IBC Model Verification

Sensitivity TXT Input

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Results: Mortality



1 ggplot(alltrtw1%>% filter(EP=="Mortality" & HT == "Fixed"),aes(x=Time,y=Mean))+annotat

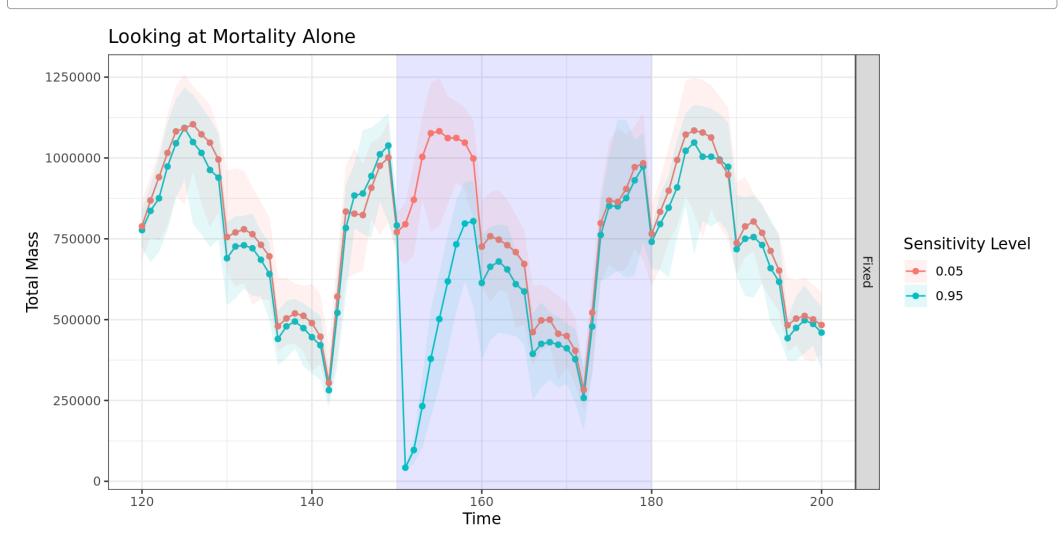


Figure 1: Total Mass Response with Mortality Effect

Results: Effect parameters



1 ggplot(alltrtw1%>% filter(HT == "Fixed"), aes(x=Time, y=Mean))+annotate("rect", xmin = 5

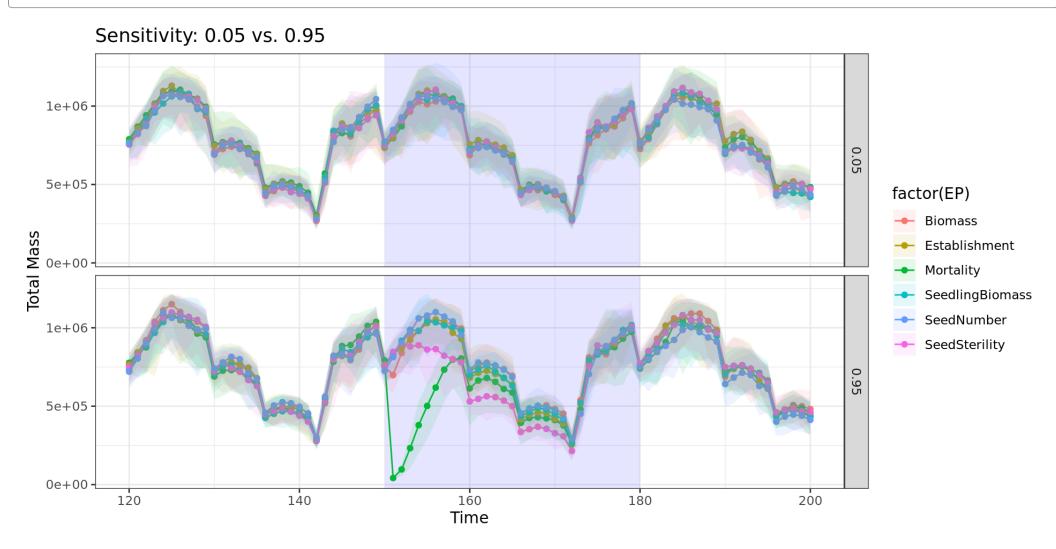


Figure 2: Total Mass Response with different Effects

Looking at each EP



1 ggplot(alltrtw1%>% filter(HT == "Fixed"), aes(x=Time, y=Mean))+annotate("rect", xmin = 5

