

# Spatiotemporal model-based index development for Bering Sea and Aleutian Islands crab stocks

Update for Crab Plan Team modeling workshop

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## Introduction

The goal of this investigation was to develop spatiotemporal model-based indices of abundance for three Bering Sea and Aleutian Islands (BSAI) crab stocks: Tanner crab (*Chionoecetes bairdi*), Norton Sound red king crab (*Paralithodes camtschaticus*), and St. Matthew Island blue king crab (*Paralithodes platypus*). Research suggests that spatiotemporal model-based indices can be more robust to survey changes than are design-based indices, though the models must be well-specified (Yalcin et al. 2023). Spatiotemporal model-based indices are used in North Pacific Fishery Management Council (NPFMC) groundfish stock assessments for species including Eastern Bering Sea (EBS) walleye pollock (*Gadus chalcogrammus*) and EBS Pacific cod (*Gadus macrocephalus*), both of which use the vector-autoregressive spatial temporal (VAST) approach (Thorson 2019) to produce indices used in the assessments (Ianelli et al. 2024; Barbeaux et al. 2024). Previous BSAI crab stock assessments have presented models using spatiotemporal model-based indices (e.g., Ianelli et al. 2017), although these models were not accepted for harvest specifications (SSC 2017).

We generated biomass and abundance estimates using the R package *sdmTMB* (Anderson et al. 2022), which uses geostatistical time series data to estimate spatial and spatiotemporal generalized linear mixed effects models. This approach allows for index standardization when the set of stations surveyed is not consistent across years: one can generate a spatial grid that covers the area of interest, predict from the model onto that grid, and sum the predicted biomass to obtain an area-weighted biomass index that is independent of sampling locations (Anderson et al. 2022).

All three stock assessments for the crab stocks presented here use data from the National Marine Fisheries Service (NMFS) EBS bottom trawl survey (Stockhausen 2024; Hamazaki 2024; Stern and Palof 2024). The St. Matthew Island blue king crab stock assessment also uses data from the Alaska Department of Fish and Game (ADF&G) St. Matthew Island blue king crab pot survey, while the Norton Sound red king crab stock assessment uses data from the NMFS Northern Bering Sea bottom trawl survey and the ADF&G Norton Sound red king crab trawl survey.

Spatiotemporal model-based index development is expected to confer distinct advantages for each of the three stocks. For the St. Matthew Island blue king crab stock, standardizing the survey indices could allow the assessment to use the existing survey data more rigorously. The NMFS EBS trawl survey is undergoing changes including dropping the high sampling density “corner stations” near St. Matthew Island from 2024

onward (DePhilippo et al. 2023; Stern & Palof 2024); index standardization will allow the assessment to continue using the full time series of data despite changes in the spatial footprint of the survey. For Norton Sound red king crab, a model-based approach could provide a more consistent way to combine the three existing trawl survey data sets into a single index of abundance.

## Methods

We fit models using the R package *sdmTMB*. After fitting a model, we used the `sanity()` command to check whether the model converged, the Hessian matrix was positive definite, and any extreme eigenvalues were detected, among other checks. For models that passed the `sanity()` checks, we used the R package *DHARMA* (Hartig 2022) for model diagnostics. After calculating the DHARMA residuals using the function `DHARMA::dharma_residuals()`, we tested for quantile deviations, under/overdispersion, outliers, and zero inflation using the functions `DHARMA::testQuantiles()`, `DHARMA::testDispersion()`, `DHARMA::testOutliers()`, and `DHARMA::testZeroInflation()`, respectively.

A number of decision points arise when fitting models using *sdmTMB*, including: - the resolution of the spatial mesh used in fitting the model. A higher number of knots, specified when creating the spatial mesh using the `make_mesh()` function, indicates a higher resolution mesh. Few guidelines exist to aid in selection of an appropriate mesh resolution for a given dataset. - the spatiotemporal random fields estimation method. The spatiotemporal random fields can be estimated as independent and identically distributed (IID), first-order autoregressive (AR1), a random walk, or fixed at zero. - the model family. Many options exist, including `tweedie()`, `delta_gamma()`, and `delta_lognormal()`.

Further decision points arise when evaluating fitted models: - the relative importance of different model diagnostics in selecting a preferred model. - the importance of visual model fit to survey observations in selecting a preferred model.

For each stock, we present a range of models to show the effects of the decision points that arise when fitting models, and discuss the evaluation of fitted models.

### Tanner crab

#### Norton Sound red king crab

We combined data from the NMFS trawl survey (1976-1991), ADF&G trawl survey (1996-2024), and NMFS NBS trawl survey (2010-2023) into a single data set to which we fit models in *sdmTMB*. We filtered the data set to ensure that it included only observations with coordinates falling within the Norton Sound Section of Statistical Area Q. For model fitting, we used spatial meshes at three resolutions, specified in terms of the number of knots (vertices): 100 knots, 50 knots, and 30 knots (Figures 11 - 9). We used a prediction grid with resolution of 5 km<sup>2</sup> (Figure 8).

#### St. Matthew Island blue king crab

For model fitting, we used spatial meshes at three resolutions, specified in terms of the number of knots (vertices): 120 knots, 90 knots, and 50 knots (Figures 22 - 20). We used a prediction grid with resolution of 4 km<sup>2</sup> (Figure 23).

## EBS Tanner estimated abundance

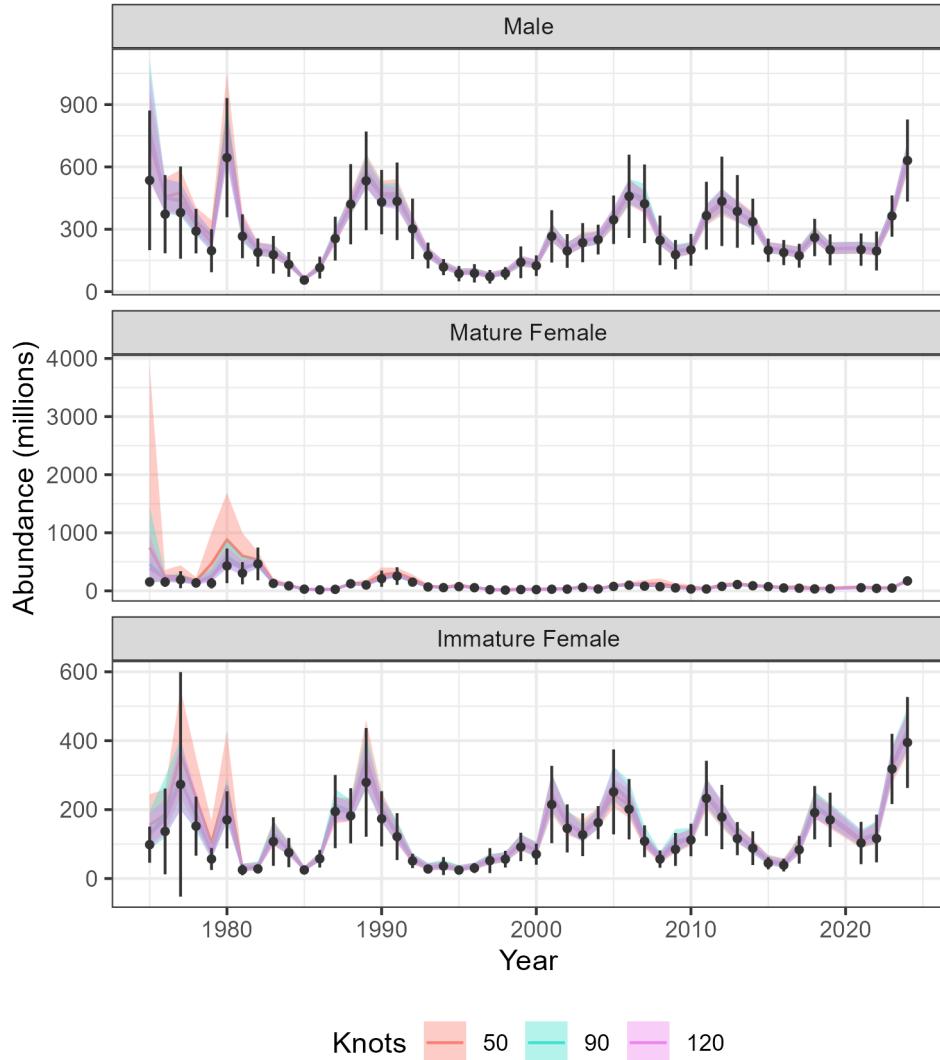


Figure 1: Estimated abundance (millions) for Eastern Bering Sea Tanner crab (*Chionoecetes bairdi*). Colored lines represent abundance ( $\pm 95\%$  CI) estimated by sdmTMB, with pink and blue denoting models fit with a 50- and 120-knot mesh, respectively. Black points represent abundance ( $\pm 95\%$  CI) estimated by the NMFS summer bottom trawl survey.

