TIM Diagnostics

MODEL STRUCTURE

Match/Mismatch: Matching

Population Structure: Metapopulation

Diagnostic Run: NO. Uses OBS values as data inputs

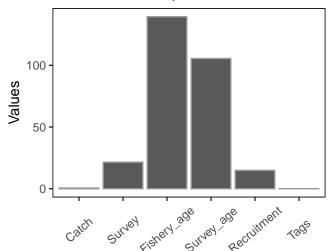
Residual plots represent RELATIVE difference; ((TRUE-ESTIMATED)/TRUE)*100

Run Time of Estimation: 0.3345 minutes

OM Error Parameters

	Parameter	1	2
1	Sigma_Rec	0.4	0.4
2	Sigma_Rec_Apport	0.2	0.2
3	Sigma_F	0.2	0.2
4	Rec_Index_sigma	0.5	0.5
5	Survey_Sigma	0.2	0.2
6	Catch_Sigma	0.05	0.05
7	SIM_N_Catch	100	100
8	SIM_N_Survey	100	100

Likelihood Components



Model Converged

num gradient component = 4.67585321861597€

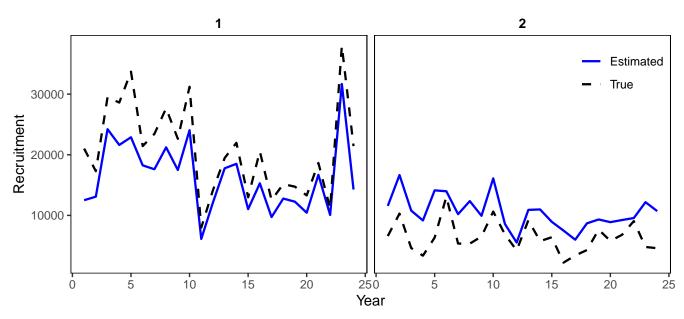
Estimated Parameter Values

	Parameter	1	2
1	q	0.00	0.00
2	R_ave	16745.30	10936.80
3	beta 1 fishery	4.86	5.22
4	beta 2 fishery	1.69	1.83
5	beta 3 fishery	NA	NA
6	beta 4 fishery	NA	NA
7	beta 1 survey	3.85	3.29
8	beta 2 survey	1.67	1.99
9	beta 3 survey	NA	NA
10	beta 4 survey	NA	NA

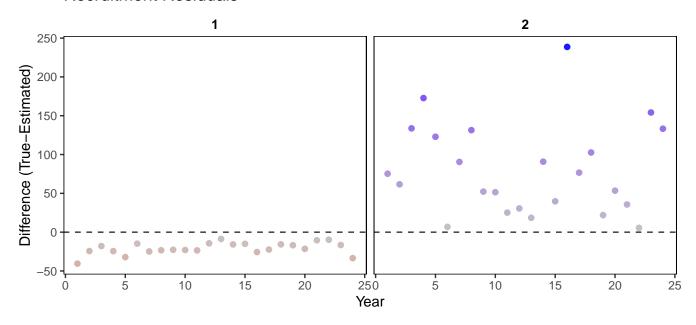
True Parameter Values

Parameter	1	2
q	0.00	0.00
R_ave	21019.00	6596.00
beta 1 fishery	1.31	2.45
beta 2 fishery	3.76	3.16
beta 3 fishery	NA	NA
beta 4 fishery	NA	NA
beta 1 survey	2.51	1.81
beta 2 survey	2.31	2.64
beta 3 survey	NA	NA
beta 4 survey	NA	NA
	q R_ave beta 1 fishery beta 2 fishery beta 3 fishery beta 4 fishery beta 1 survey	q 0.00 R_ave 21019.00 beta 1 fishery 1.31 beta 2 fishery 3.76 beta 3 fishery NA beta 4 fishery NA beta 1 survey 2.51 beta 2 survey 2.31 beta 3 survey NA

