0.5	I or II	I or II	[II]	I or II
	P _{G2b}	0.5	I or II	I or II	[II]	I or II
	S _A	0.020	I or II	I or II	I or II	I or II
	S _R	0.008	I or II	I or II	I or II	I or II
Low survival, high detection probability at Chipps Island – Parameter Set 14	S _{R0}	0.4	I or II	I or II	I or II	I or II
	S _{A1}	0.3	I or II	I or II	[II]	I or II
	S _{A2}	0.07	I or II	I or II	I or II	I or II
	S _B	0.07	I or II	I or II	I or II	I or II
	S _F	0.05	I or II	I or II	I	I or II
	Ψ _{A1}	0.4	I or II	I or II	I or II	I or II
	Ψ _{A2}	0.75	I or II	I or II	I	I or II
	P _{G1a}	0.9	I or II	I or II	[I]	I or II
	P _{G2b}	0.9	I or II	I or II	[I or II]	I or II
	S _A	0.020	I or II	I or II	I or II	I or II
	S _R	0.008	I or II	I or II	I or II	I or II
Low Mixed survival, low detection probability at Chipps Island – Parameter Set 26	S _{R0}	0.4	I or II	I or II	I or II	I or II
	S _{A1}	0.3	I or II	I or II	[II]	I or II
	S _{A2}	0.15	I or II	I or II	[I]	I or II
	S _B	0.2	I or II	I or II	[II]	I or II
	S _F	0.05	I or II	I or II	[I]	I or II
	Ψ _{A1}	0.4	I or II	I or II	I or II	I or II
	Ψ _{A2}	0.75	I or II	I or II	I	I or II
	P _{G1a}	0.5	I or II	I or II	[II]	I or II
	P _{G2b}	0.5	I or II	I or II	[II]	I or II
	S _A	0.038	I or II	I or II	I or II	I or II
	S _R	0.016	I or II	I or II	I or II	I or II
Low Mixed survival, high detection probability at Chipps Island – Parameter Set 38	S _{R0}	0.4	I or II	I or II	I or II	I or II
	S _{A1}	0.3	I or II	I or II	[II]	I or II
	S _{A2}	0.15	I or II	I or II	I	I or II
	S _B	0.2	I or II	I or II	I or II	I or II
	S _F	0.05	I or II	I or II	I	I or II
	Ψ _{A1}	0.4	I or II	I or II	I or II	I or II
	Ψ _{A2}	0.75	I or II	I or II	I	I or II
	P _{G1a}	0.9	I or II	I or II	[II]	I or II
	P _{G2b}	0.9	I or II	I or II	[II]	I or II
	S _A	0.038	I or II	I or II	I or II	I or II
	S _R	0.016	I or II	I or II	I or II	I or II

References

Buchanan, R. A. 2014. Sample Size for 2015 Chinook Tagging Study, prepared for Pat Brandes, 11 December 2014.

Appendix A

Table 6. Simulation results for survival parameter sets ($s_{R0}=0.4$, $s_{A1}=0.3$, $s_{A2}=0.07$ or 0.15 , $s_B=0.07$ or 0.2 , $p_{A1a}=p_{A1b}=p_{B1a}=p_{B1b}=0.90$ or 0.85 , and $p_{A2a}=p_{A2b}=p_{F1a}=p_{F1b}=0.85$). Parameter sets 6 and 18 correspond to the “Low” survival scenario; parameter sets 30 and 42 correspond to the “Low Mixed” survival scenario (see Table 2). C1 = % of simulations in which parameter was successfully estimated. C2 = % of simulations in which parameter was estimated ≤ 1.1 . C3 = standard error of estimate (standard deviation of sampling distribution). C4 = absolute difference between average estimate and true value. Mean = average estimate. Max = maximum estimate. Statistics are calculated over all 10,000 simulations. Highlighted results are summarized in Table 4 and Table 5.

Set	Parameter		R1 = R2 = 324						R1 = 500, R2 = 148					
	Name	Value	C1	C2	C3	C4	Mean	Max	C1	C2	C3	C4	Mean	Max
1	S _{R0}	0.4	100	0	0.028	0.008	0.408	0.534	100	0	0.023	0.007	0.407	0.489
	S _{A1}	0.3	100	0	0.065	0.005	0.295	0.538	100	0	0.052	0.006	0.294	0.494
	S _{A2}	0.07	95	0	0.046	0.006	0.076	0.486	95	0	0.053	0.006	0.076	0.525
	S _B	0.07	95	0	0.046	0.006	0.076	0.503	95	0	0.040	0.006	0.076	0.378
	S _F	0.01	95	0	0.026	0.001	0.011	0.344	95	0	0.034	0.001	0.011	0.400
	Ψ_{A1}	0.4	100	0	0.044	0.005	0.405	0.596	100	0	0.035	0.006	0.406	0.538
	Ψ_{A2}	0.75	100	0	0.042	0.007	0.757	0.911	100	0	0.055	0.014	0.764	0.970
	P _{G1a}	0.5	99.7	0	0.231	0.001	0.501	1.000	99.7	0	0.228	0.002	0.498	1.000
	P _{G2b}	0.5	99.7	0	0.230	0.000	0.500	1.000	99.7	0	0.227	0.003	0.497	1.000
	S _A	0.016	95	0	0.011	0.002	0.018	0.126	95	0	0.013	0.002	0.018	0.115
2	S _R	0.019	95	0	0.012	0.005	0.021	0.125	95	0	0.011	0.005	0.021	0.100
	S _{R0}	0.4	100	0	0.028	0.008	0.408	0.523	100	0	0.023	0.008	0.408	0.510
	S _{A1}	0.3	100	0	0.065	0.005	0.295	0.603	100	0	0.052	0.004	0.296	0.535
	S _{A2}	0.07	96	0	0.046	0.006	0.076	0.535	96	0	0.053	0.005	0.075	0.552
	S _B	0.07	96	0	0.047	0.006	0.076	0.467	96	0	0.040	0.006	0.076	0.473
	S _F	0.05	96	0	0.058	0.004	0.054	0.597	96	0	0.076	0.005	0.055	0.700
	Ψ_{A1}	0.4	100	0	0.044	0.006	0.406	0.578	100	0	0.035	0.005	0.405	0.554
	Ψ_{A2}	0.75	100	0	0.042	0.008	0.758	0.907	100	0	0.054	0.016	0.766	0.946
	P _{G1a}	0.5	99.8	0	0.221	0.000	0.500	1.000	99.8	0	0.217	0.001	0.499	1.000
	P _{G2b}	0.5	99.8	0	0.222	0.000	0.500	1.000	99.8	0	0.219	0.001	0.499	1.000
3	S _A	0.020	96	0	0.013	0.001	0.021	0.155	96	0	0.014	0.001	0.021	0.166
	S _R	0.020	96	0	0.012	0.002	0.022	0.141	96	0	0.011	0.002	0.022	0.121
	S _{R0}	0.4	100	0	0.029	0.009	0.409	0.530	100	0	0.023	0.009	0.409	0.492
	S _{A1}	0.3	100	0	0.053	0.005	0.295	0.532	100	0	0.042	0.005	0.295	0.461
	S _{A2}	0.07	93	0	0.045	0.006	0.076	0.538	92	0	0.051	0.006	0.076	0.463
	S _B	0.07	93	0	0.053	0.006	0.076	0.495	92	0	0.046	0.006	0.076	0.551

