# TIM Diagnostics

#### MODEL STRUCTURE

Match/Mismatch: Matching

Population Structure: Metapopulation

Diagnostic Run: NO. Uses OBS values as data inputs

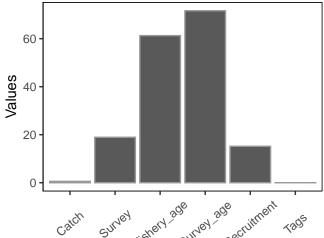
Residual plots represent RELATIVE difference; ((TRUE-ESTIMATED)/TRUE)\*100

Run Time of Estimation: 0.23667 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 = 2.86611874602967€

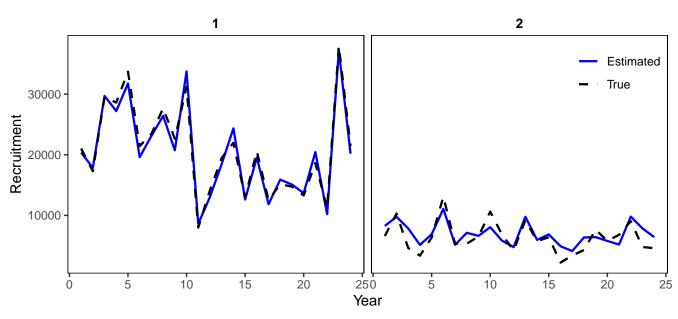
#### **Estimated Parameter Values**

	Parameter	1	2
1	q	0.00	0.00
2	R_ave	20692.40	7190.84
3	beta 1 fishery	4.87	5.37
4	beta 2 fishery	1.74	1.84
5	beta 3 fishery	NA	NA
6	beta 4 fishery	NA	NA
7	beta 1 survey	4.41	5.53
8	beta 2 survey	1.66	1.55
9	beta 3 survey	NA	NA
10	beta 4 survey	NA	NA

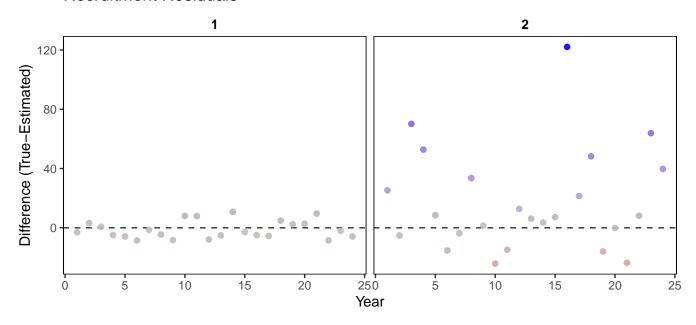
True Parameter Values

	Parameter	1	2
1	q	0.00	0.00
2	R_ave	21019.00	6596.00
3	beta 1 fishery	1.31	2.45
4	beta 2 fishery	3.76	3.16
5	beta 3 fishery	NA	NA
6	beta 4 fishery	NA	NA
7	beta 1 survey	2.51	1.81
8	beta 2 survey	2.31	2.64
9	beta 3 survey	NA	NA
10	beta 4 survey	NA	NA