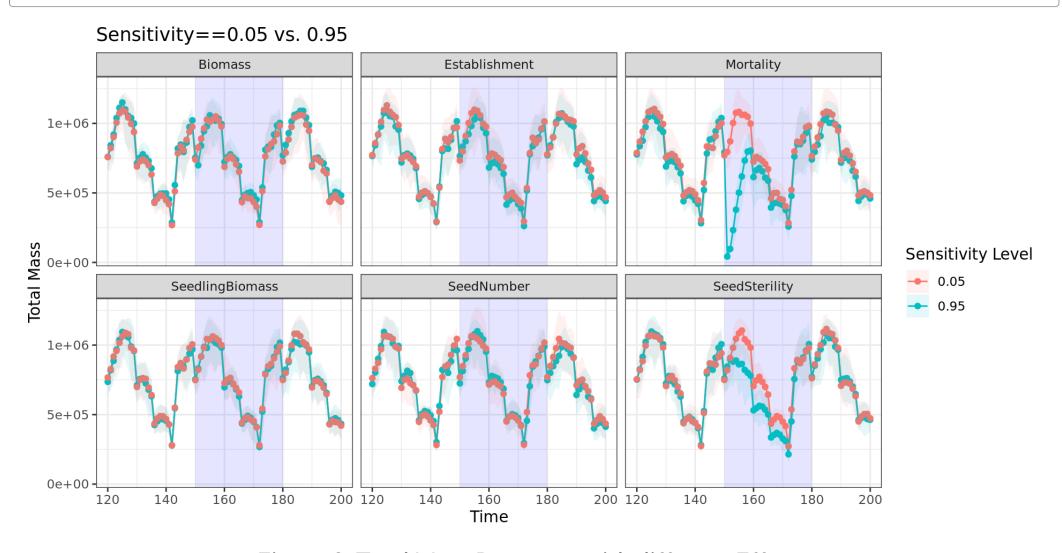


Figure 3: Total Mass Response with different Effects

Above Mass



1 ggplot(alltrtw1%>% filter(HT == "Fixed"), aes(x=Time, y=Mean))+annotate("rect", xmin = 5

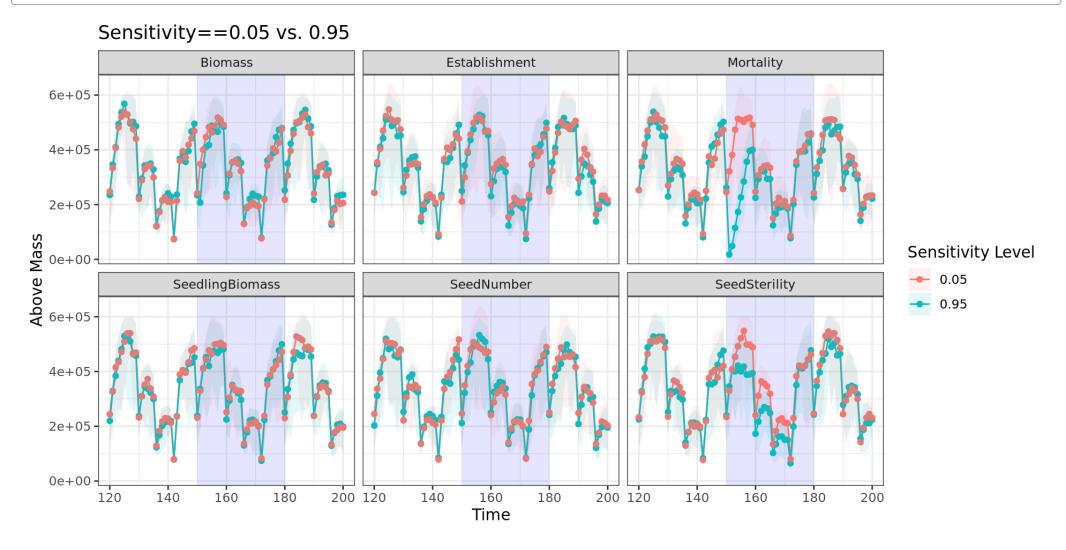


Figure 4: Total Mass Response with different Effects

Below Mass



1 ggplot(alltrtw1%>% filter(HT == "Fixed"), aes(x=Time, y=Mean))+annotate("rect", xmin = 5