	S _F	0.01	93	0	0.024	0.001	0.011	0.250		92	0	0.030	0.001	0.011	0.286
	Ψ _{A1}	0.6	100	0	0.044	0.004	0.604	0.746		100	0	0.035	0.005	0.605	0.738
	Ψ _{A2}	0.75	100	0	0.041	0.009	0.759	0.599		100	0	0.050	0.017	0.767	0.979
	P _{G1a}	0.5	99	0	0.251	0.001	0.501	1.000		99	0	0.253	0.000	0.500	1.000
	P _{G2b}	0.5	99	0	0.248	0.001	0.499	1.000		99	0	0.254	0.000	0.500	1.000
	S _A	0.016	93	0	0.011	0.003	0.018	0.118		92	0	0.012	0.003	0.018	0.122
	S _R	0.015	93	0	0.010	0.000	0.017	0.094		92	0	0.009	0.000	0.017	0.086
4	S _{R0}	0.4	100	0	0.028	0.009	0.409	0.521		100	0	0.023	0.009	0.409	0.496
	S _{A1}	0.3	100	0	0.053	0.005	0.295	0.511		100	0	0.043	0.005	0.295	0.456
	S _{A2}	0.07	95	0	0.044	0.006	0.076	0.671		94	0	0.051	0.005	0.075	0.479
	S _B	0.07	95	0	0.053	0.006	0.076	0.677		94	0	0.045	0.006	0.076	0.524
	S _F	0.05	95	0	0.056	0.006	0.055	0.583		94	0	0.067	0.004	0.054	0.750
	Ψ _{A1}	0.6	100	0	0.043	0.006	0.606	0.770		100	0	0.035	0.005	0.605	0.723
	Ψ _{A2}	0.75	100	0	0.040	0.010	0.760	0.885		100	0	0.050	0.017	0.767	0.938
	P _{G1a}	0.5	99.7	0	0.231	0.004	0.504	1.000		99.6	0	0.241	0.006	0.506	1.000
	P _{G2b}	0.5	99.6	0	0.234	0.005	0.505	1.000		99.6	0	0.240	0.002	0.502	1.000
	S _A	0.020	95	0	0.012	0.005	0.021	0.166		94	0	0.013	0.005	0.021	0.135
	S _R	0.016	95	0	0.010	0.002	0.017	0.111		94	0	0.009	0.002	0.017	0.101
5	S _{R0}	0.4	88	0	0.029	0.013	0.413	0.524		96	0	0.023	0.012	0.412	0.511
	S _{A1}	0.3	100	0	0.041	0.005	0.295	0.461		100	0	0.033	0.005	0.295	0.423
	S _{A2}	0.07	85	0	0.042	0.006	0.076	0.524		77	0	0.046	0.005	0.075	0.479
	S _B	0.07	79	1.3	0.231	0.006	0.076	2.500		76	0.4	0.177	0.005	0.075	1.800
	S _F	0.01	85	0	0.022	0.001	0.011	0.200		77	0	0.024	0.001	0.011	0.242
	Ψ _{A1}	0.98	91	0	0.171	0.033	0.947	0.994		97	0	0.105	0.012	0.968	0.996
	Ψ _{A2}	0.75	100	0	0.038	0.012	0.762	0.904		100	0	0.044	0.021	0.771	0.922
	P _{G1a}	0.5	98	0	0.295	0.002	0.498	1.000		95	0	0.331	0.001	0.501	1.000
	P _{G2b}	0.5	98	0	0.294	0.000	0.500	1.000		95	0	0.327	0.001	0.499	1.000
	S _A	0.016	85	0	0.010	0.011	0.018	0.138		77	0	0.011	0.011	0.018	0.112
	S _R	0.007	75	0	0.005	0.009	0.008	0.065		74	0	0.005	0.009	0.008	0.042
6	S _{R0}	0.4	88	0	0.029	0.014	0.414	0.519		96	0	0.023	0.013	0.413	0.495
	S _{A1}	0.3	100	0	0.041	0.005	0.295	0.465		100	0	0.033	0.005	0.295	0.413
	S _{A2}	0.07	90	0	0.042	0.006	0.076	0.412		82	0	0.046	0.005	0.075	0.463
	S _B	0.07	83	1	0.239	0.011	0.081	2.667		81	1	0.193	0.005	0.075	3.000
	S _F	0.05	90	0	0.053	0.004	0.054	0.477		82	0	0.059	0.003	0.053	0.635
	Ψ _{A1}	0.98	91	0	0.157	0.028	0.952	0.994		97	0	0.110	0.013	0.967	0.996
	Ψ _{A2}	0.75	100	0	0.038	0.012	0.762	0.894		100	0	0.044	0.021	0.771	0.934

	P _{G1a}	0.5	99	0	0.267	0.004	0.496	1.000		97	0	0.308	0.001	0.499	1.000
	P _{G2b}	0.5	99	0	0.268	0.002	0.498	1.000		97	0	0.307	0.005	0.495	1.000
	S _A	0.020	90	0	0.011	0.013	0.021	0.123		82	0	0.012	0.013	0.021	0.136
	S _R	0.008	78	0	0.005	0.010	0.009	0.055		79	0	0.005	0.010	0.009	0.054
7	S _{R0}	0.4	100	0	0.028	0.008	0.408	0.530		100	0	0.023	0.007	0.407	0.495
	S _{A1}	0.3	98	0	0.065	0.004	0.296	0.563		92	0	0.053	0.004	0.296	0.529
	S _{A2}	0.07	96	0	0.043	0.006	0.076	0.560		96	0	0.048	0.006	0.076	0.490
	S _B	0.07	96	0	0.048	0.007	0.077	0.465		96	0	0.040	0.007	0.077	0.489
	S _F	0.01	96	0.02	0.067	0.002	0.012	1.667		92	0.2	0.094	0.003	0.0123	1.800
	Ψ _{A1}	0.4	100	0	0.043	0.005	0.405	0.584		100	0	0.035	0.005	0.405	0.538
	Ψ _{A2}	0.95	98	0	0.043	0.004	0.954	1.008		92	0	0.082	0.005	0.955	1.033
	P _{G1a}	0.5	99.8	0	0.219	0.002	0.498	1.000		99.9	0	0.218	0.000	0.500	1.000
	P _{G2b}	0.5	99.9	0	0.220	0.003	0.497	1.000		99.8	0	0.218	0.001	0.499	1.000
	S _A	0.020	94	0	0.013	0.002	0.022	0.178		88	0	0.015	0.002	0.022	0.146
	S _R	0.020	94	0	0.012	0.002	0.022	0.135		88	0	0.011	0.002	0.022	0.150
8	S _{R0}	0.4	100	0	0.028	0.008	0.408	0.516		100	0	0.023	0.007	0.407	0.498
	S _{A1}	0.3	98	0	0.066	0.004	0.296	0.609		92	0	0.052	0.003	0.297	0.487
	S _{A2}	0.07	97	0	0.041	0.005	0.075	0.596		96	0	0.050	0.006	0.076	0.552
	S _B	0.07	97	0	0.046	0.006	0.076	0.560		96	0	0.040	0.006	0.076	0.490
	S _F	0.05	96	0.2	0.133	0.002	0.052	1.731		93	0.4	0.178	0.006	0.056	2.909
	Ψ _{A1}	0.4	100	0	0.044	0.006	0.406	0.566		100	0	0.035	0.006	0.406	0.541
	Ψ _{A2}	0.95	98	0	0.038	0.005	0.955	1.008		92	0	0.075	0.007	0.957	1.033
	P _{G1a}	0.5	99.9	0	0.213	0.002	0.502	1.000		99.8	0	0.215	0.003	0.503	1.000
	P _{G2b}	0.5	99.9	0	0.214	0.002	0.502	1.000		99.9	0	0.214	0.002	0.498	1.000
	S _A	0.021	95	0	0.013	0.002	0.022	0.281		88	0	0.015	0.002	0.022	0.188
	S _R	0.020	95	0	0.012	0.001	0.022	0.163		88	0	0.011	0.001	0.022	0.115
9	S _{R0}	0.4	100	0	0.028	0.010	0.410	0.515		100	0	0.023	0.009	0.409	0.494
	S _{A1}	0.3	99	0	0.053	0.004	0.296	0.504		95	0	0.043	0.003	0.297	0.470
	S _{A2}	0.07	95	0	0.041	0.006	0.076	0.467		94	0	0.046	0.006	0.076	0.492
	S _B	0.07	95	0	0.053	0.006	0.076	0.556		94	0	0.046	0.006	0.076	0.473
	S _F	0.01	95	0.01	0.060	0.001	0.011	1.375		92	0.04	0.067	0.000	0.010	1.312
	Ψ _{A1}	0.6	100	0	0.044	0.005	0.605	0.761		100	0	0.035	0.005	0.605	0.731
	Ψ _{A2}	0.95	99	0	0.051	0.006	0.956	1.014		95	0	0.079	0.010	0.960	1.033
	P _{G1a}	0.5	99.6	0	0.230	0.000	0.500	1.000		99.5	0	0.239	0.000	0.500	1.000
	P _{G2b}	0.5	99.7	0	0.229	0.001	0.499	1.000		99.5	0	0.239	0.001	0.501	1.000
	S _A	0.020	94	0	0.012	0.006	0.022	0.140		89	0	0.014	0.006	0.022	0.137