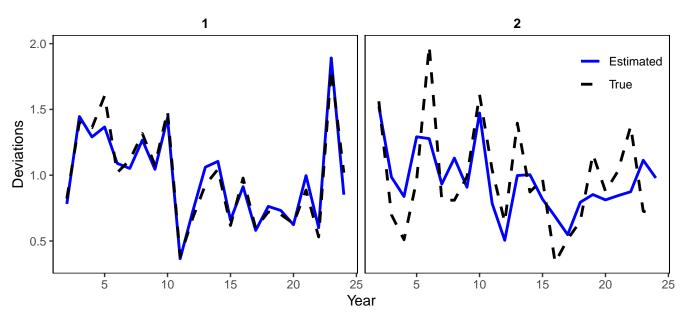
Recruitment Total



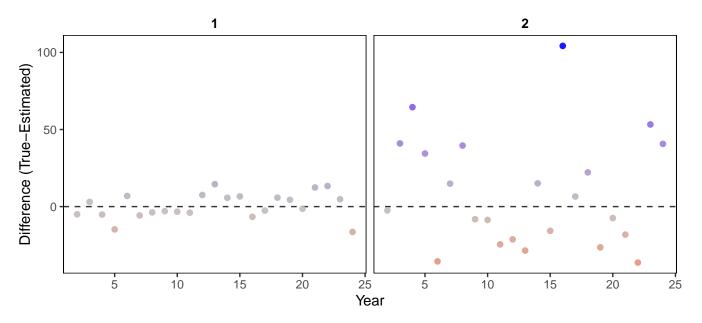
Recruitment Residuals



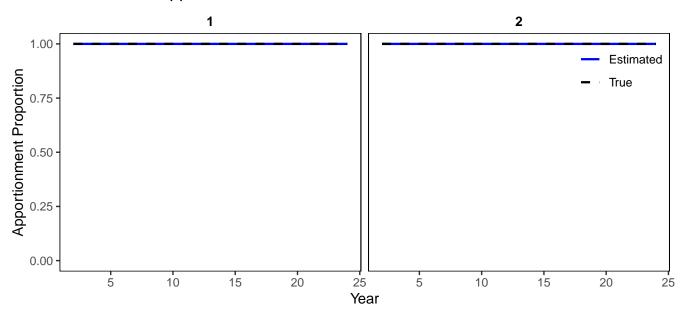
Recruitment Deviations



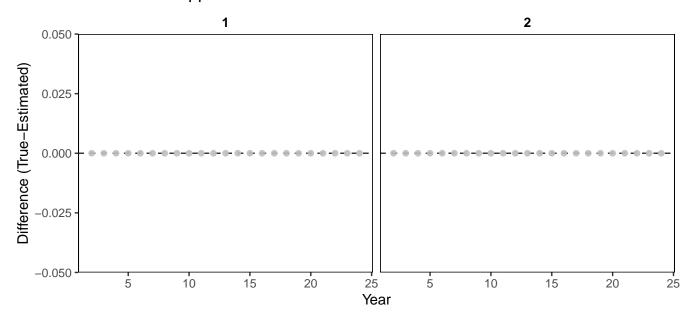
Recruitment Devs Residuals



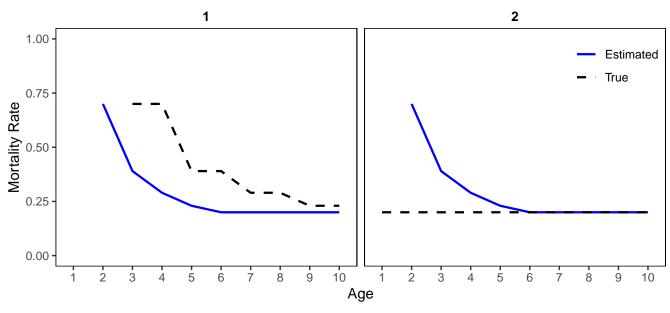