Sensitivity: 0.05 vs. 0.95 Establishment Biomass Mortality 6e+05 4e+05 2e+05 Below Mass 6e+90 Sensitivity Level **0.05** SeedlingBiomass SeedNumber SeedSterility **0**.95 4e+05 2e+05 0e+00-160 160 200 120 140 140 180 200 120 180 140 160 180 200 120 Time

Figure 5: Above Mass Response with different Effects

Shannon



1 ggplot(alltrtw1%>% filter(HT == "Fixed"), aes(x=Time, y=Mean))+annotate("rect", xmin = 5

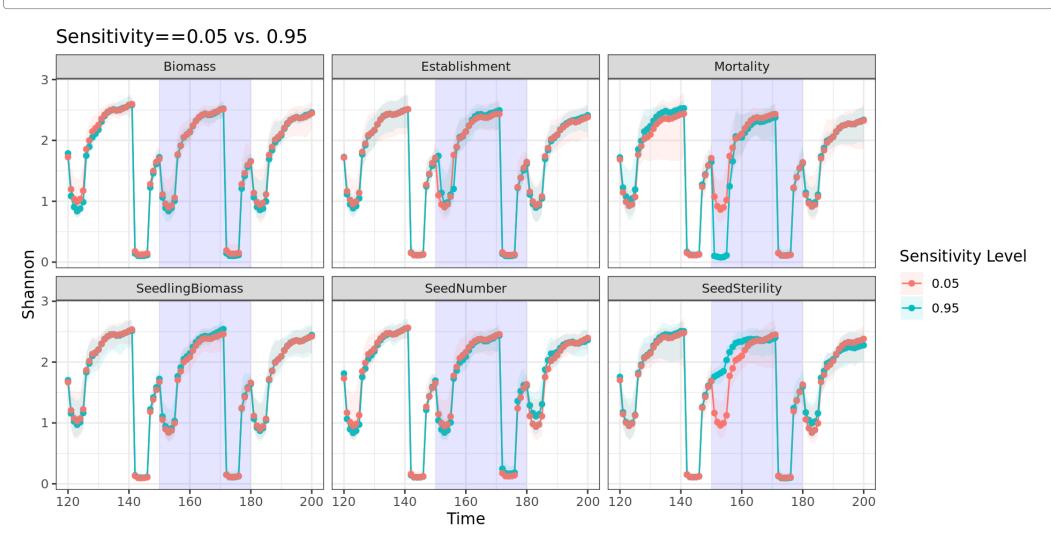


Figure 6: Shannon Response with different Effects





1 ggplot(alltrtw1%>% filter(HT == "Fixed"), aes(x=Time, y=Mean))+annotate("rect", xmin = 5

Sensitivity: 0.05 vs. 0.95

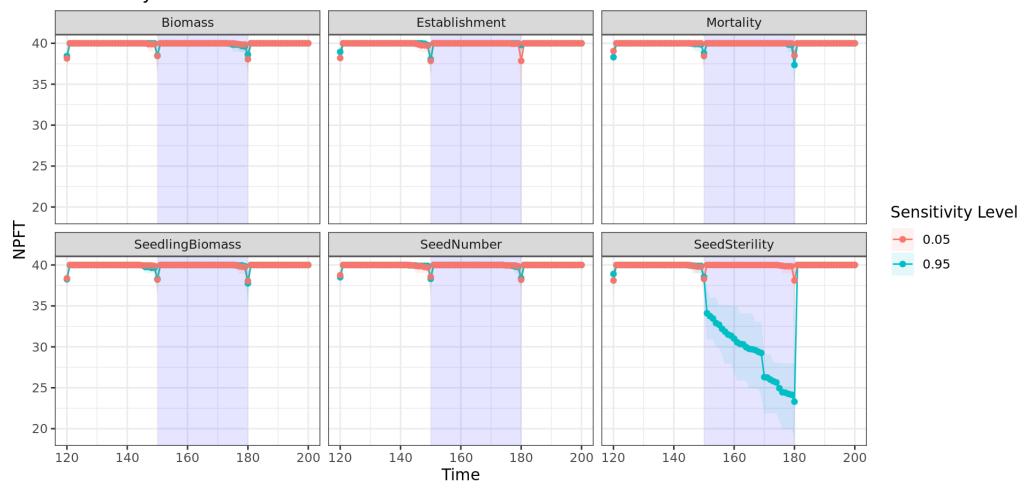


Figure 7: NPFT Response with different Effects

Issues