	S _R	0.016	94	0	0.010	0.002	0.018	0.123		89	0	0.009	0.002	0.018	0.100
10	S _{R0}	0.4	100	0	0.029	0.009	0.409	0.521		100	0	0.023	0.009	0.409	0.487
	S _{A1}	0.3	99	0	0.053	0.004	0.296	0.516		94	0	0.043	0.004	0.296	0.465
	S _{A2}	0.07	95	0	0.041	0.006	0.076	0.505		94	0	0.047	0.006	0.076	0.529
	S _B	0.07	95	0	0.054	0.006	0.076	0.656		94	0	0.046	0.006	0.076	0.535
	S _F	0.05	94	0.1	0.132	0.005	0.055	2.143		92	0.4	0.166	0.006	0.056	2.800
	Ψ _{A1}	0.6	100	0	0.043	0.005	0.605	0.755		100	0	0.035	0.005	0.605	0.735
	Ψ _{A2}	0.95	99	0	0.051	0.005	0.955	1.012		95	0	0.070	0.012	0.962	1.043
	P _{G1a}	0.5	99.8	0	0.230	0.000	0.500	1.000		99.6	0	0.239	0.002	0.502	1.000
	P _{G2b}	0.5	99.8	0	0.230	0.000	0.500	1.000		99.6	0	0.237	0.001	0.501	1.000
	S _A	0.021	93	0	0.013	0.006	0.022	0.159		89	0	0.014	0.006	0.022	0.142
	S _R	0.016	93	0	0.010	0.003	0.018	0.109		89	0	0.009	0.003	0.018	0.094
11	S _{R0}	0.4	88	0	0.029	0.014	0.414	0.513		96	0	0.023	0.013	0.413	0.493
	S _{A1}	0.3	99	0	0.041	0.004	0.296	0.466		98	0	0.034	0.004	0.296	0.440
	S _{A2}	0.07	90	0	0.039	0.005	0.075	0.401		84	0	0.042	0.005	0.075	0.460
	S _B	0.07	83	1.3	0.237	0.008	0.078	3.000		82	4.1	0.180	0.004	0.074	2.000
	S _F	0.01	90	0	0.051	0.000	0.010	0.781		83	0.01	0.057	0.000	0.010	1.429
	Ψ _{A1}	0.98	90	0	0.162	0.030	0.950	0.994		97	0	0.116	0.015	0.965	0.996
	Ψ _{A2}	0.95	99	0	0.040	0.010	0.960	1.018		98	0	0.060	0.019	0.969	1.034
	P _{G1a}	0.5	99	0	0.265	0.000	0.500	1.000		97	0	0.302	0.000	0.500	1.000
	P _{G2b}	0.5	99	0	0.267	0.001	0.501	1.000		97	0	0.302	0.001	0.499	1.000
	S _A	0.020	89	0	0.012	0.013	0.021	0.123		82	0	0.012	0.013	0.022	0.135
	S _R	0.008	78	0	0.005	0.011	0.009	0.052		78	0	0.005	0.011	0.009	0.058
12	S _{R0}	0.4	89	0	0.029	0.013	0.413	0.544		96	0	0.023	0.013	0.413	0.501
	S _{A1}	0.3	99	0	0.042	0.005	0.295	0.455		98	0	0.034	0.003	0.297	0.423
	S _{A2}	0.07	91	0	0.039	0.005	0.075	0.494		84	0	0.042	0.005	0.075	0.409
	S _B	0.07	84	1.2	0.221	0.002	0.072	2.286		84	0.5	0.184	0.005	0.075	2.000
	S _F	0.05	91	0.07	0.123	0.004	0.054	1.604		84	0.2	0.141	0.004	0.054	1.600
	Ψ _{A1}	0.98	91	0	0.151	0.026	0.954	0.994		98	0	0.109	0.013	0.967	0.996
	Ψ _{A2}	0.95	99	0	0.038	0.010	0.960	1.021		98	0	0.061	0.019	0.969	1.037
	P _{G1a}	0.5	99	0	0.264	0.000	0.500	1.000		98	0	0.296	0.004	0.504	1.000
	P _{G2b}	0.5	99	0	0.261	0.001	0.499	1.000		98	0	0.297	0.006	0.506	1.000
	S _A	0.021	90	0	0.012	0.013	0.022	0.171		83	0	0.012	0.013	0.022	0.119
	S _R	0.009	80	0	0.005	0.011	0.010	0.058		80	0	0.005	0.011	0.010	0.046
13	S _{R0}	0.4	100	0	0.028	0.008	0.408	0.537		100	0	0.023	0.007	0.407	0.492
	S _{A1}	0.3	100	0	0.065	0.004	0.296	0.569		100	0	0.052	0.005	0.295	0.526

	S _{A2}	0.07	100	0	0.029	0.000	0.070	0.190		100	0	0.037	0.001	0.071	0.244
	S _B	0.07	100	0	0.029	0.000	0.070	0.196		100	0	0.024	0.000	0.070	0.165
	S _F	0.01	100	0	0.019	0.000	0.010	0.150		100	0	0.025	0.000	0.010	0.222
	Ψ _{A1}	0.4	100	0	0.044	0.005	0.405	0.576		100	0	0.035	0.006	0.406	0.543
	Ψ _{A2}	0.75	100	0	0.041	0.008	0.758	0.907		100	0	0.054	0.014	0.764	0.938
	P _{G1a}	0.9	100	0	0.096	0.002	0.902	1.000		100	0	0.095	0.001	0.901	1.000
	P _{G2b}	0.9	100	0	0.099	0.002	0.898	1.000		100	0	0.095	0.000	0.900	1.000
	S _A	0.016	100	0	0.008	0.003	0.016	0.062		100	0	0.009	0.003	0.017	0.061
	S _R	0.019	100	0	0.007	0.003	0.020	0.053		100	0	0.006	0.003	0.020	0.046
14	S _{R0}	0.4	100	0	0.028	0.008	0.408	0.510		100	0	0.023	0.007	0.407	0.492
	S _{A1}	0.3	100	0	0.065	0.005	0.295	0.563		100	0	0.052	0.004	0.296	0.525
	S _{A2}	0.07	100	0	0.028	0.000	0.070	0.206		100	0	0.036	0.000	0.070	0.224
	S _B	0.07	100	0	0.030	0.000	0.070	0.209		100	0	0.024	0.000	0.070	0.184
	S _F	0.05	100	0	0.042	0.001	0.049	0.304		100	0	0.056	0.001	0.051	0.377
	Ψ _{A1}	0.4	100	0	0.044	0.005	0.405	0.594		100	0	0.035	0.005	0.405	0.532
	Ψ _{A2}	0.75	100	0	0.042	0.007	0.757	0.903		100	0	0.055	0.014	0.764	0.968
	P _{G1a}	0.9	100	0	0.093	0.001	0.899	1.000		100	0	0.094	0.000	0.900	1.000
	P _{G2b}	0.9	100	0	0.093	0.001	0.899	1.000		100	0	0.093	0.000	0.900	1.000
	S _A	0.020	100	0	0.008	0.001	0.019	0.058		100	0	0.010	0.001	0.019	0.068
	S _R	0.020	100	0	0.007	0.001	0.020	0.053		100	0	0.006	0.001	0.020	0.047
15	S _{R0}	0.4	100	0	0.029	0.010	0.410	0.509		100	0	0.023	0.009	0.409	0.495
	S _{A1}	0.3	100	0	0.053	0.005	0.295	0.530		100	0	0.042	0.005	0.295	0.506
	S _{A2}	0.07	100	0	0.028	0.000	0.070	0.182		100	0	0.034	0.000	0.070	0.240
	S _B	0.07	100	0	0.036	0.001	0.071	0.275		100	0	0.029	0.000	0.070	0.217
	S _F	0.01	100	0	0.018	0.000	0.010	0.158		100	0	0.024	0.000	0.010	0.312
	Ψ _{A1}	0.6	100	0	0.043	0.005	0.605	0.764		100	0	0.035	0.005	0.605	0.746
	Ψ _{A2}	0.75	100	0	0.040	0.009	0.759	0.890		100	0	0.051	0.016	0.766	0.935
	P _{G1a}	0.9	100	0	0.104	0.000	0.900	1.000		100	0	0.106	0.001	0.599	1.000
	P _{G2b}	0.9	100	0	0.105	0.001	0.901	1.000		100	0	0.105	0.002	0.902	1.000
	S _A	0.016	100	0	0.007	0.001	0.016	0.051		100	0	0.008	0.001	0.017	0.057
	S _R	0.015	100	0	0.006	0.001	0.015	0.044		100	0	0.005	0.001	0.015	0.042
16	S _{R0}	0.4	100	0	0.029	0.009	0.409	0.533		100	0	0.023	0.009	0.409	0.491
	S _{A1}	0.3	100	0	0.053	0.005	0.295	0.513		100	0	0.042	0.005	0.295	0.466
	S _{A2}	0.07	100	0	0.027	0.000	0.070	0.215		100	0	0.033	0.000	0.070	0.238
	S _B	0.07	100	0	0.037	0.000	0.070	0.229		100	0	0.029	0.0000	0.070	0.190
	S _F	0.05	100	0	0.041	0.000	0.050	0.286		100	0	0.051	.000	0.050	0.300