Recruitment Apportionment



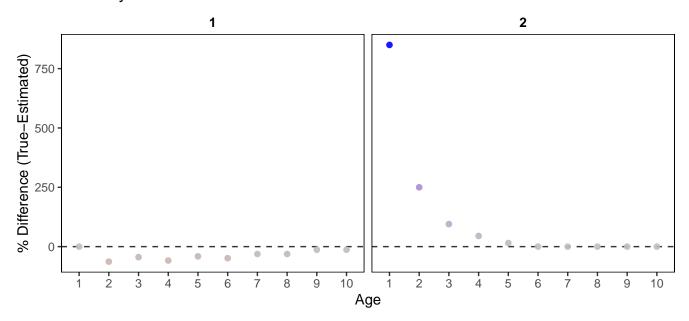
Recruitment Apportionment Residuals



Mortality Rate



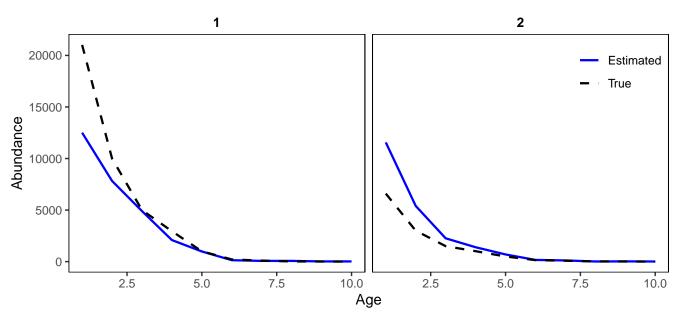
Mortality Rate Residuals



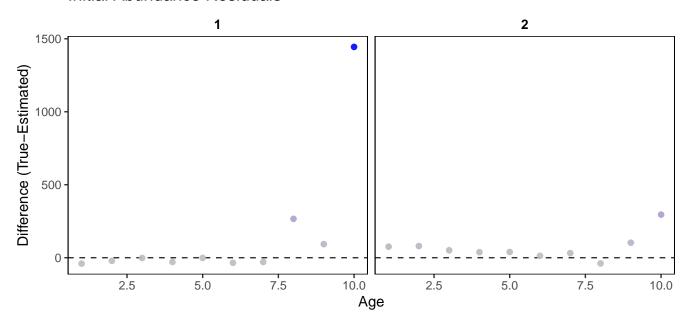




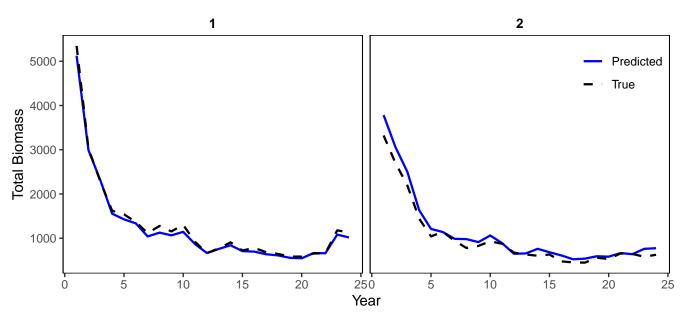