## EBS Tanner estimated biomass

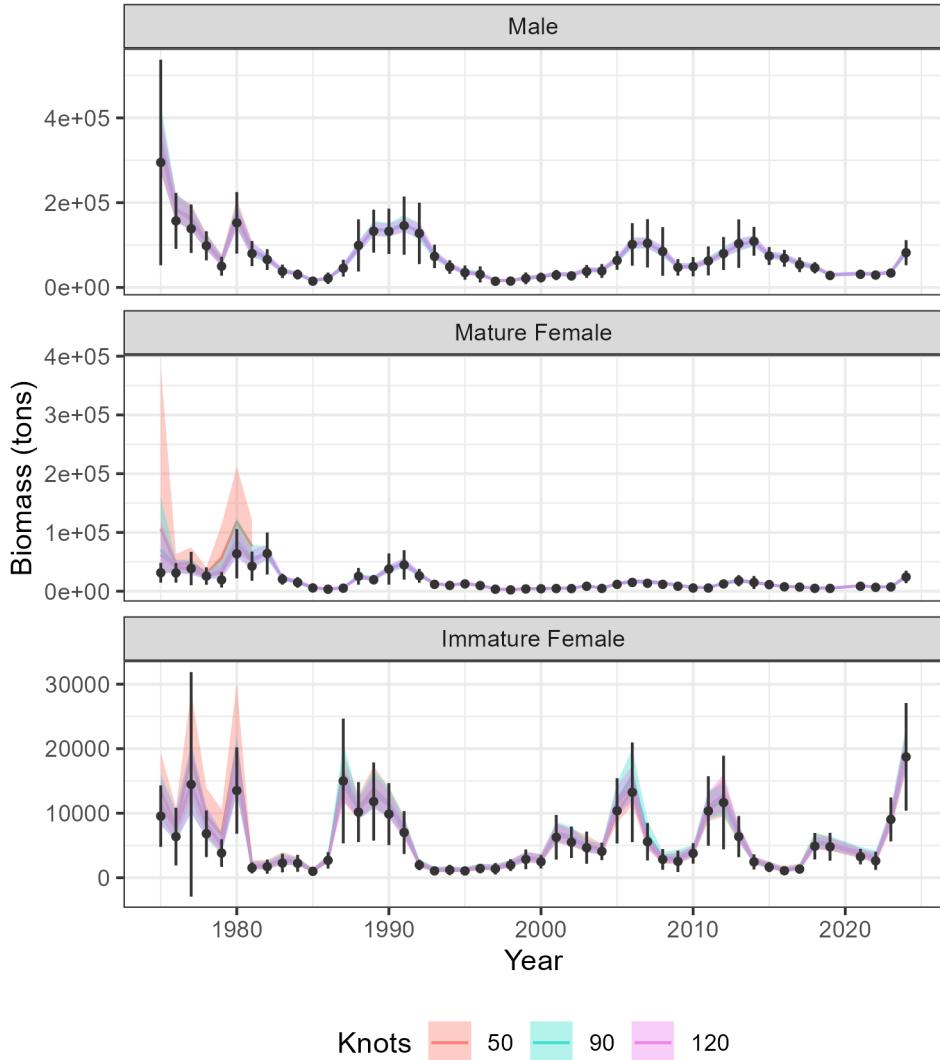


Figure 2: Estimated biomass (tons) for Eastern Bering Sea Tanner crab (*Chionoecetes bairdi*). Colored lines represent abundance ( $\pm 95\%$  CI) estimated by sdmTMB, with pink and blue denoting models fit with a 50- and 120-knot mesh, respectively. Black points represent abundance ( $\pm 95\%$  CI) estimated by the NMFS summer bottom trawl survey.

# **Results**

## **Tanner crab**

### **1 ## Tanner crab**

#### **Norton Sound red king crab**

##### **Model diagnostics**

The DHARMA residuals diagnostic plots show evidence of quantile deviations for all three NSRKC models (Figures 12 - 14). The models with 100 knots and 50 knots showed evidence of underdispersion, with observed data less dispersed than expected under the fitted models, while the model with 30 knots did not. None of the models showed evidence of outliers or zero inflation.

##### **Predicted abundance**

Heat maps of predicted NSRKC abundance for the three models are show in figures 15 - 17.

#### **St. Matthew Island blue king crab**

##### **Model diagnostics**

Examination of DHARMA residuals showed similar patterns for the three SMBKC models (Figures 24 - 26). All three models showed evidence of underdispersion, with observed data less dispersed than expected under the fitted models. None of the models showed evidence of outliers. All three models showed evidence of quantile deviations. The model with 120 knots showed evidence of zero inflation, with the observed data containing more zeros than would be expected under the fitted model, but the models with 50 and 90 knots did not show evidence of zero inflation.

##### **Predicted abundance**

Heat maps of predicted SMBKC abundance for the three models are show in figures 27 - 29.

##### **Predicted index fits to observations**

The model-predicted indices varied in their fits to the survey biomass observations, with the model fit using a mesh with an intermediate number of knots seeming to fit the survey observations more closely than the models fit to meshes with either higher or lower numbers of knots (Figure 30).

## **Conclusions**

## **Acknowledgements**

The authors thank Katie Palof and Mike Litzow for their support and feedback on this work.

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## Tables

## 2 Figures

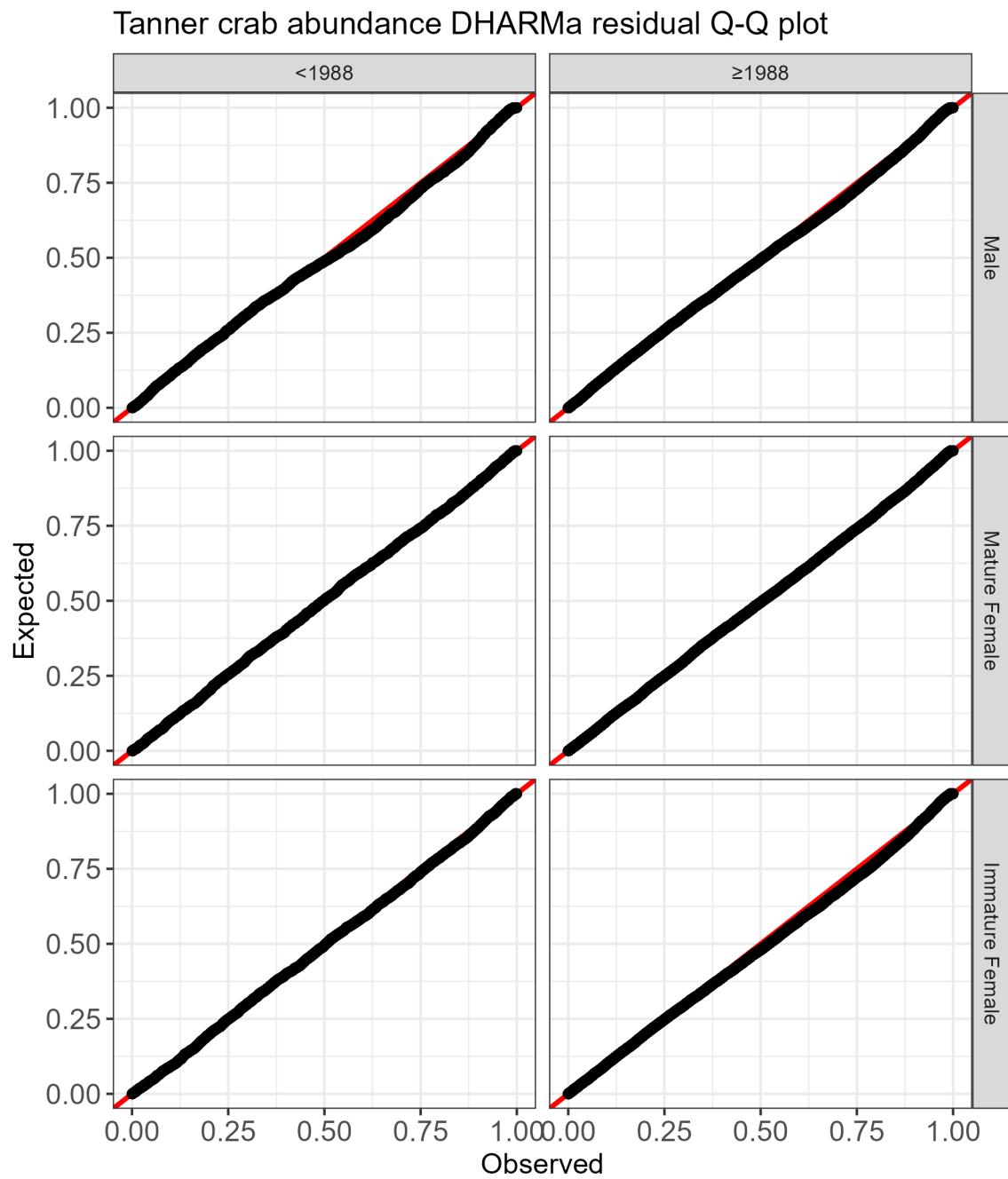


Figure 3: Q-Q plot of DHARMa residuals for EBS-wide abundance models fit using NMFS summer bottom trawl survey data before 1988 (left) and 1988 onward (right) using 120 knots in the model mesh.