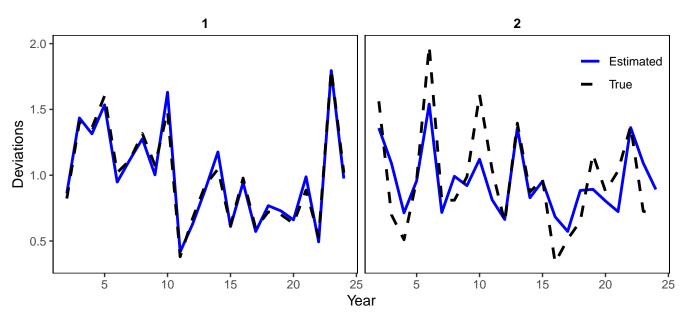
#### Recruitment Total



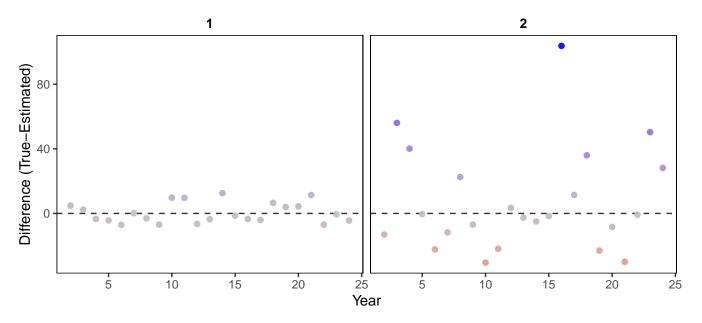
## Recruitment Residuals



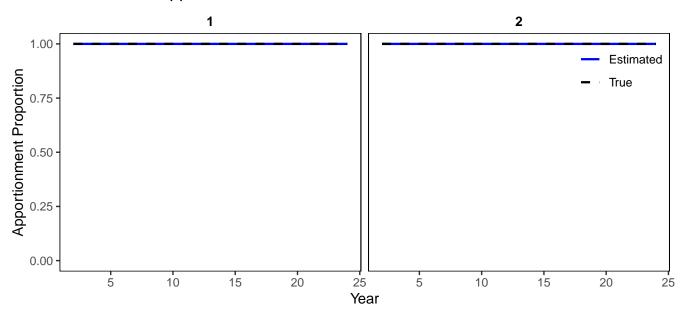
#### **Recruitment Deviations**



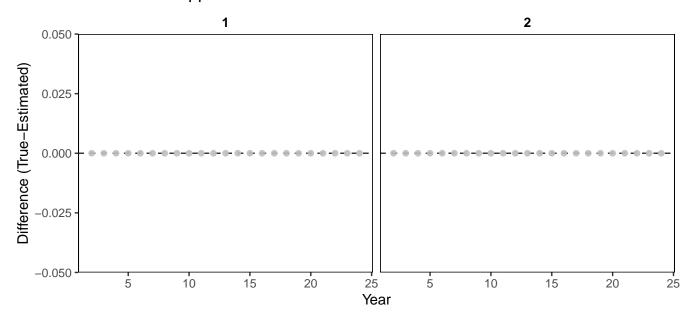
# Recruitment Devs Residuals



## **Recruitment Apportionment**



## Recruitment Apportionment Residuals







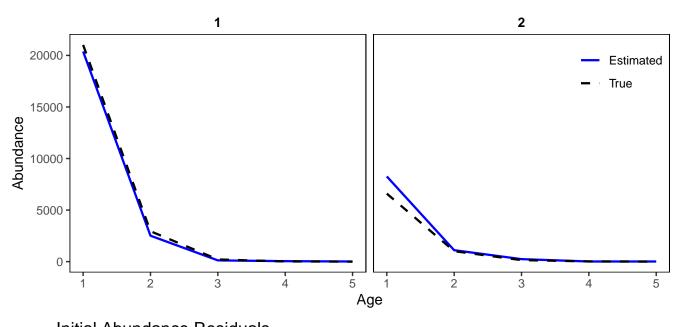