Initial Abundance



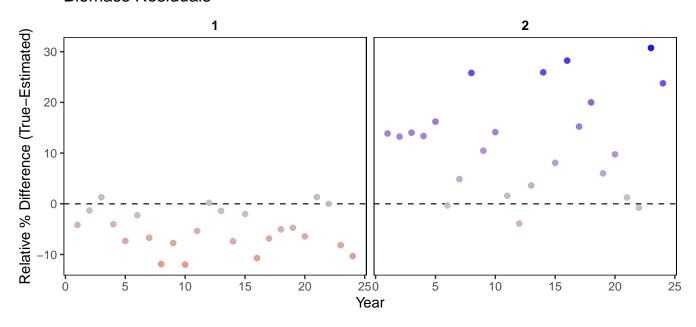
Initial Abundance Residuals

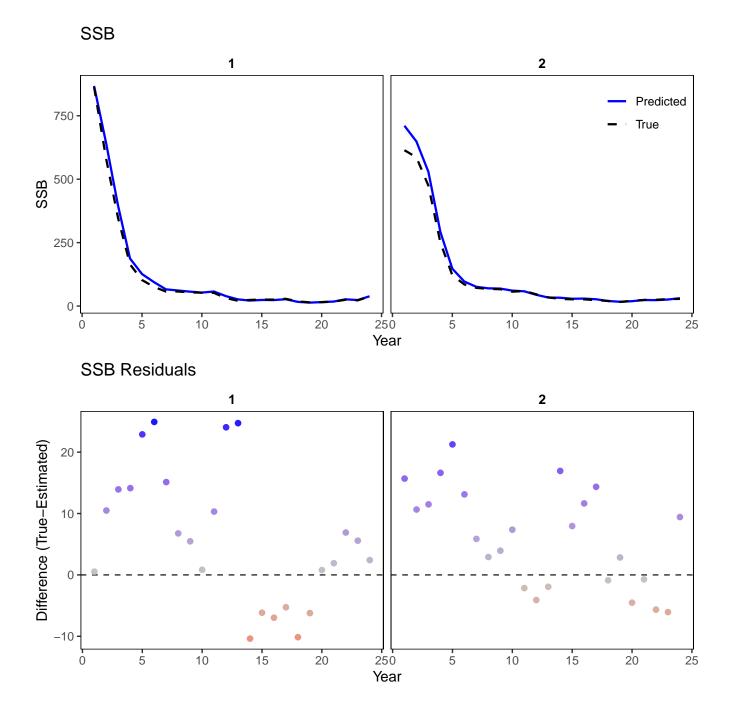


Total Biomass

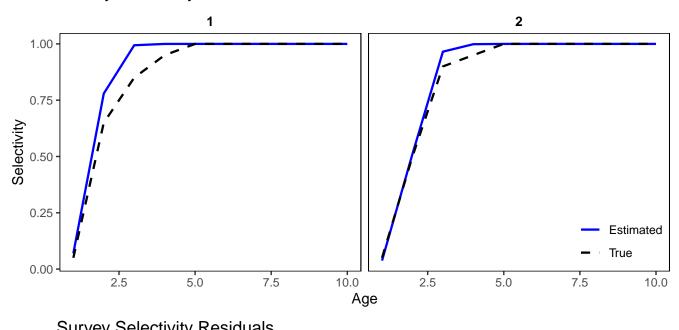


Biomass Residuals

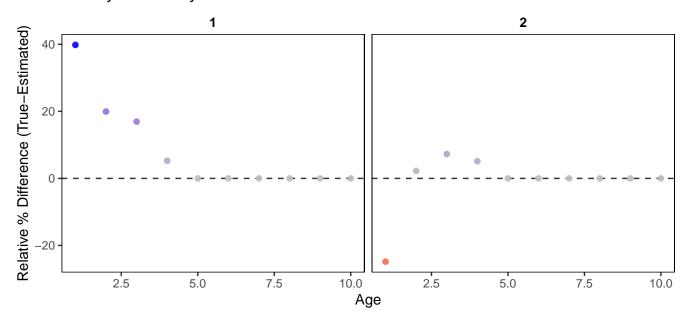




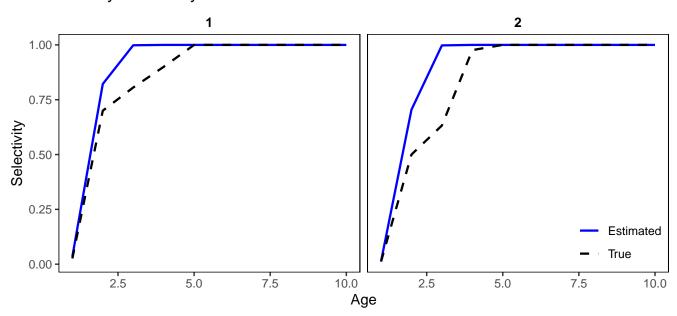