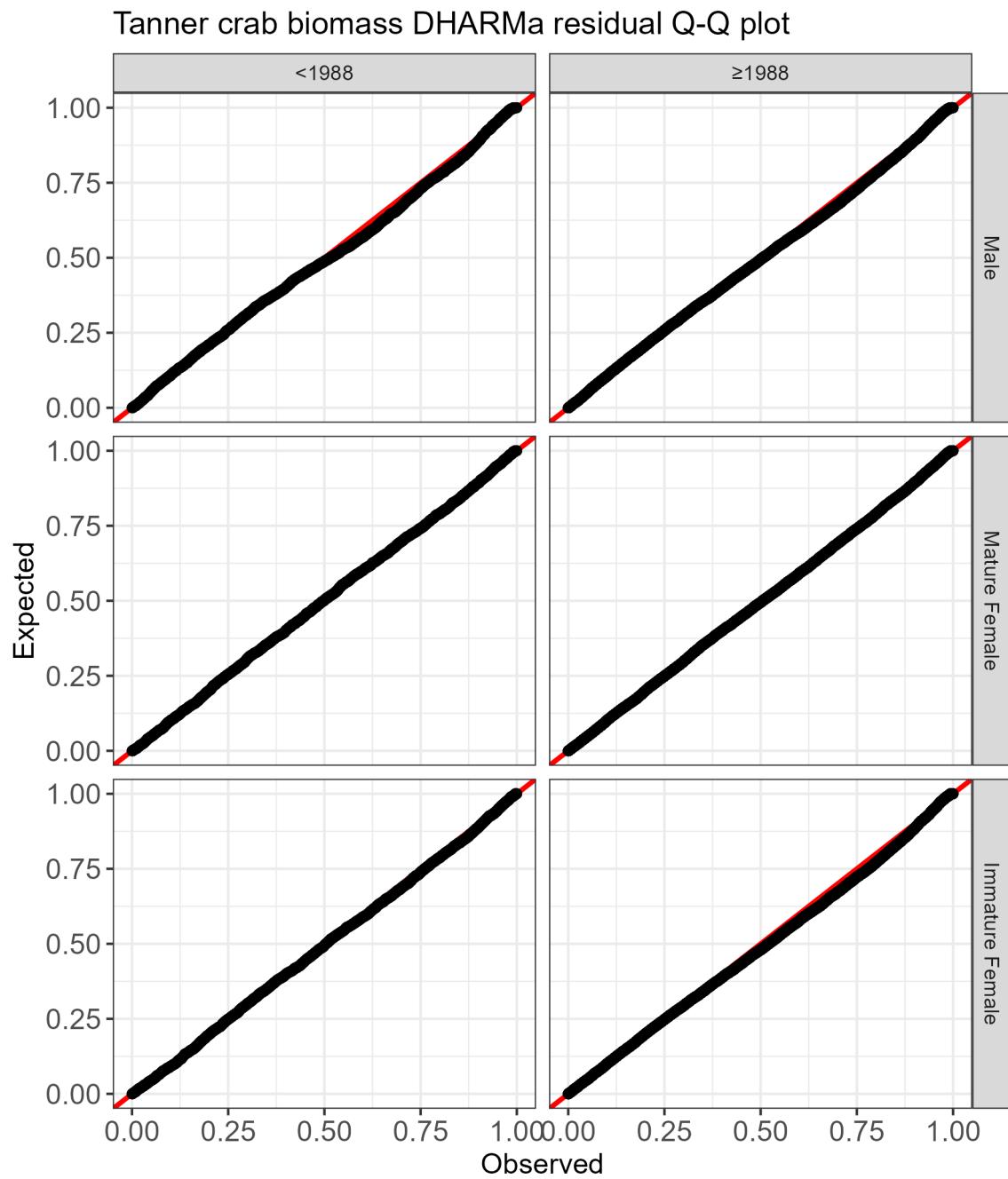


Figure 4: Q-Q plot of DHARMA residuals for EBS-wide biomass models fit using NMFS summer bottom trawl survey data before 1988 (left) and 1988 onward (right) using 120 knots in the model mesh.

### All Male abundance residuals (knots=120)

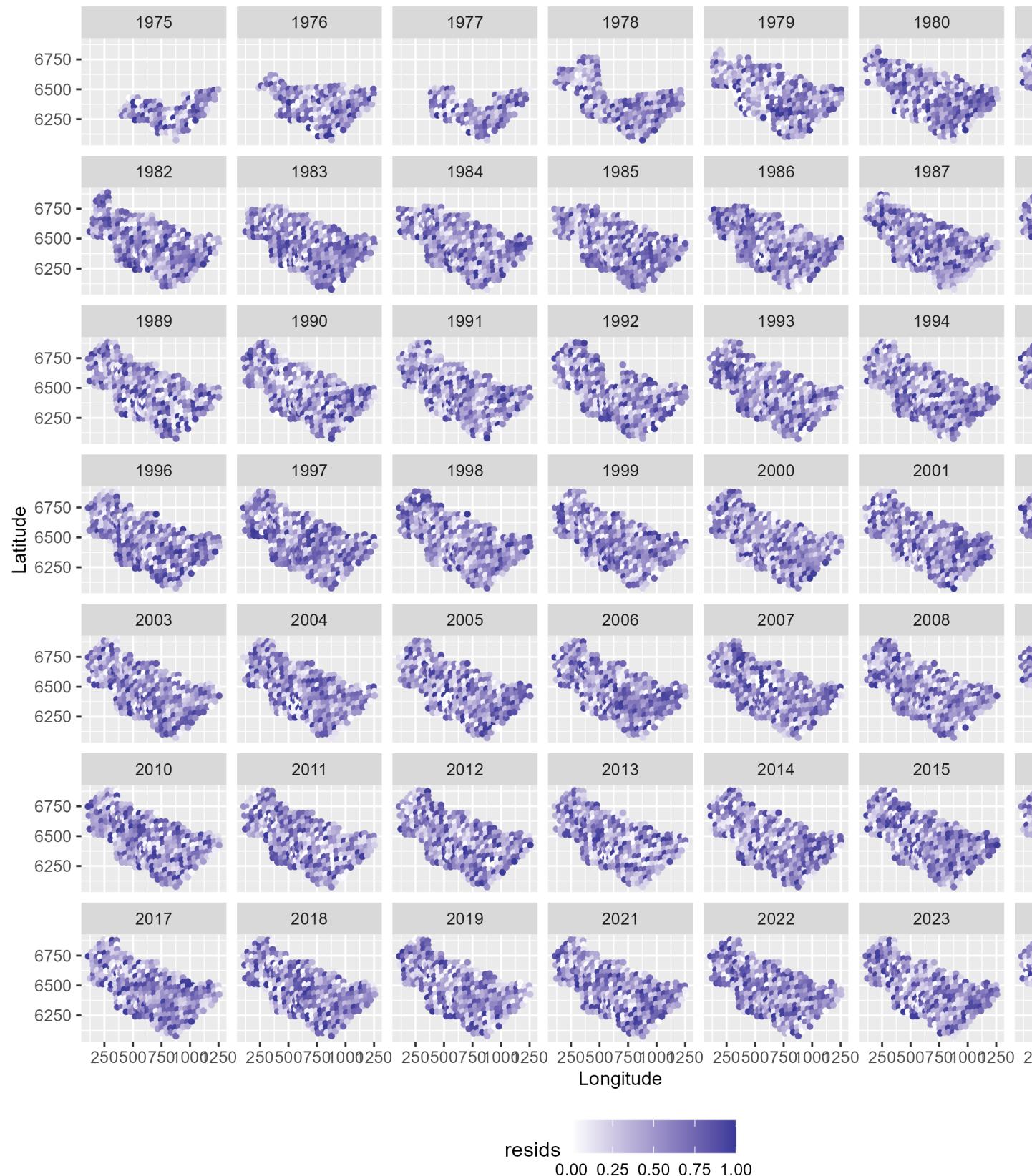


Figure 5: Spatial plot of DHARMA residuals for EBS-wide male abundance models fit using NMFS summer bottom trawl survey data before 1988 and 1988 onward. Predictions from both these periods/models are combined in this figure.