# Mortality Rate Residuals



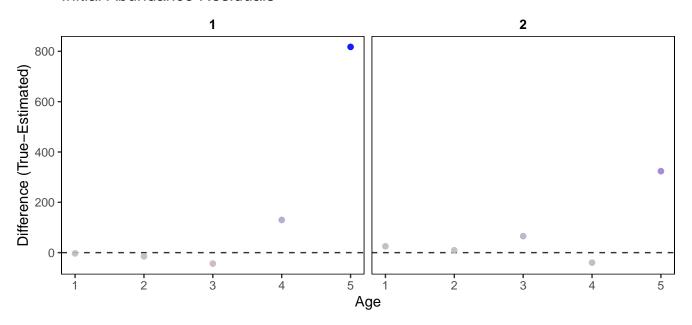




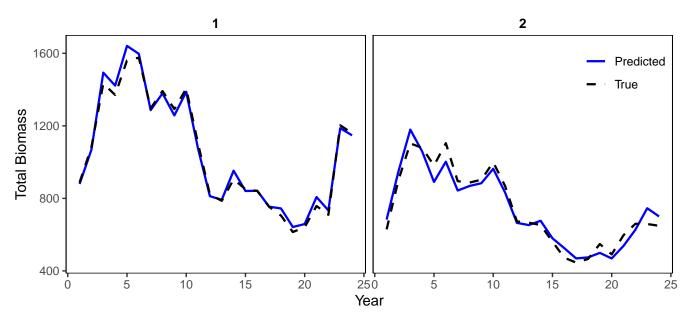




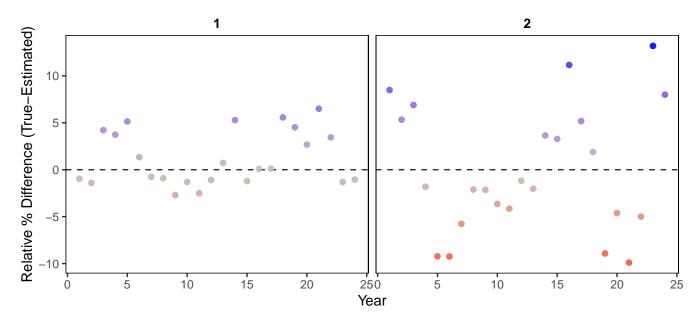
#### Initial Abundance Residuals

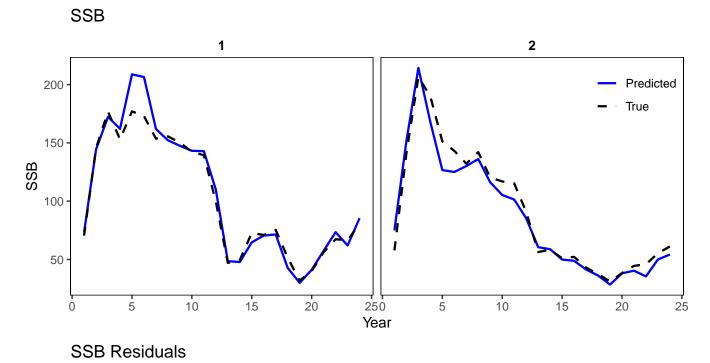


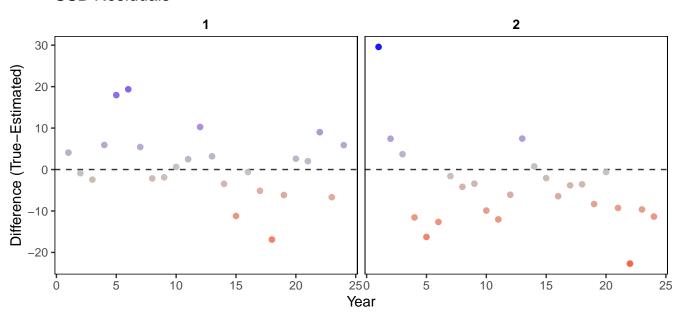
#### **Total Biomass**



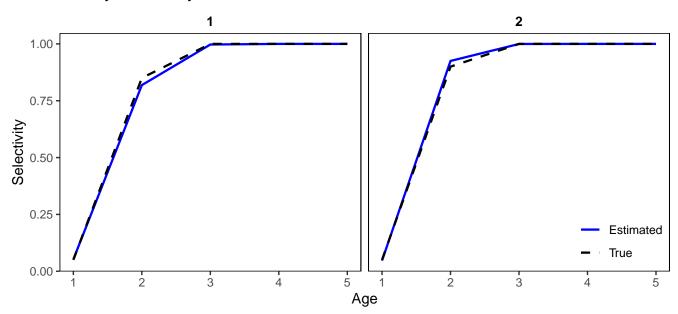