	Ψ_{A1}	0.6	100	0	0.043	0.005	0.605	0.770		100	0	0.035	0.005	0.605	0.733
	Ψ_{A2}	0.75	100	0	0.040	0.010	0.760	0.908		100	0	0.050	0.017	0.767	0.930
	P_{G1a}	0.9	100	0	0.099	0.000	0.900	1.000		100	0	0.102	0.001	0.899	1.000
	P_{G2b}	0.9	100	0	0.098	0.001	0.901	1.000		100	0	0.103	0.002	0.898	1.000
	S_A	0.020	100	0	0.008	0.003	0.019	0.054		100	0	0.009	0.004	0.019	0.068
	S_R	0.016	100	0	0.006	0.003	0.016	0.046		100	0	0.005	0.003	0.016	0.043
17	S_{R0}	0.4	88	0	0.029	0.013	0.413	0.535		96	0	0.023	0.013	0.413	0.511
	S_{A1}	0.3	100	0	0.041	0.005	0.295	0.46*9		100	0	0.033	0.005	0.295	0.408
	S_{A2}	0.07	99.8	0	0.026	0.000	0.070	0.203		99	0	0.030	0.001	0.071	0.190
	S_B	0.07	92	0.03	0.167	0.005	0.065	1.333		97	0.02	0.149	0.000	0.070	1.125
	S_F	0.01	99.8	0	0.018	0.000	0.010	0.154		99	0	0.021	0.000	0.010	0.192
	Ψ_{A1}	0.98	90	0	0.162	0.030	0.950	0.994		97	0	0.108	0.013	0.967	0.996
	Ψ_{A2}	0.75	100	0	0.038	0.011	0.761	0.891		100	0	0.044	0.021	0.771	0.913
	P_{G1a}	0.9	100	0	0.129	0.000	0.900	1.000		99.5	0	0.145	0.001	0.901	1.000
	P_{G2b}	0.9	99.9	0	0.127	0.000	0.900	1.000		99.6	0	0.145	0.001	0.901	1.000
	S_A	0.016	99.8	0	0.006	0.009	0.016	0.045		99	0	0.007	0.010	0.017	0.052
	S_R	0.007	87	0	0.003	0.009	0.007	0.023		95	0	0.003	0.009	0.007	0.022
18	S_{R0}	0.4	88	0	0.029	0.013	0.413	0.522		96	0	0.023	0.012	0.412	0.508
	S_{A1}	0.3	100	0	0.041	0.005	0.295	0.458		100	0	0.033	0.004	0.296	0.444
	S_{A2}	0.07	99.9	0	0.026	0.001	0.071	0.207		99.6	0	0.030	0.000	0.070	0.214
	S_B	0.07	92	0	0.178	0.001	0.069	1.089		98	0.01	0.149	0.000	0.070	1.111
	S_F	0.05	99.9	0	0.039	0.000	0.050	0.245		99.6	0	0.045	0.001	0.051	0.295
	Ψ_{A1}	0.98	90	0	0.159	0.029	0.951	0.994		97	0	0.097	0.010	0.970	0.996
	Ψ_{A2}	0.75	100	0	0.038	0.012	0.762	0.892		100	0	0.044	0.021	0.771	0.916
	P_{G1a}	0.9	99.9	0	0.115	0.001	0.899	1.000		99.8	0	0.133	0.000	0.900	1.000
	P_{G2b}	0.9	100	0	0.115	0.001	0.899	1.000		99.8	0	0.132	0.002	0.902	1.000
	S_A	0.020	99.9	0	0.007	0.011	0.019	0.051		99.6	0	0.008	0.011	0.019	0.058
	S_R	0.008	88	0	0.003	0.011	0.008	0.026		96	0	0.003	0.011	0.008	0.025
19	S_{R0}	0.4	100	0	0.029	0.008	0.408	0.514		100	0	0.023	0.008	0.408	0.497
	S_{A1}	0.3	99	0	0.066	0.004	0.296	0.569		92	0	0.053	0.004	0.296	0.493
	S_{A2}	0.07	100	0	0.025	0.000	0.070	0.178		100	0	0.033	0.000	0.070	0.251
	S_B	0.07	100	0	0.029	0.000	0.070	0.202		100	0	0.024	0.000	0.070	0.171
	S_F	0.01	99.6	0	0.046	0.001	0.009	1.048		96	0	0.067	0.001	0.011	1.071
	Ψ_{A1}	0.4	100	0	0.044	0.005	0.405	0.560		100	0	0.036	0.006	0.406	0.542
	Ψ_{A2}	0.95	99	0	0.034	0.005	0.955	1.012		92	0	0.089	0.004	0.954	1.030
	P_{G1a}	0.9	100	0	0.092	0.001	0.901	1.000		100	0	0.091	0.001	0.901	1.000

	P _{G2b}	0.9	100	0	0.092	0.001	0.899	1.000		100	0	0.092	0.000	0.900	1.000
	S _A	0.020	99	0	0.009	0.000	0.020	0.064		91	0	0.010	0.000	0.020	0.082
	S _R	0.020	99	0	0.007	0.000	0.020	0.060		91	0	0.006	0.000	0.020	0.046
20	S _{R0}	0.4	100	0	0.028	0.007	0.407	0.520		100	0	0.022	0.007	0.407	0.496
	S _{A1}	0.3	98	0	0.065	0.003	0.297	0.612		92	0	0.052	0.003	0.297	0.491
	S _{A2}	0.07	100	0	0.025	0.000	0.070	0.189		100	0	0.032	0.000	0.070	0.239
	S _B	0.07	100	0	0.029	0.000	0.070	0.194		100	0	0.024	0.000	0.070	0.171
	S _F	0.05	99.6	0	0.104	0.001	0.051	1.024		97	0.01	0.134	0.000	0.050	1.250
	Ψ _{A1}	0.4	100	0	0.044	0.006	0.406	0.578		100	0	0.035	0.006	0.406	0.537
	Ψ _{A2}	0.95	99	0	0.044	0.004	0.954	1.010		92	0	0.083	0.005	0.955	1.035
	P _{G1a}	0.9	100	0	0.092	0.000	0.900	1.000		100	0	0.092	0.001	0.899	1.000
	P _{G2b}	0.9	100	0	0.090	0.000	0.900	1.000		100	0	0.091	0.001	0.901	1.000
	S _A	0.021	98	0	0.009	0.001	0.021	0.070		92	0	0.010	0.000	0.020	0.085
	S _R	0.020	98	0	0.007	0.000	0.020	0.052		92	0	0.006	0.000	0.020	0.051
21	S _{R0}	0.4	100	0	0.029	0.009	0.409	0.516		100	0	0.023	0.009	0.409	0.499
	S _{A1}	0.3	99	0	0.054	0.004	0.296	0.515		95	0	0.043	0.003	0.297	0.486
	S _{A2}	0.07	100	0	0.024	0.000	0.070	0.188		100	0	0.030	0.000	0.070	0.208
	S _B	0.07	100	0	0.036	0.000	0.070	0.225		100	0	0.029	0.001	0.071	0.190
	S _F	0.01	99.7	0	0.049	0.001	0.011	1.013		98	0	0.057	0.000	0.010	1.018
	Ψ _{A1}	0.6	100	0	0.043	0.005	0.605	0.803		100	0	0.035	0.004	0.605	0.758
	Ψ _{A2}	0.95	99	0	0.038	0.007	0.957	1.016		95	0	0.071	0.012	0.962	1.039
	P _{G1a}	0.9	100	0	0.099	0.002	0.898	1.000		100	0	0.101	0.000	0.900	1.000
	P _{G2b}	0.9	100	0	0.099	0.000	0.900	1.000		100	0	0.100	0.000	0.900	1.000
	S _A	0.020	99	0	0.008	0.000	0.020	0.076		95	0	0.009	0.000	0.020	0.073
	S _R	0.016	99	0	0.006	0.000	0.016	0.045		95	0	0.005	0.000	0.016	0.037
22	S _{R0}	0.4	100	0	0.029	0.009	0.409	0.526		100	0	0.023	0.009	0.409	0.498
	S _{A1}	0.3	99	0	0.053	0.004	0.296	0.498		95	0	0.042	0.003	0.297	0.462
	S _{A2}	0.07	100	0	0.024	0.000	0.070	0.188		100	0	0.030	0.000	0.070	0.195
	S _B	0.07	100	0	0.036	0.000	0.070	0.263		100	0	0.029	0.000	0.070	0.197
	S _F	0.05	99.7	0	0.100	0.001	0.049	1.046		98	0	0.127	0.000	0.050	1.091
	Ψ _{A1}	0.6	100	0	0.044	0.005	0.605	0.767		100	0	0.035	0.005	0.605	0.736
	Ψ _{A2}	0.95	99	0	0.041	0.006	0.956	1.012		95	0	0.075	0.011	0.961	1.032
	P _{G1a}	0.9	100	0	0.096	0.000	0.900	1.000		100	0	0.100	0.000	0.900	1.000
	P _{G2b}	0.9	100	0	0.096	0.001	0.901	1.000		100	0	0.100	0.001	0.901	1.000
	S _A	0.021	99	0	0.008	0.000	0.020	0.059		95	0	0.009	0.000	0.021	0.068
	S _R	0.016	99	0	0.006	0.000	0.016	0.051		95	0	0.005	0.000	0.016	0.037