### EBS Tanner estimated abundance

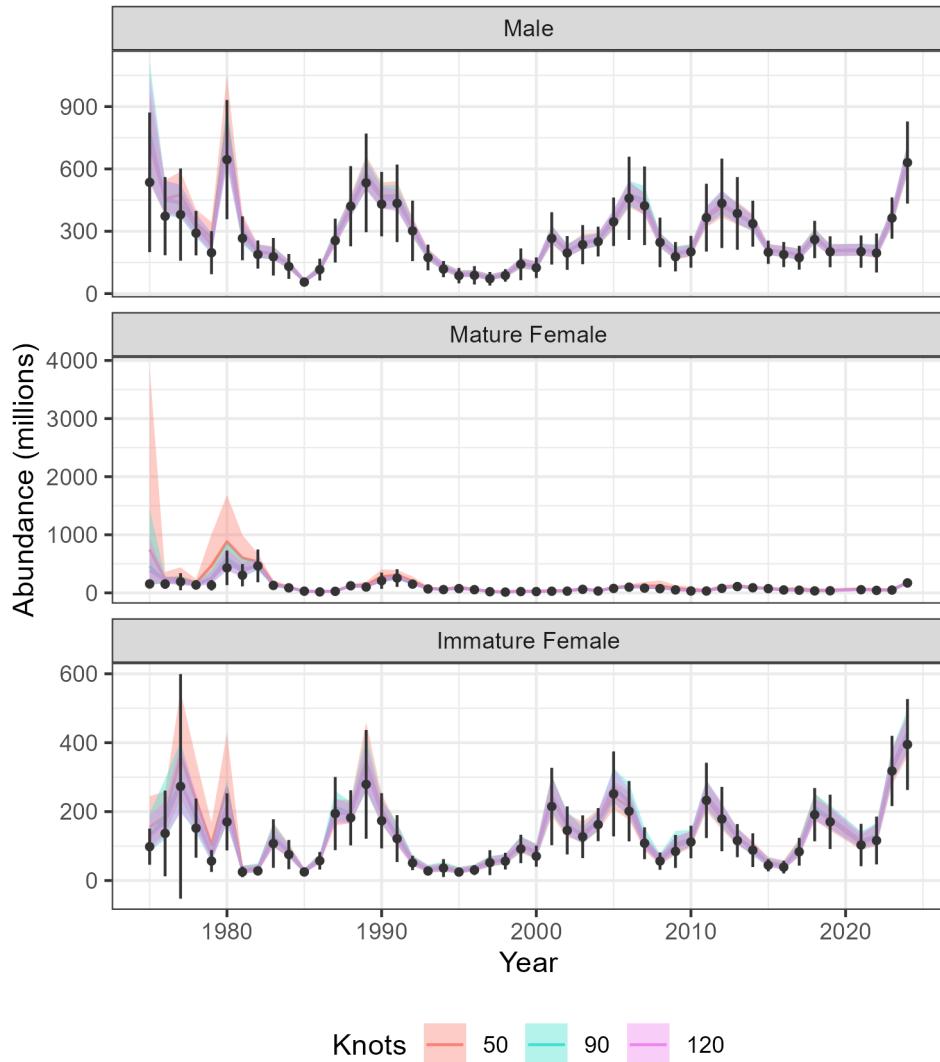


Figure 6: Estimated abundance (millions) for Eastern Bering Sea Tanner crab (*Chionoecetes bairdi*). Colored lines represent abundance ( $\pm 95\%$  CI) estimated by sdmTMB, with pink and blue denoting models fit with a 50- and 120-knot mesh, respectively. Black points represent abundance ( $\pm 95\%$  CI) estimated by the NMFS summer bottom trawl survey.

## EBS Tanner estimated biomass

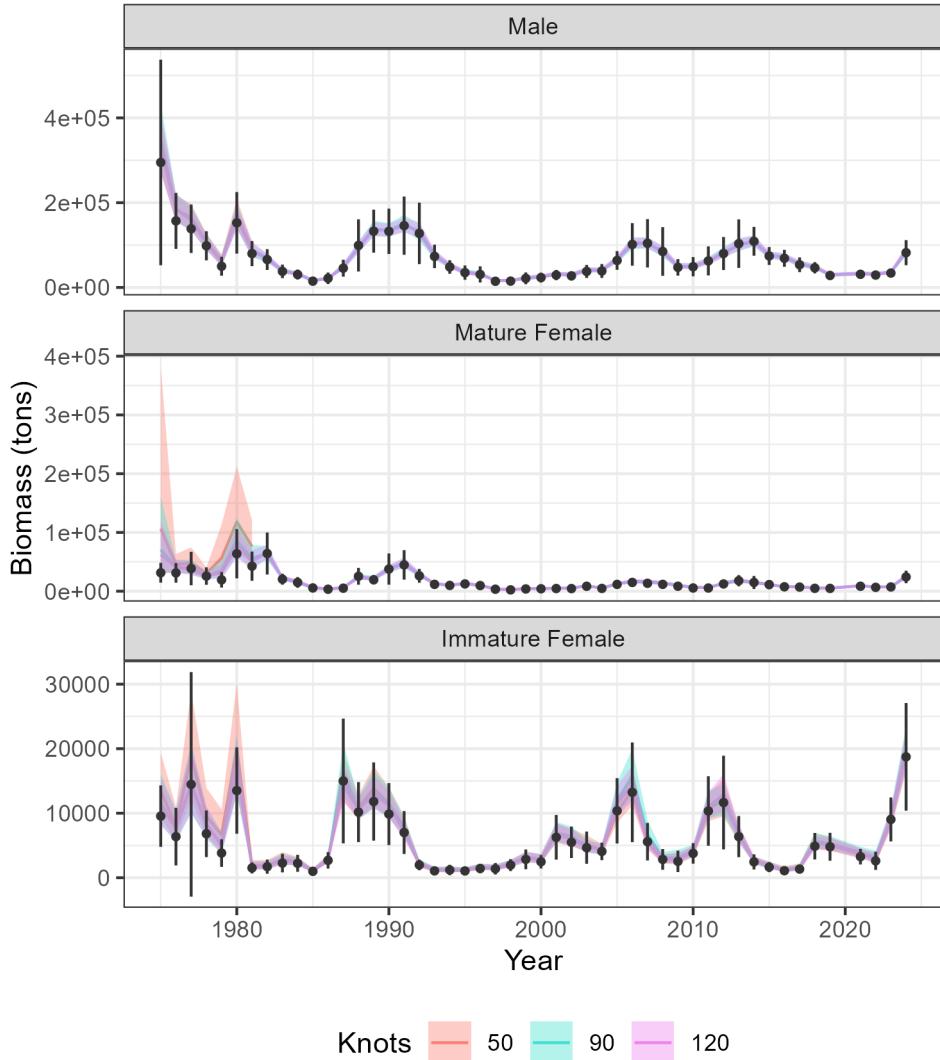


Figure 7: Estimated biomass (tons) for Eastern Bering Sea Tanner crab (*Chionoecetes bairdi*). Colored lines represent abundance ( $\pm 95\%$  CI) estimated by sdmTMB, with pink and blue denoting models fit with a 50- and 120-knot mesh, respectively. Black points represent abundance ( $\pm 95\%$  CI) estimated by the NMFS summer bottom trawl survey.