## Biomass Residuals



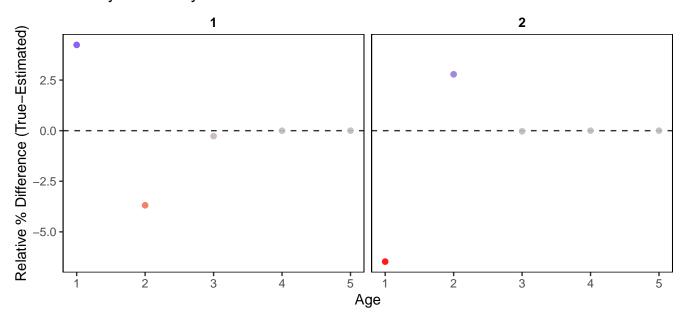


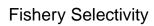


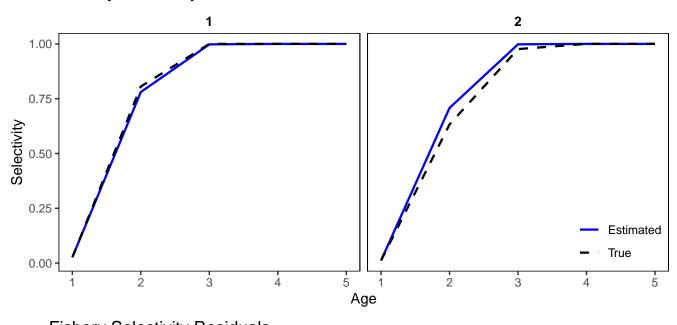




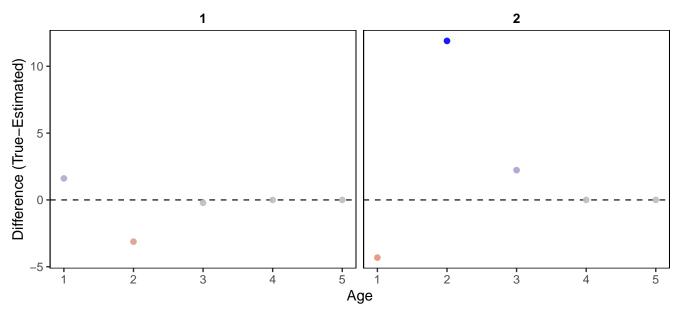
# Survey Selectivity Residuals





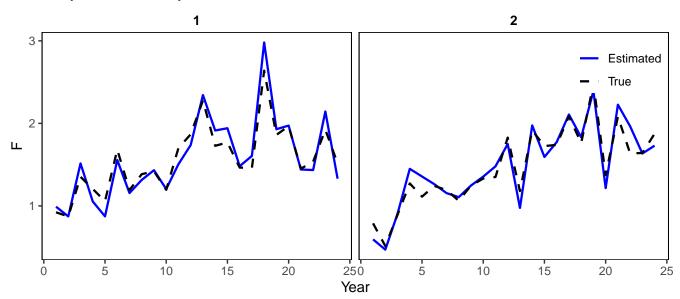


# Fishery Selectivity Residuals

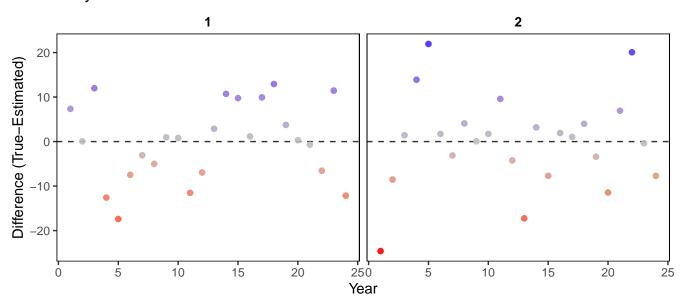


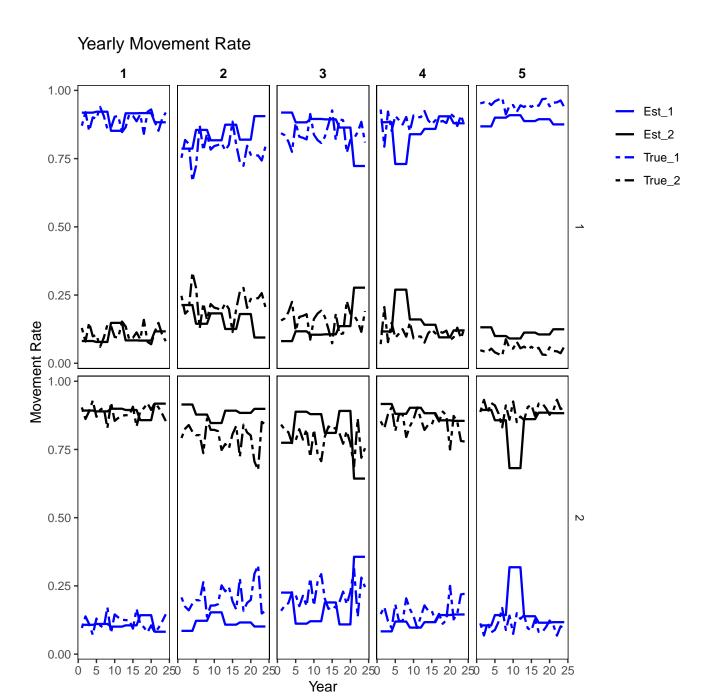
Fishing Mortality