23	S _{R0}	0.4	88	0	0.029	0.014	0.414	0.552		96	0	0.023	0.013	0.413	0.509
	S _{A1}	0.3	99	0	0.042	0.005	0.4295	0.468		98	0	0.034	0.003	0.297	0.426
	S _{A2}	0.07	100	0	0.023	0.000	0.070	0.179		99.7	0	0.026	0.000	0.070	0.184
	S _B	0.07	93	0.04	0.179	0.001	0.069	1.250		98	0.02	0.150	0.001	0.071	1.125
	S _F	0.01	99.7	0	0.045	0.001	0.011	1.037		99	0	0.051	0.000	0.010	1.000
	Ψ _{A1}	0.98	91	0	0.165	0.031	0.949	0.994		97	0	0.109	0.013	0.967	0.996
	Ψ _{A2}	0.95	99	0	0.043	0.010	0.960	1.017		98	0	0.068	0.018	0.968	1.038
	P _{G1a}	0.9	99.9	0	0.113	0.000	0.900	1.000		99.8	0	0.130	0.000	0.900	1.000
	P _{G2b}	0.9	100	0	0.112	0.000	0.900	1.000		99.9	0	0.132	0.002	0.898	1.000
	S _A	0.020	99	0	0.007	0.000	0.020	0.052		97	0	0.008	0.000	0.020	0.056
	S _R	0.008	87	0	0.003	0.000	0.009	0.031		93	0	0.003	0.000	0.009	0.023
24	S _{R0}	0.4	88	0	0.029	0.013	0.413	0.515		96	0	0.023	0.013	0.413	0.503
	S _{A1}	0.3	99	0	0.041	0.003	0.297	0.477		97	0	0.034	0.004	0.296	0.432
	S _{A2}	0.07	100	0	0.023	0.000	0.070	0.162		99.7	0	0.026	0.000	0.070	0.192
	S _B	0.07	93	0.08	0.187	0.004	0.074	1.200		98	0.01	0.151	0.002	0.072	1.111
	S _F	0.05	99.8	0	0.093	0.000	0.050	1.029		99	0	0.110	0.000	0.050	1.059
	Ψ _{A1}	0.98	91	0	0.168	0.032	0.948	0.994		97	0	0.104	0.012	0.968	0.996
	Ψ _{A2}	0.95	99.5	0	0.037	0.010	0.960	1.018		98	0	0.065	0.018	0.968	1.038
	P _{G1a}	0.9	100	0	0.109	0.000	0.900	1.000		99.8	0	0.129	0.001	0.899	1.000
	P _{G2b}	0.9	100	0	0.110	0.001	0.899	1.000		99.8	0	0.127	0.002	0.902	1.000
	S _A	0.021	99	0	0.007	0.000	0.021	0.056		97	0	0.008	0.000	0.021	0.063
	S _R	0.009	87	0	0.003	0.000	0.009	0.028		93	0	0.003	0.000	0.009	0.029
25	S _{R0}	0.4	100	0	0.029	0.013	0.413	0.540		100	0	0.023	0.013	0.413	0.508
	S _{A1}	0.3	100	0	0.064	0.005	0.295	0.564		100	0	0.052	0.005	0.295	0.521
	S _{A2}	0.15	99.9	0	0.064	0.008	0.158	1.063		100	0	0.073	0.008	0.158	0.932
	S _B	0.2	99.9	0.01	0.078	0.011	0.211	1.286		100	0	0.064	0.009	0.209	1.065
	S _F	0.01	99.9	0	0.024	0.000	0.010	0.257		100	0	0.031	0.001	0.011	0.321
	Ψ _{A1}	0.4	100	0	0.044	0.000	0.400	0.583		100	0	0.035	0.000	0.400	0.547
	Ψ _{A2}	0.75	100	0	0.041	0.008	0.758	0.915		100	0	0.055	0.014	0.764	0.963
	P _{G1a}	0.5	100	0	0.138	0.000	0.500	1.000		100	0	0.129	0.001	0.499	1.000
	P _{G2b}	0.5	100	0	0.138	0.001	0.499	1.000		100	0	0.129	0.002	0.498	1.000
	S _A	0.034	99.9	0	0.017	0.002	0.036	0.265		100	0	0.018	0.002	0.036	0.239
	S _R	0.054	99.9	0	0.021	0.005	0.058	0.335		100	0	0.017	0.004	0.058	0.296
26	S _{R0}	0.4	100	0	0.029	0.013	0.413	0.521		100	0	0.023	0.013	0.413	0.502
	S _{A1}	0.3	100	0	0.064	0.006	0.294	0.552		100	0	0.053	0.006	0.294	0.491

	S _{A2}	0.15	100	0	0.060	0.007	0.157	0.751	100	0.01	0.073	0.006	0.156	1.195
	S _B	0.2	100	0.02	0.075	0.009	0.209	1.243	100	0	0.065	0.010	0.210	0.990
	S _F	0.05	100	0	0.054	0.003	0.053	0.533	100	0	0.071	0.003	0.053	0.808
	Ψ _{A1}	0.4	100	0	0.044	0.000	0.400	0.570	100	0	0.035	0.001	0.399	0.556
	Ψ _{A2}	0.75	100	0	0.041	0.007	0.757	0.907	100	0	0.054	0.015	0.765	0.963
	P _{G1a}	0.5	100	0	0.134	0.002	0.502	1.000	100	0	0.130	0.003	0.497	1.000
	P _{G2b}	0.5	100	0	0.135	0.002	0.502	1.000	100	0	0.129	0.001	0.499	1.000
	S _A	0.038	100	0	0.017	0.001	0.039	0.205	100	0	0.019	0.001	0.039	0.210
	S _R	0.054	100	0	0.020	0.004	0.058	0.381	100	0	0.018	0.004	0.058	0.271
27	S _{R0}	0.4	100	0	0.029	0.012	0.412	0.527	100	0	0.023	0.013	0.413	0.504
	S _{A1}	0.3	100	0	0.053	0.005	0.295	0.511	100	0	0.042	0.004	0.296	0.451
	S _{A2}	0.15	99.7	0	0.065	0.009	0.159	0.909	99.8	0	0.070	0.008	0.158	0.847
	S _B	0.2	99.7	0	0.091	0.014	0.214	1.043	99.8	0	0.079	0.011	0.211	1.028
	S _F	0.01	99.7	0	0.024	0.001	0.011	0.245	99.8	0	0.029	0.000	0.010	0.407
	Ψ _{A1}	0.6	100	0	0.044	0.000	0.600	0.759	100	0	0.035	0.001	0.599	0.730
	Ψ _{A2}	0.75	100	0	0.041	0.009	0.759	0.923	100	0	0.050	0.017	0.767	0.951
	P _{G1a}	0.5	100	0	0.150	0.003	0.497	1.000	100	0	0.147	0.003	0.503	1.000
	P _{G2b}	0.5	100	0	0.150	0.000	0.500	1.000	100	0	0.147	0.001	0.501	1.000
	S _A	0.034	99.7	0	0.016	0.002	0.036	0.215	99.8	0	0.017	0.002	0.037	0.215
	S _R	0.040	99.7	0	0.017	0.004	0.044	0.224	99.8	0	0.015	0.004	0.044	0.233
28	S _{R0}	0.4	100	0	0.029	0.013	0.413	0.529	100	0	0.023	0.013	0.413	0.493
	S _{A1}	0.3	100	0	0.052	0.005	0.295	0.488	100	0	0.043	0.005	0.295	0.455
	S _{A2}	0.15	99.9	0	0.064	0.008	0.158	0.804	99.7	0	0.073	0.008	0.158	1.067
	S _B	0.2	99.9	0.03	0.092	0.011	0.211	1.250	99.7	0.01	0.079	0.012	0.212	1.106
	S _F	0.05	99.9	0	0.053	0.002	0.052	0.515	99.7	0	0.065	0.003	0.053	0.891
	Ψ _{A1}	0.6	100	0	0.044	0.000	0.600	0.769	100	0	0.035	0.000	0.600	0.733
	Ψ _{A2}	0.75	100	0	0.040	0.010	0.760	0.896	100	0	0.050	0.017	0.767	0.942
	P _{G1a}	0.5	100	0	0.147	0.001	0.501	1.000	100	0	0.147	0.002	0.498	1.000
	P _{G2b}	0.5	100	0	0.149	0.003	0.503	1.000	100	0	0.147	0.001	0.499	1.000
	S _A	0.038	99.9	0	0.017	0.002	0.039	0.188	99.7	0	0.018	0.002	0.039	0.290
	S _R	0.041	99.9	0	0.017	0.004	0.045	0.269	99.7	0	0.016	0.004	0.045	0.248
29	S _{R0}	0.4	95	0	0.029	0.013	0.413	0.545	94	0	0.024	0.013	0.413	0.508
	S _{A1}	0.3	100	0	0.042	0.005	0.295	0.449	100	0	0.033	0.005	0.295	0.436
	S _{A2}	0.15	98	0	0.070	0.013	0.163	0.794	96	0	0.079	0.014	0.164	1.014
	S _B	0.2	91	4.4	0.384	0.017	0.217	4.267	94	2.6	0.318	0.018	0.218	3.273
	S _F	0.01	98	0	0.024	0.001	0.011	0.312	96	0	0.027	0.001	0.011	0.328