Survey Selectivity



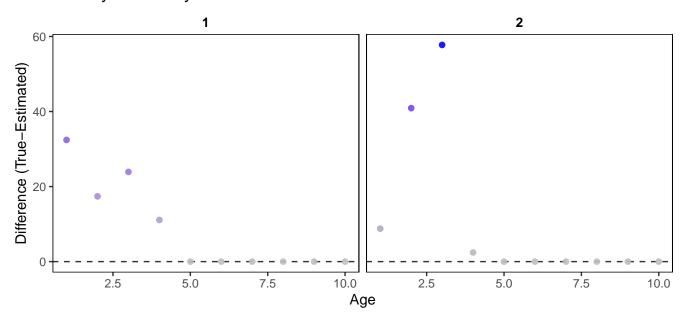
Survey Selectivity Residuals



Fishery Selectivity

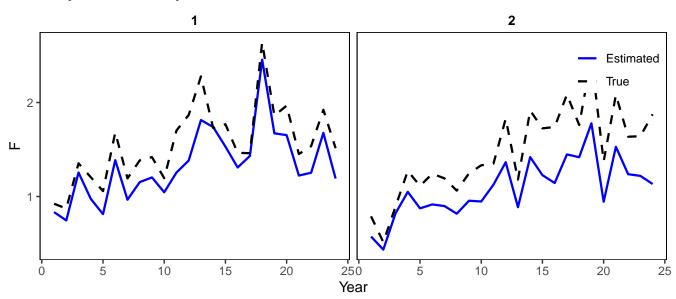


Fishery Selectivity Residuals

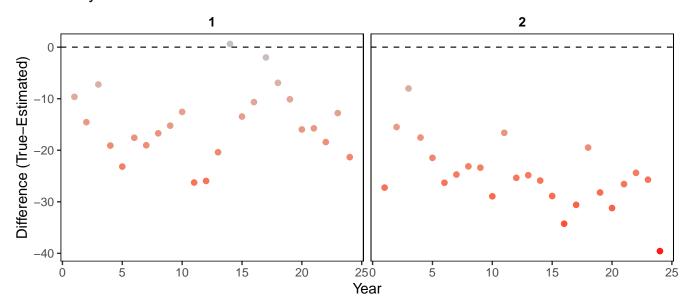


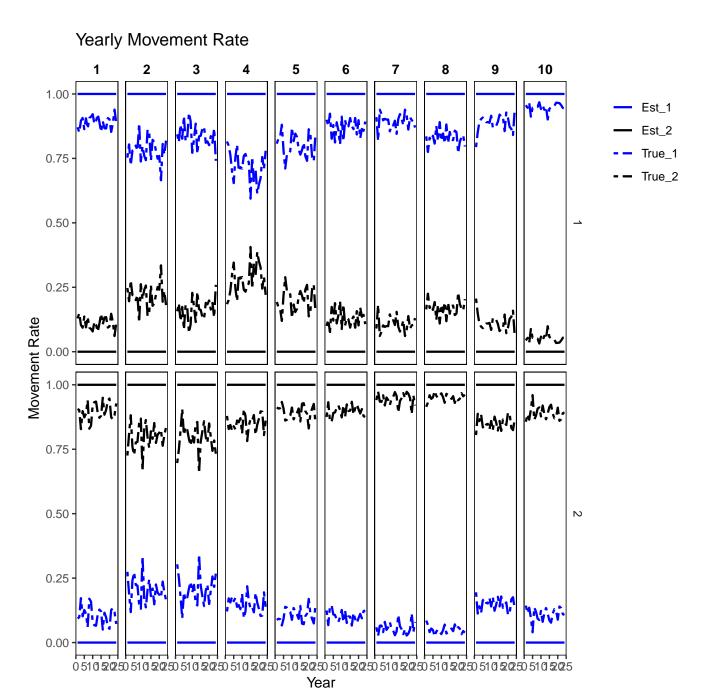


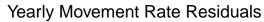