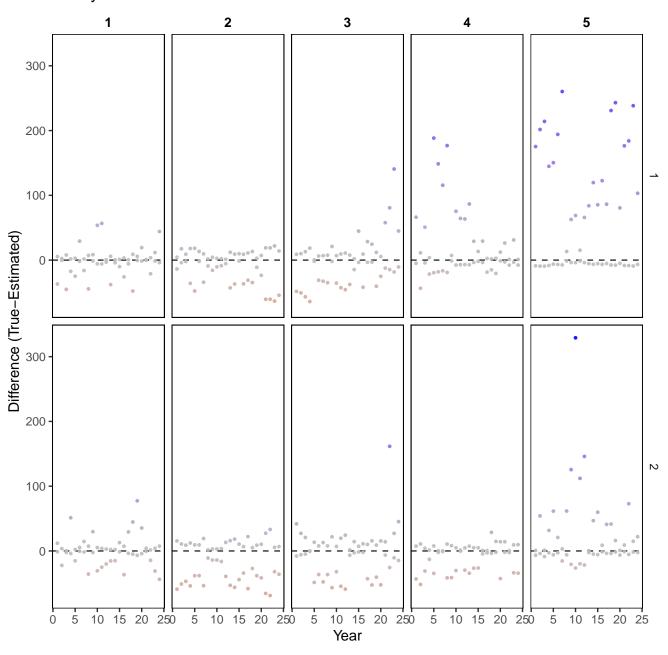


Fully Selected F Residuals

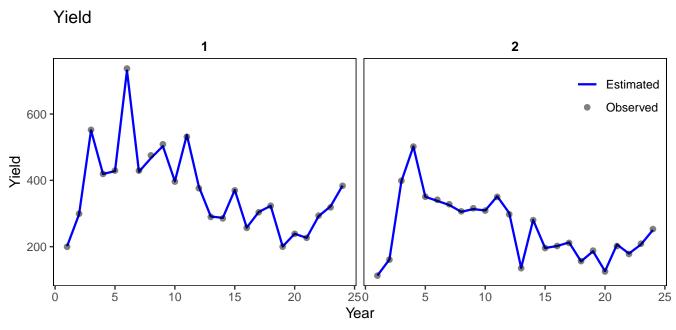




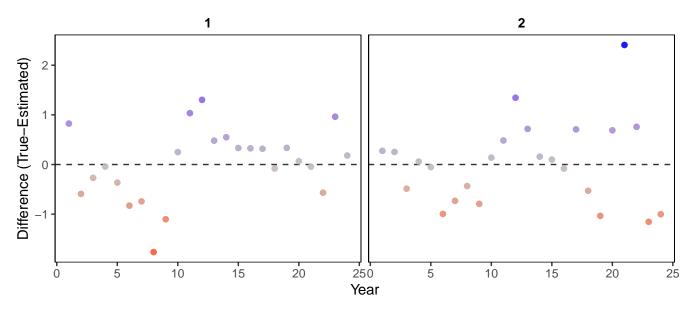
## Yearly Movement Rate Residuals





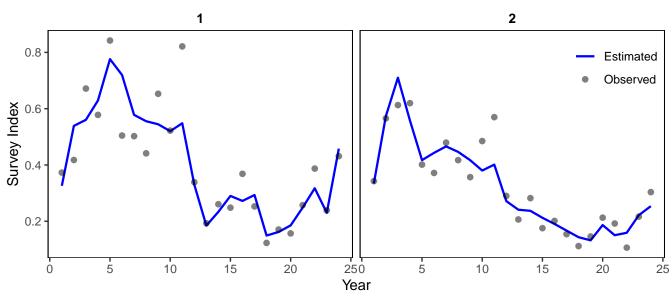


# Yield Residuals

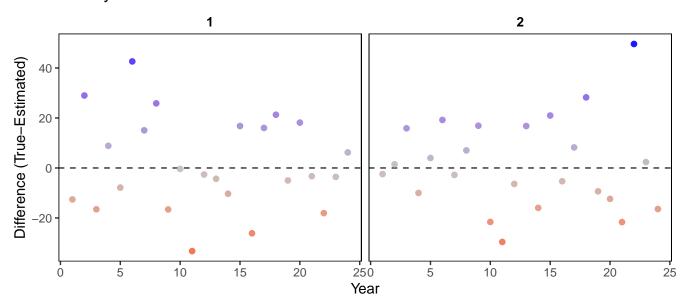


Fits to Data continued





# Survey Index Residuals



Fits to Data continued