	Ψ_{A1}	0.98	90	0	0.218	0.055	0.925	0.994		97	0	0.150	0.025	0.955	0.996
	Ψ_{A2}	0.75	100	0	0.038	0.011	0.761	0.892		100	0	0.045	0.020	0.770	0.932
	P_{G1a}	0.5	100	0	0.190	0.001	0.501	1.000		99.8	0	0.221	0.000	0.500	1.000
	P_{G2b}	0.5	100	0	0.190	0.000	0.500	1.000		99.8	0	0.220	0.002	0.498	1.000
	S_A	0.034	98	0	0.017	0.003	0.037	0.205		96	0	0.019	0.004	0.038	0.231
	S_R	0.015	83	0	0.008	0.002	0.017	0.107		90	0	0.009	0.002	0.017	0.101
30	S_{R0}	0.4	85	0	0.029	0.014	0.414	0.518		96	0	0.024	0.013	0.413	0.497
	S_{A1}	0.3	100	0	0.041	0.005	0.295	0.469		100	0	0.034	0.004	0.296	0.429
	S_{A2}	0.15	99	0	0.072	0.012	0.162	0.953		97	0	0.077	0.013	0.163	0.860
	S_B	0.2	91	4.6	0.390	0.024	0.224	4.267		95	2.2	0.313	0.017	0.217	3.273
	S_F	0.05	99	0	0.054	0.004	0.054	0.735		97	0	0.061	0.004	0.054	0.625
	Ψ_{A1}	0.98	90	0	0.208	0.050	0.930	0.994		97	0	0.141	0.022	0.958	0.996
	Ψ_{A2}	0.75	100	0	0.038	0.012	0.762	0.886		100	0	0.045	0.021	0.771	0.934
	P_{G1a}	0.5	100	0	0.183	0.001	0.499	1.000		99.9	0	0.210	0.003	0.503	1.000
	P_{G2b}	0.5	100	0	0.183	0.003	0.497	1.000		100	0	0.208	0.000	0.500	1.000
	S_A	0.038	99	0	0.019	0.003	0.040	0.261		97	0	0.019	0.003	0.041	0.222
	S_R	0.016	84	0	0.009	0.002	0.018	0.092		92	0	0.008	0.002	0.018	0.085
31	S_{R0}	0.4	100	0	0.029	0.013	0.413	0.525		100	0	0.023	0.013	0.413	0.501
	S_{A1}	0.3	98	0	0.065	0.004	0.296	0.554		92	0	0.052	0.004	0.296	0.506
	S_{A2}	0.15	100	0	0.055	0.007	0.157	0.770		100	0.01	0.064	0.006	0.156	1.212
	S_B	0.2	100	0.01	0.072	0.009	0.209	1.189		100	0.01	0.062	0.007	0.207	1.345
	S_F	0.01	99.6	0.04	0.063	0.001	0.011	1.560		97	0.10	0.080	0.001	0.011	2.333
	Ψ_{A1}	0.4	100	0	0.044	0.000	0.400	0.585		100	0	0.036	0.000	0.400	0.549
	Ψ_{A2}	0.95	99	0	0.049	0.003	0.953	1.017		92	0	0.079	0.005	0.955	1.032
	P_{G1a}	0.5	100	0	0.129	0.002	0.498	1.000		100	0	0.125	0.001	0.501	1.000
	P_{G2b}	0.5	100	0	0.132	0.000	0.500	1.000		100	0	0.126	0.002	0.502	1.000
	S_A	0.043	98	0	0.019	0.002	0.044	0.284		92	0	0.020	0.002	0.044	0.305
	S_R	0.055	98	0	0.019	0.004	0.059	0.289		92	0	0.017	0.004	0.059	0.390
32	S_{R0}	0.4	100	0	0.029	0.013	0.413	0.520		100	0	0.023	0.013	0.413	0.507
	S_{A1}	0.3	99	0	0.065	0.005	0.295	0.556		91	0	0.052	0.004	0.296	0.500
	S_{A2}	0.15	100	0	0.057	0.007	0.157	0.755		100	0	0.063	0.005	0.155	0.799
	S_B	0.2	100	0.03	0.076	0.009	0.209	1.414		100	0	0.062	0.007	0.207	1.024
	S_F	0.05	99.6	0.2	0.149	0.004	0.054	5.400		96	0.6	0.175	0.006	0.056	1.965
	Ψ_{A1}	0.4	100	0	0.044	0.001	0.401	0.568		100	0	0.035	0.000	0.400	0.522
	Ψ_{A2}	0.95	99	0	0.039	0.004	0.954	1.011		92	0	0.085	0.005	0.955	1.030
	P_{G1a}	0.5	100	0	0.131	0.000	0.500	1.000		100	0	0.125	0.002	0.502	0.917

	P _{G2b}	0.5	100	0	0.130	0.000	0.500	1.000		100	0	0.125	0.002	0.498	1.000
	S _A	0.044	99	0	0.019	0.001	0.045	0.323		91	0	0.020	0.001	0.045	0.216
	S _R	0.055	99	0	0.021	0.004	0.059	0.372		91	0	0.017	0.004	0.059	0.256
33	S _{R0}	0.4	100	0	0.029	0.013	0.413	0.523		100	0	0.024	0.013	0.413	0.506
	S _{A1}	0.3	99	0	0.053	0.004	0.296	0.508		95	0	0.043	0.003	0.297	0.478
	S _{A2}	0.15	99.9	0	0.057	0.008	0.158	0.984		99.8	0.01	0.066	0.007	0.157	1.186
	S _B	0.2	99.9	0	0.086	0.010	0.210	0.868		99.8	0	0.077	0.009	0.209	1.027
	S _F	0.01	99.6	0.01	0.055	0.000	0.010	1.250		98	0.03	0.066	0.000	0.010	1.500
	Ψ _{A1}	0.6	100	0	0.043	0.000	0.600	0.770		100	0	0.035	0.001	0.599	0.735
	Ψ _{A2}	0.95	99	0	0.039	0.007	0.957	1.013		95	0	0.079	0.010	0.960	1.040
	P _{G1a}	0.5	100	0	0.139	0.000	0.500	1.000		100	0	0.142	0.000	0.500	1.000
	P _{G2b}	0.5	100	0	0.140	0.001	0.501	1.000		100	0	0.141	0.000	0.500	1.000
	S _A	0.043	99	0	0.018	0.002	0.045	0.277		95	0	0.020	0.002	0.045	0.253
	S _R	0.042	99	0	0.017	0.003	0.046	0.181		95	0	0.015	0.004	0.046	0.207
34	S _{R0}	0.4	100	0	0.029	0.013	0.413	0.529		100	0	0.024	0.013	0.413	0.504
	S _{A1}	0.3	99	0	0.053	0.005	0.295	0.518		94	0	0.043	0.003	0.297	0.478
	S _{A2}	0.15	99.9	0	0.056	0.007	0.157	0.922		100	0	0.064	0.009	0.159	0.913
	S _B	0.2	99.9	0.03	0.087	0.009	0.209	1.445		100	0.01	0.076	0.011	0.211	1.141
	S _F	0.05	99.8	0.1	0.128	0.004	0.054	1.670		98	0.5	0.166	0.005	0.055	3.000
	Ψ _{A1}	0.6	100	0	0.045	0.001	0.601	0.763		100	0	0.035	0.000	0.600	0.736
	Ψ _{A2}	0.95	99	0	0.047	0.006	0.956	1.013		95	0	0.079	0.010	0.960	1.035
	P _{G1a}	0.5	100	0	0.138	0.000	0.500	1.000		100	0	0.141	0.001	0.501	1.000
	P _{G2b}	0.5	100	0	0.138	0.001	0.501	1.000		100	0	0.139	0.001	0.499	1.000
	S _A	0.044	99	0	0.018	0.002	0.045	0.302		94	0	0.020	0.003	0.046	0.277
	S _R	0.042	99	0	0.017	0.003	0.046	0.317		94	0	0.015	0.004	0.046	0.214
35	S _{R0}	0.4	85	0	0.029	0.013	0.413	0.521		95	0	0.023	0.013	0.413	0.506
	S _{A1}	0.3	99	0	0.042	0.004	0.296	0.475		98	0	0.034	0.004	0.296	0.414
	S _{A2}	0.15	99	0.01	0.064	0.011	0.161	1.130		98	0	0.070	0.013	0.163	0.990
	S _B	0.2	92	3.9	0.366	0.005	0.205	4.529		96	2.3	0.318	0.019	0.219	3.765
	S _F	0.01	99	0.01	0.056	0.001	0.011	2.032		97	0.03	0.063	0.000	0.010	1.296
	Ψ _{A1}	0.98	90	0	0.211	0.051	0.929	0.994		97	0	0.146	0.024	0.956	0.996
	Ψ _{A2}	0.95	99	0	0.048	0.009	0.959	1.018		98	0	0.063	0.018	0.968	1.037
	P _{G1a}	0.5	100	0	0.169	0.004	0.504	1.000		99.9	0	0.194	0.002	0.498	1.000
	P _{G2b}	0.5	100	0	0.167	0.002	0.502	1.000		100	0	0.194	0.002	0.498	1.000
	S _A	0.043	99	0	0.020	0.003	0.046	0.355		95	0	0.021	0.004	0.047	0.287
	S _R	0.018	84	0	0.009	0.002	0.021	0.141		90	0	0.009	0.003	0.021	0.115