## Figures

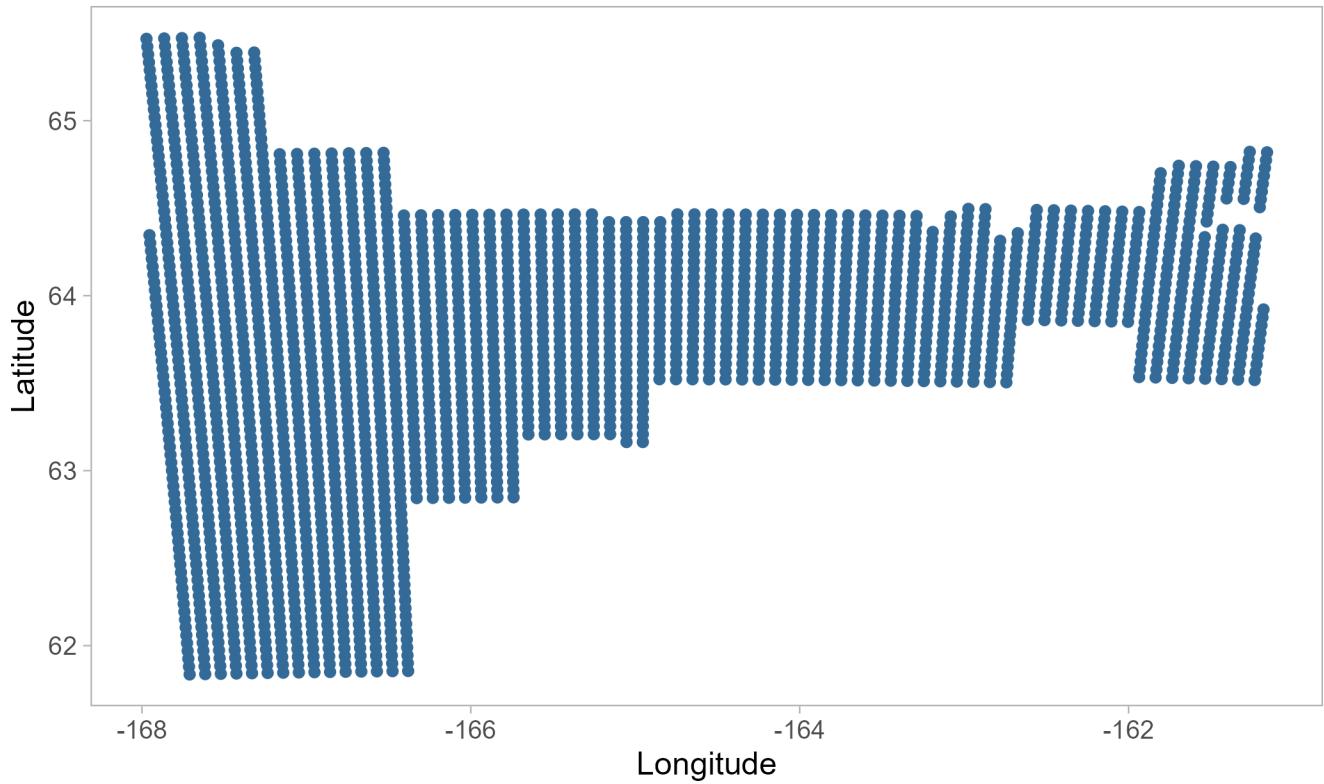


Figure 8: Prediction grid used for Norton Sound red king crab spatial abundance predictions. Spatial resolution is  $5 \text{ km}^2$ .

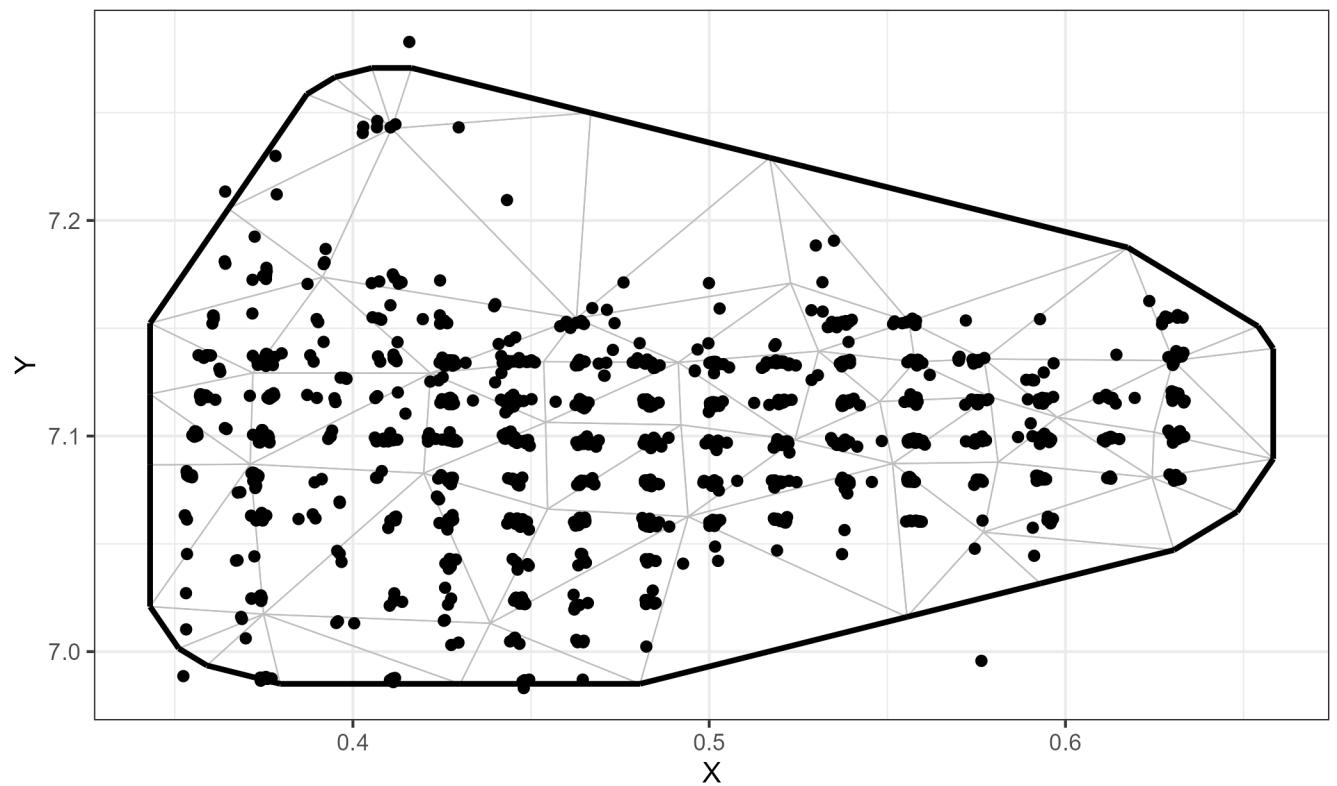


Figure 9: Spatial mesh with 30 knots used for fitting Norton Sound red king crab spatial models. Points represent observations and vertices represent knot locations.