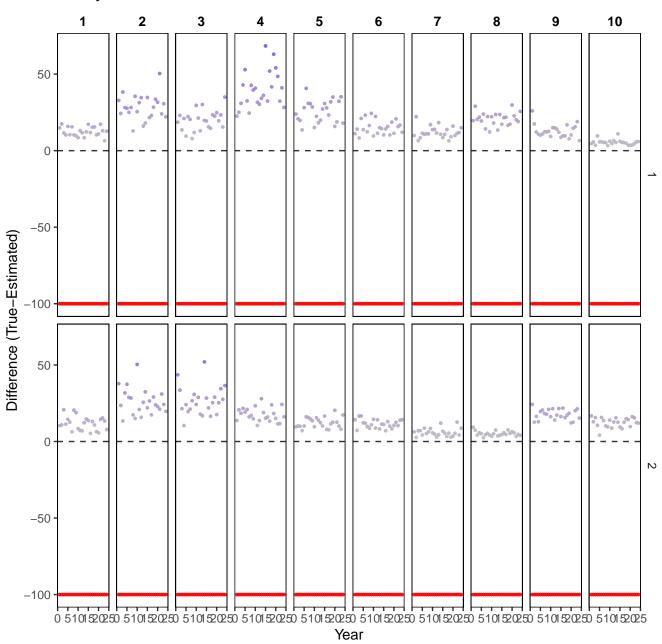


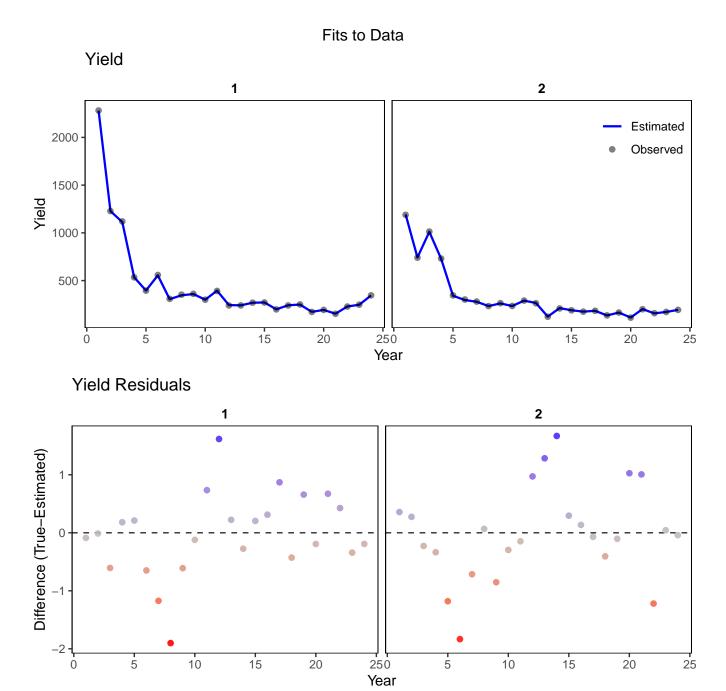
Fully Selected F Residuals





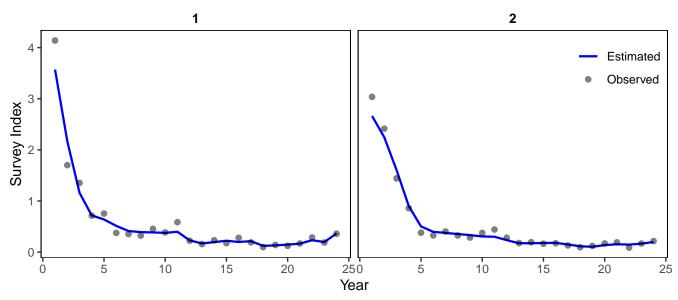




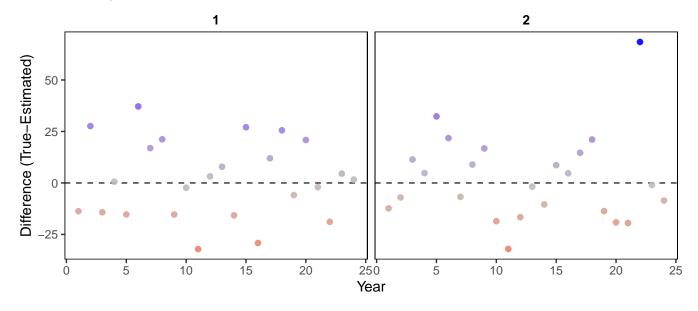


Fits to Data continued





Survey Index Residuals



Fits to Data continued