36	S _{R0}	0.4	85	0	0.029	0.013	0.413	0.521		95	0	0.023	0.013	0.413	0.502
	S _{A1}	0.3	99	0	0.042	0.004	0.296	0.476		98	0	0.034	0.003	0.297	0.440
	S _{A2}	0.15	99	0	0.062	0.011	0.161	0.916		98	0	0.070	0.012	0.162	0.833
	S _B	0.2	92	3.9	0.364	0.009	0.209	3.500		96	2.5	0.325	0.023	0.223	3.462
	S _F	0.05	99	0.04	0.123	0.004	0.054	2.000		97	0.2	0.143	0.004	0.054	2.000
	Ψ _{A1}	0.98	90	0	0.205	0.048	0.932	0.994		97	0	0.147	0.024	0.956	0.996
	Ψ _{A2}	0.95	99	0	0.046	0.009	0.959	1.027		98	0	0.067	0.018	0.968	1.034
	P _{G1a}	0.5	100	0	0.167	0.000	0.500	1.000		100	0	0.194	0.002	0.502	1.000
	P _{G2b}	0.5	100	0	0.167	0.001	0.499	1.000		100	0	0.194	0.000	0.500	1.000
	S _A	0.044	99	0	0.019	0.003	0.046	0.221		96	0	0.021	0.004	0.047	0.226
	S _R	0.019	84	0	0.009	0.002	0.021	0.116		91	0	0.009	0.002	0.021	0.120
37	S _{R0}	0.4	100	0	0.029	0.013	0.413	0.522		100	0	0.023	0.013	0.413	0.506
	S _{A1}	0.3	100	0	0.064	0.005	0.295	0.560		100	0	0.052	0.005	0.295	0.482
	S _{A2}	0.15	100	0	0.040	0.000	0.150	0.318		100	0	0.051	0.001	0.151	0.376
	S _B	0.2	100	0	0.046	0.000	0.200	0.403		100	0	0.037	0.000	0.200	0.365
	S _F	0.01	100	0	0.019	0.000	0.010	0.151		100	0	0.026	0.001	0.011	0.251
	Ψ _{A1}	0.4	100	0	0.043	0.001	0.401	0.571		100	0	0.036	0.000	0.400	0.533
	Ψ _{A2}	0.75	100	0	0.041	0.007	0.757	0.895		100	0	0.055	0.014	0.764	0.945
	P _{G1a}	0.9	100	0	0.060	0.001	0.901	1.000		100	0	0.057	0.000	0.900	1.000
	P _{G2b}	0.9	100	0	0.061	0.001	0.899	1.000		100	0	0.056	0.000	0.900	1.000
	S _A	0.034	100	0	0.012	0.000	0.034	0.108		100	0	0.013	0.000	0.035	0.108
	S _R	0.054	100	0	0.013	0.002	0.055	0.105		100	0	0.010	0.002	0.055	0.110
38	S _{R0}	0.4	100	0	0.029	0.013	0.413	0.541		100	0	0.023	0.013	0.413	0.501
	S _{A1}	0.3	100	0	0.064	0.006	0.294	0.584		100	0	0.052	0.004	0.296	0.493
	S _{A2}	0.15	100	0	0.040	0.000	0.150	0.343		100	0	0.051	0.001	0.149	0.352
	S _B	0.2	100	0	0.046	0.000	0.200	0.409		100	0	0.037	0.000	0.200	0.348
	S _F	0.05	100	0	0.042	0.000	0.050	0.306		100	0	0.055	0.000	0.050	0.303
	Ψ _{A1}	0.4	100	0	0.044	0.000	0.400	0.571		100	0	0.035	0.000	0.400	0.544
	Ψ _{A2}	0.75	100	0	0.042	0.007	0.757	0.892		100	0	0.054	0.015	0.765	0.958
	P _{G1a}	0.9	100	0	0.060	0.001	0.899	1.000		100	0	0.056	0.000	0.900	1.000
	P _{G2b}	0.9	100	0	0.060	0.000	0.900	1.000		100	0	0.056	0.000	0.900	1.000
	S _A	0.038	100	0	0.013	0.001	0.037	0.113		100	0	0.014	0.000	0.037	0.102
	S _R	0.054	100	0	0.012	0.002	0.056	0.107		100	0	0.010	0.002	0.056	0.100
39	S _{R0}	0.4	100	0	0.029	0.013	0.413	0.521		100	0	0.024	0.013	0.413	0.508
	S _{A1}	0.3	100	0	0.053	0.006	0.294	0.501		100	0	0.043	0.004	0.296	0.473
	S _{A2}	0.15	100	0	0.038	0.001	0.151	0.354		100	0	0.047	0.000	0.150	0.357

	S _B	0.2	100	0	0.057	0.001	0.201	0.434		100	0	0.046	0.000	0.200	0.374
	S _F	0.01	100	0	0.019	0.000	0.010	0.138		100	0	0.023	0.000	0.010	0.182
	Ψ _{A1}	0.6	100	0	0.044	0.001	0.599	0.778		100	0	0.036	0.000	0.600	0.729
	Ψ _{A2}	0.75	100	0	0.041	0.010	0.760	0.894		100	0	0.050	0.017	0.767	0.948
	P _{G1a}	0.9	100	0	0.066	0.001	0.899	1.000		100	0	0.065	0.001	0.599	1.000
	P _{G2b}	0.9	100	0	0.066	0.000	0.900	1.000		100	0	0.065	0.000	0.900	1.000
	S _A	0.034	100	0	0.011	0.000	0.034	0.084		100	0	0.012	0.000	0.035	0.100
	S _R	0.040	100	0	0.011	0.001	0.042	0.086		100	0	0.009	0.001	0.042	0.081
40	S _{R0}	0.4	100	0	0.029	0.013	0.413	0.528		100	0	0.023	0.013	0.413	0.514
	S _{A1}	0.3	100	0	0.053	0.004	0.296	0.503		100	0	0.042	0.004	0.296	0.457
	S _{A2}	0.15	100	0	0.038	0.000	0.150	0.329		100	0	0.047	0.001	0.149	0.356
	S _B	0.2	100	0	0.057	0.000	0.200	0.475		100	0	0.046	0.000	0.200	0.420
	S _F	0.05	100	0	0.041	0.000	0.050	0.267		100	0	0.050	0.000	0.050	0.336
	Ψ _{A1}	0.6	100	0	0.044	0.001	0.599	0.762		100	0	0.035	0.000	0.600	0.729
	Ψ _{A2}	0.75	100	0	0.040	0.010	0.760	0.895		100	0	0.050	0.017	0.767	0.929
	P _{G1a}	0.9	100	0	0.064	0.000	0.900	1.000		100	0	0.063	0.001	0.901	1.000
	P _{G2b}	0.9	100	0	0.064	0.001	0.901	1.000		100	0	0.064	0.001	0.901	1.000
	S _A	0.038	100	0	0.011	0.000	0.037	0.088		100	0	0.012	0.000	0.037	0.101
	S _R	0.041	100	0	0.011	0.001	0.042	0.103		100	0	0.009	0.001	0.042	0.081
41	S _{R0}	0.4	84	0	0.029	0.014	0.414	0.520		95	0	0.023	0.013	0.413	0.501
	S _{A1}	0.3	100	0	0.041	0.004	0.296	0.449		100	0	0.033	0.005	0.295	0.438
	S _{A2}	0.15	100	0	0.036	0.000	0.150	0.302		100	0	0.042	0.001	0.151	0.333
	S _B	0.2	92	0.02	0.290	0.005	0.205	1.333		98	0.03	0.235	0.001	0.201	1.200
	S _F	0.01	100	0	0.018	0.000	0.010	0.127		100	0	0.020	0.000	0.010	0.175
	Ψ _{A1}	0.98	89	0	0.216	0.053	0.927	0.994		97	0	0.147	0.024	0.956	0.996
	Ψ _{A2}	0.75	100	0	0.038	0.012	0.762	0.908		100	0	0.044	0.021	0.771	0.931
	P _{G1a}	0.9	100	0	0.082	0.001	0.899	1.000		100	0	0.094	0.000	0.900	1.000
	P _{G2b}	0.9	100	0	0.082	0.000	0.900	1.000		100	0	0.094	0.002	0.898	1.000
	S _A	0.035	100	0	0.010	0.000	0.035	0.085		100	0	0.011	0.000	0.035	0.088
	S _R	0.015	84	0	0.005	0.001	0.016	0.043		95	0	0.005	0.001	0.016	0.037
42	S _{R0}	0.4	85	0	0.029	0.013	0.413	0.535		94	0	0.023	0.013	0.413	0.517
	S _{A1}	0.3	100	0	0.041	0.005	0.295	0.456		100	0	0.034	0.005	0.295	0.429
	S _{A2}	0.15	100	0	0.036	0.000	0.150	0.306		100	0	0.042	0.000	0.150	0.343
	S _B	0.2	92	0.01	0.283	0.000	0.200	1.800		98	0.01	0.234	0.001	0.201	1.111
	S _F	0.05	100	0	0.039	0.000	0.050	0.311		100	0	0.044	0.000	0.050	0.404
	Ψ _{A1}	0.98	89	0	0.207	0.049	0.931	0.994		96	0	0.147	0.024	0.956	0.996