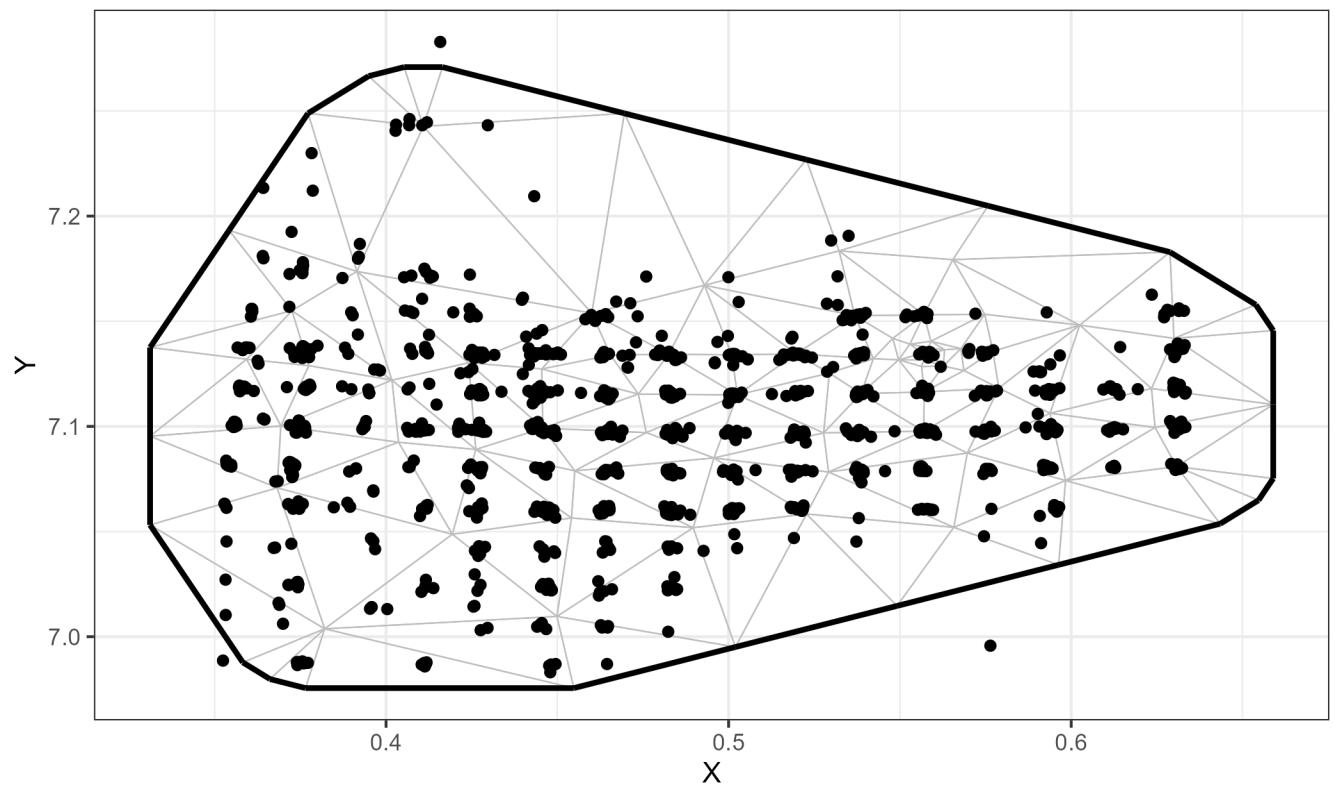


Figure 10: Spatial mesh with 50 knots used for fitting Norton Sound red king crab spatial models. Points represent observations and vertices represent knot locations.

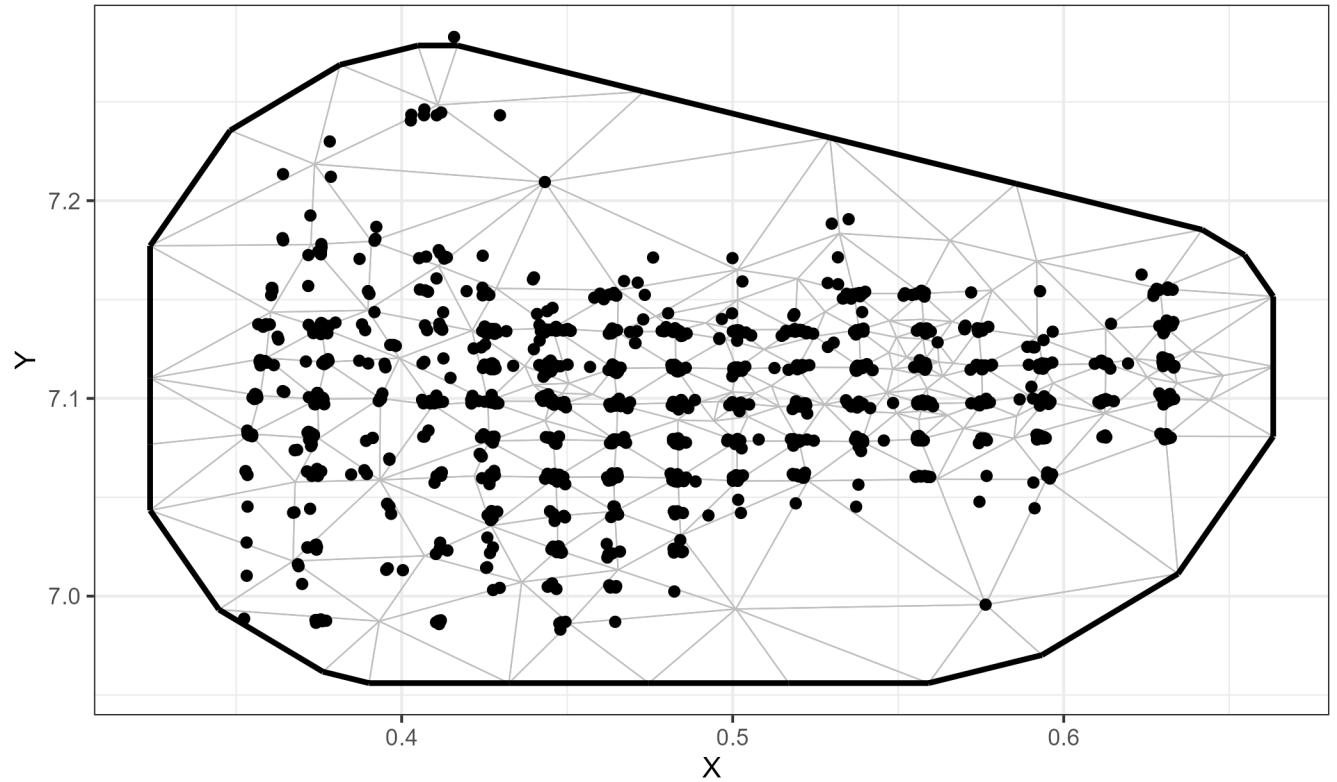


Figure 11: Spatial mesh with 100 knots used for fitting Norton Sound red king crab spatial models. Points represent observations and vertices represent knot locations.

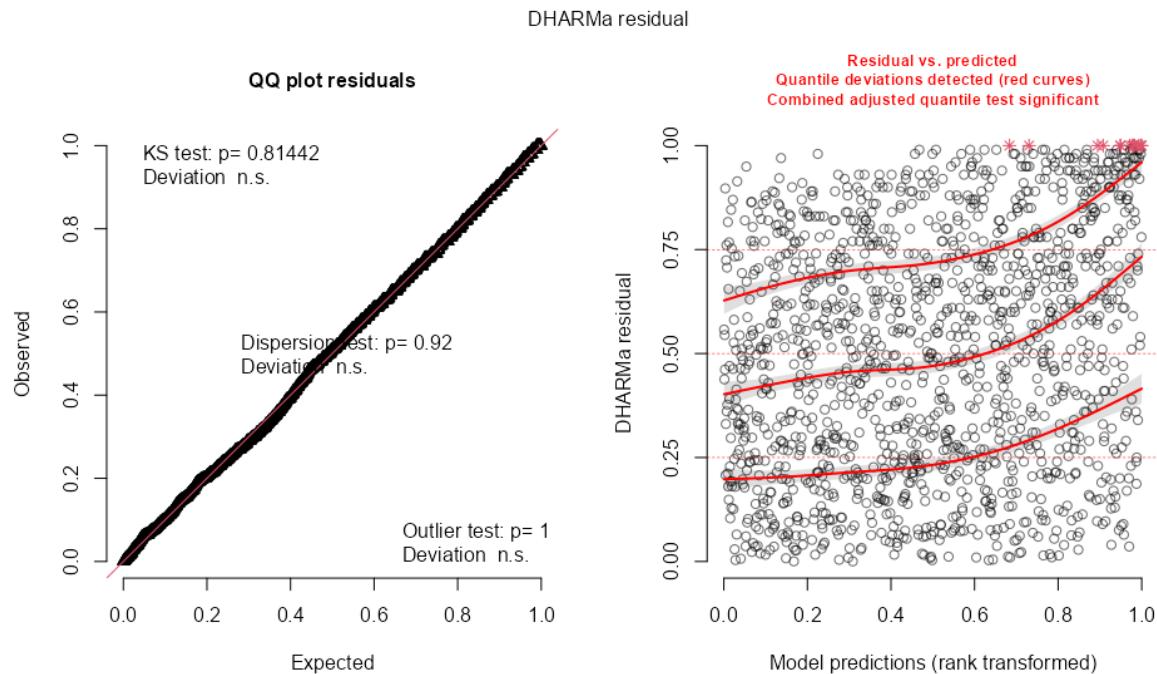


Figure 12: Model diagnostic plots using DHARMA residuals for the Norton Sound red king crab model with 30 knots.