	Ψ_{A2}	0.75	100	0	0.038	0.012	0.762	0.892		100	0	0.044	0.021	0.771	0.920
	P_{G1a}	0.9	100	0	0.079	0.001	0.899	1.000		100	0	0.088	0.001	0.901	1.000
	P_{G2b}	0.9	100	0	0.079	0.000	0.900	1.000		100	0	0.089	0.001	0.901	1.000
	S_A	0.038	100	0	0.010	0.000	0.037	0.082		100	0	0.011	0.000	0.038	0.087
	S_R	0.016	85	0	0.005	0.001	0.017	0.043		94	0	0.005	0.001	0.017	0.039
43	S_{R0}	0.4	100	0	0.029	0.014	0.414	0.525		100	0	0.023	0.013	0.413	0.525
	S_{A1}	0.3	98	0	0.065	0.004	0.296	0.536		92	0	0.052	0.004	0.296	0.498
	S_{A2}	0.15	100	0	0.035	0.000	0.150	0.304		100	0	0.045	0.000	0.150	0.333
	S_B	0.2	100	0	0.046	0.001	0.199	0.412		100	0	0.037	0.000	0.200	0.346
	S_F	0.01	99.5	0	0.046	0.001	0.009	1.012		97	0	0.058	0.001	0.009	1.014
	Ψ_{A1}	0.4	100	0	0.044	0.000	0.400	0.584		100	0	0.035	0.000	0.400	0.543
	Ψ_{A2}	0.95	98	0	0.044	0.004	0.954	1.009		93	0	0.082	0.005	0.955	1.039
	P_{G1a}	0.9	100	0	0.057	0.001	0.899	1.000		100	0	0.056	0.000	0.900	1.000
	P_{G2b}	0.9	100	0	0.057	0.000	0.900	1.000		100	0	0.056	0.000	0.900	1.000
	S_A	0.043	98	0	0.014	0.000	0.043	0.129		92	0	0.015	0.000	0.043	0.120
	S_R	0.055	98	0	0.013	0.002	0.057	0.117		92	0	0.010	0.002	0.056	0.096
44	S_{R0}	0.4	100	0	0.029	0.013	0.413	0.526		100	0	0.023	0.013	0.413	0.508
	S_{A1}	0.3	98	0	0.065	0.004	0.296	0.584		92	0	0.053	0.003	0.297	0.492
	S_{A2}	0.15	100	0	0.035	0.000	0.150	0.287		100	0	0.045	0.001	0.151	0.324
	S_B	0.2	100	0	0.046	0.000	0.200	0.429		100	0	0.037	0.000	0.200	0.347
	S_F	0.05	99.6	0	0.106	0.000	0.050	1.026		97	0	0.140	0.002	0.052	1.073
	Ψ_{A1}	0.4	100	0	0.044	0.000	0.400	0.573		100	0	0.035	0.000	0.400	0.547
	Ψ_{A2}	0.95	98	0	0.039	0.005	0.955	1.015		92	0	0.081	0.005	0.955	1.034
	P_{G1a}	0.9	100	0	0.057	0.000	0.900	1.000		100	0	0.056	0.000	0.900	1.000
	P_{G2b}	0.9	100	0	0.057	0.001	0.901	1.000		100	0	0.055	0.001	0.899	1.000
	S_A	0.044	98	0	0.014	0.000	0.043	0.118		92	0	0.015	0.000	0.044	0.144
	S_R	0.055	98	0	0.013	0.002	0.057	0.113		92	0	0.010	0.002	0.057	0.103
45	S_{R0}	0.4	100	0	0.029	0.013	0.413	0.517		100	0	0.023	0.013	0.413	0.494
	S_{A1}	0.3	99	0	0.053	0.003	0.297	0.512		94	0	0.043	0.003	0.297	0.464
	S_{A2}	0.15	100	0	0.034	0.000	0.150	0.303		100	0	0.042	0.000	0.150	0.331
	S_B	0.2	100	0	0.056	0.000	0.200	0.439		100	0	0.045	0.000	0.200	0.382
	S_F	0.01	99.7	0	0.045	0.001	0.009	1.008		98	0	0.053	0.001	0.009	1.033
	Ψ_{A1}	0.6	100	0	0.044	0.000	0.600	0.777		100	0	0.035	0.000	0.600	0.730
	Ψ_{A2}	0.95	99	0	0.048	0.006	0.956	1.011		95	0	0.079	0.010	0.960	1.044
	P_{G1a}	0.9	100	0	0.061	0.001	0.899	1.000		100	0	0.062	0.000	0.900	1.000
	P_{G2b}	0.9	100	0	0.062	0.000	0.900	1.000		100	0	0.062	0.001	0.899	1.000

	S _A	0.043	99	0	0.013	0.000	0.043	0.096		94	0	0.014	0.000	0.043	0.108
	S _R	0.042	99	0	0.011	0.001	0.044	0.092		94	0	0.009	0.001	0.044	0.083
46	S _{R0}	0.4	100	0	0.029	0.013	0.413	0.517		100	0	0.023	0.012	0.412	0.509
	S _{A1}	0.3	99	0	0.053	0.002	0.298	0.520		95	0	0.043	0.003	0.297	0.475
	S _{A2}	0.15	100	0	0.035	0.001	0.151	0.317		100	0	0.041	0.001	0.151	0.326
	S _B	0.2	100	0	0.057	0.001	0.199	0.447		100	0	0.045	0.000	0.200	0.371
	S _F	0.05	99.8	0	0.099	0.001	0.049	1.064		98	0	0.125	0.001	0.049	1.051
	Ψ _{A1}	0.6	100	0	0.044	0.002	0.598	0.763		100	0	0.036	0.000	0.600	0.728
	Ψ _{A2}	0.95	99	0	0.039	0.007	0.957	1.013		95	0	0.074	0.011	0.961	1.035
	P _{G1a}	0.9	100	0	0.061	0.002	0.898	1.000		100	0	0.061	0.001	0.901	1.000
	P _{G2b}	0.9	100	0	0.061	0.001	0.899	1.000		100	0	0.061	0.000	0.900	1.000
	S _A	0.044	99	0	0.013	0.000	0.044	0.103		95	0	0.014	0.000	0.044	0.112
	S _R	0.042	99	0	0.011	0.001	0.044	0.090		95	0	0.009	0.001	0.044	0.080
47	S _{R0}	0.4	85	0	0.029	0.014	0.414	0.527		95	0	0.023	0.013	0.413	0.513
	S _{A1}	0.3	99	0	0.041	0.004	0.296	0.443		98	0	0.034	0.003	0.297	0.453
	S _{A2}	0.15	100	0	0.032	0.000	0.150	0.309		100	0	0.037	0.000	0.150	0.292
	S _B	0.2	93	0.01	0.284	0.007	0.207	1.122		98	0	0.233	0.001	0.201	1.083
	S _F	0.01	99.9	0	0.043	0.000	0.010	0.513		99	0	0.050	0.000	0.010	1.048
	Ψ _{A1}	0.98	90	0	0.210	0.051	0.929	0.994		97	0	0.136	0.021	0.959	0.996
	Ψ _{A2}	0.95	99	0	0.037	0.010	0.960	1.021		98	0	0.058	0.019	0.969	1.041
	P _{G1a}	0.9	100	0	0.074	0.000	0.900	1.000		100	0	0.084	0.001	0.899	1.000
	P _{G2b}	0.9	100	0	0.073	0.000	0.900	1.000		100	0	0.084	0.000	0.900	1.000
	S _A	0.043	99	0	0.011	0.000	0.043	0.088		98	0	0.012	0.000	0.043	0.098
	S _R	0.018	85	0	0.005	0.001	0.019	0.042		93	0	0.005	0.001	0.019	0.043
48	S _{R0}	0.4	85	0	0.029	0.014	0.414	0.532		95	0	0.023	0.014	0.414	0.502
	S _{A1}	0.3	99	0	0.041	0.004	0.296	0.465		97	0	0.034	0.004	0.296	0.429
	S _{A2}	0.15	100	0	0.032	0.000	0.150	0.283		100	0	0.037	10.000	0.150	0.332
	S _B	0.2	92	0	0.282	0.000	0.200	1.096		98	0	0.233	0.002	0.202	1.100
	S _F	0.05	99.9	0	0.093	0.001	0.051	1.011		99	0	0.107	0.001	0.049	1.032
	Ψ _{A1}	0.98	89	0	0.211	0.051	0.929	0.994		97	0	0.145	0.024	0.956	0.996
	Ψ _{A2}	0.95	99.5	0	0.047	0.009	0.959	1.021		98	0	0.061	0.019	0.969	1.034
	P _{G1a}	0.9	100	0	0.073	0.001	0.901	1.000		100	0	0.084	0.000	0.900	1.000
	P _{G2b}	0.9	100	0	0.072	0.000	0.900	1.000		100	0	0.084	0.001	0.899	1.000
	S _A	0.044	99	0	0.011	0.000	0.043	0.097		97	0	0.012	0.000	0.044	0.100
	S _R	0.019	84	0	0.005	0.001	0.019	0.050		92	0	0.005	0.001	0.019	0.043