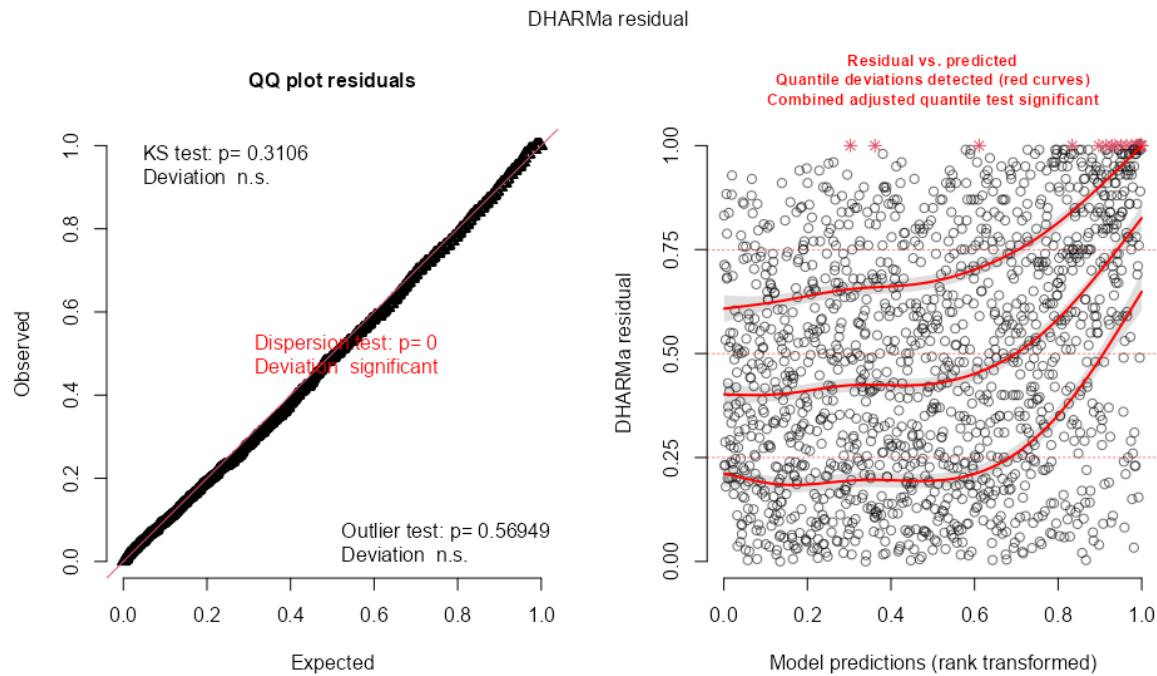


Figure 13: Model diagnostic plots using DHARMA residuals for the Norton Sound red king crab model with 50 knots.

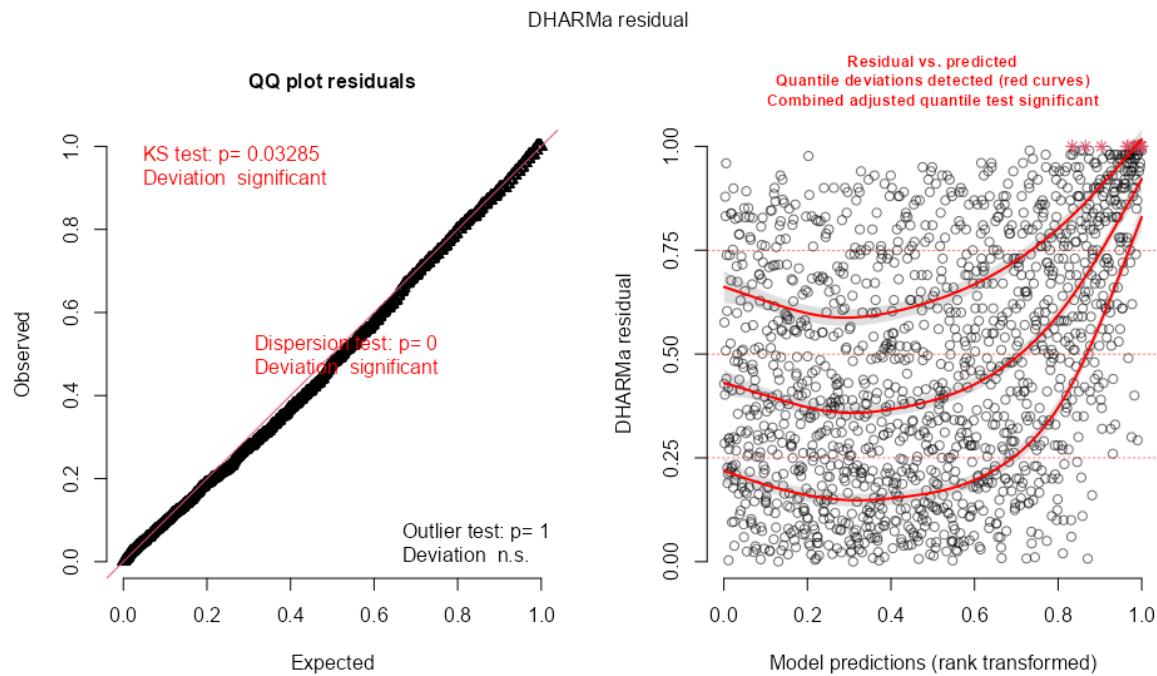


Figure 14: Model diagnostic plots using DHARMA residuals for Norton Sound red king crab model with 100 knots.

## NSRKC predicted abundance, 30 knots



Figure 15: Heat map of Norton Sound red king crab predicted abundance generated using the model with 50 knots.

## NSRKC predicted abundance, 50 knots



Figure 16: Heat map of Norton Sound red blue king crab predicted abundance generated using the model with 90 knots.

## NSRKC predicted abundance, 100 knots

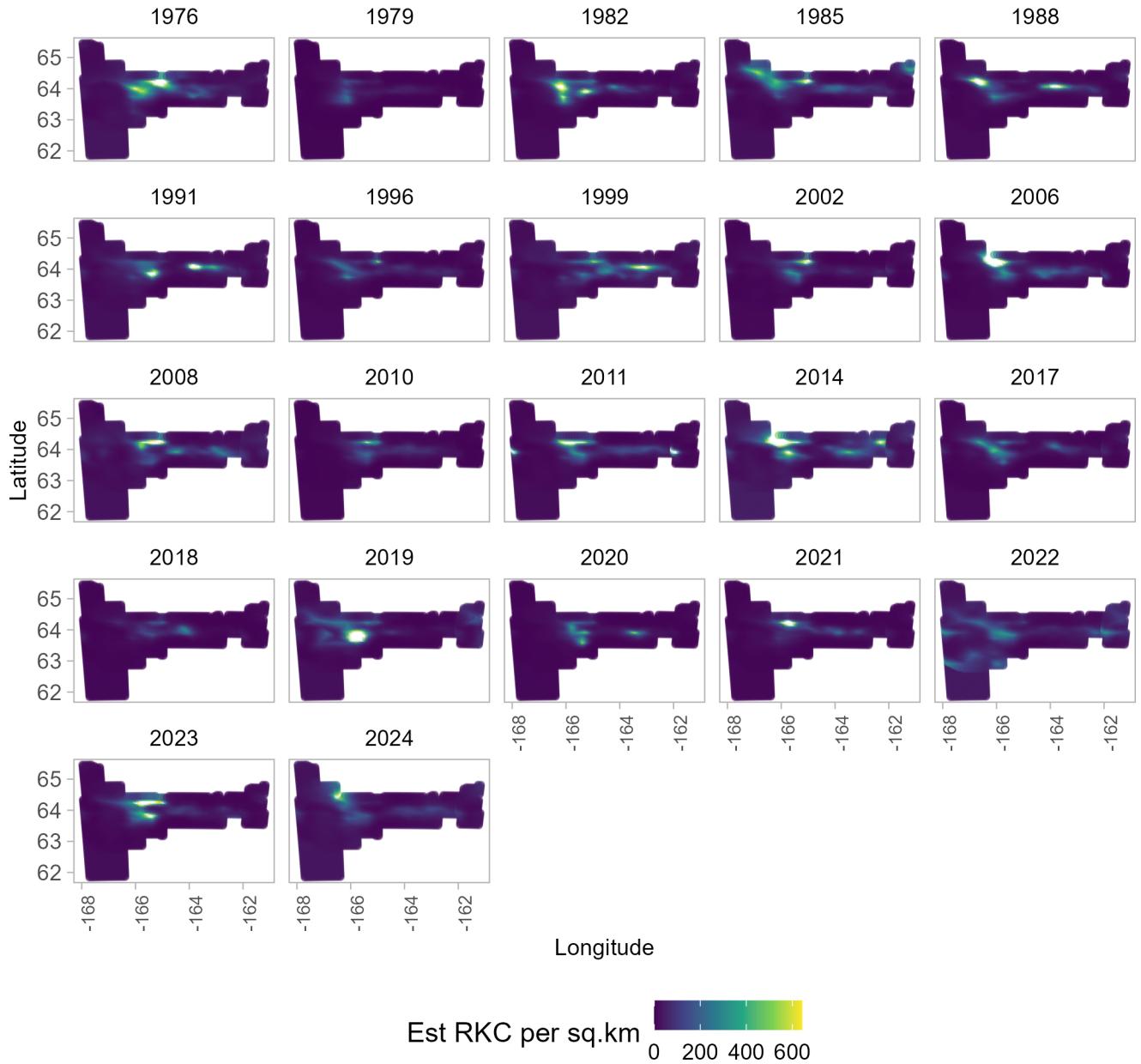


Figure 17: Heat map of Norton Sound red blue king crab predicted abundance generated using the model with 120 knots.

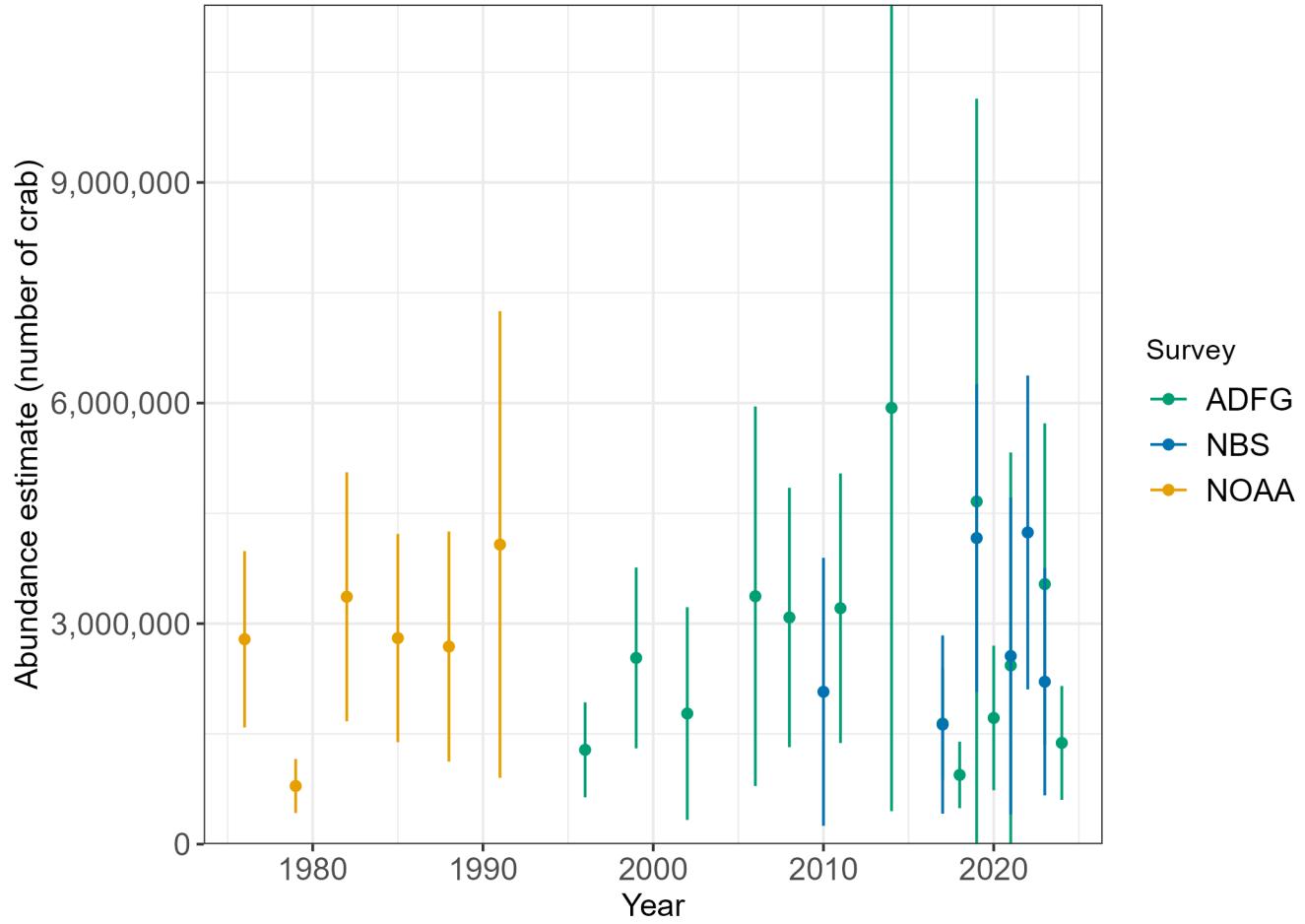


Figure 18: Estimated abundance in number of crab for Norton Sound red king crab. Colored points represent abundance ( $\pm 95\%$  CI) estimated by the trawl surveys.

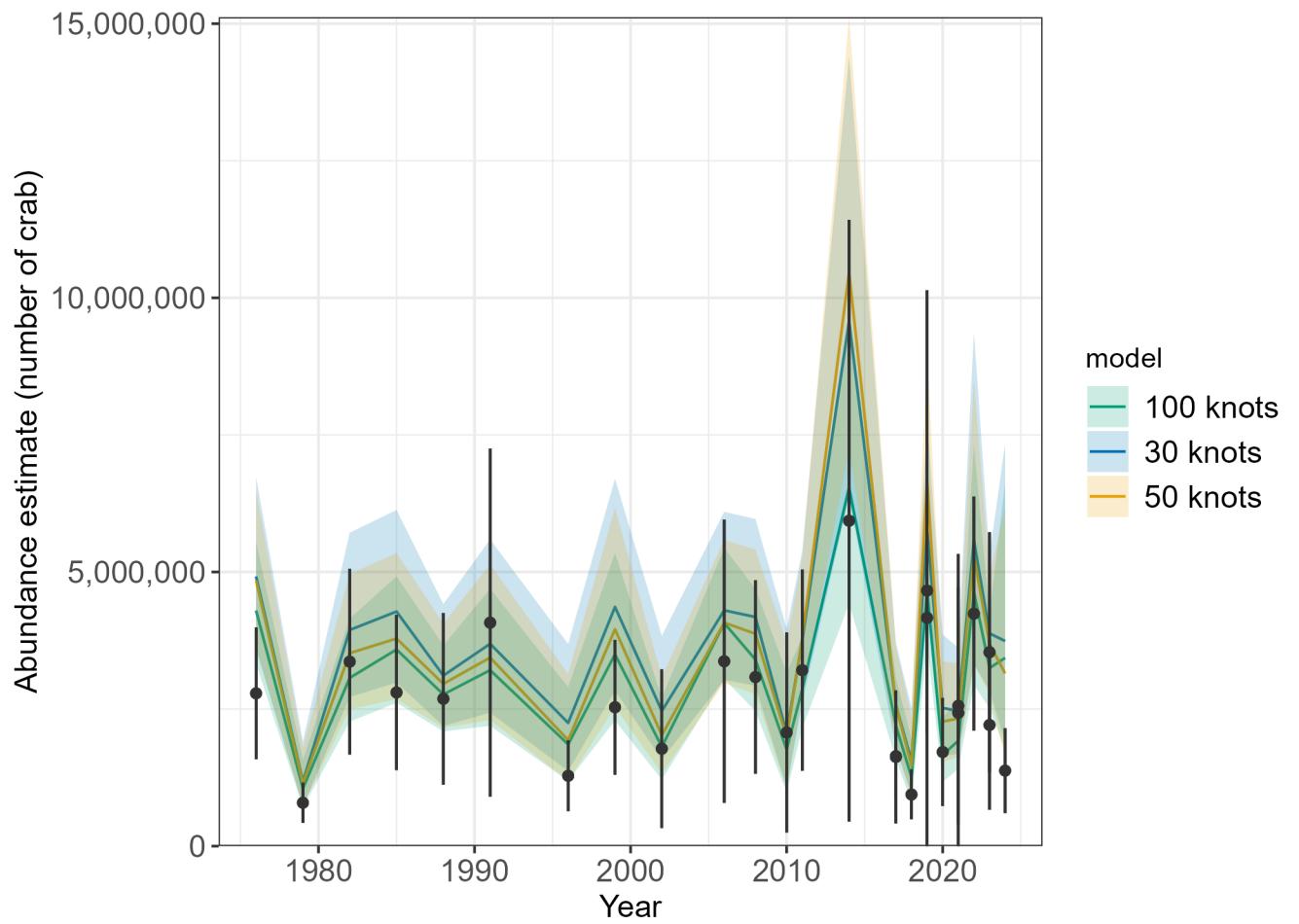


Figure 19: Estimated abundance in number of crab for Norton Sound red king crab. Colored lines represent abundance ( $\pm 95\%$  CI) estimated using sdmTMB. Black points represent abundance ( $\pm 95\%$  CI) estimated by the trawl surveys.

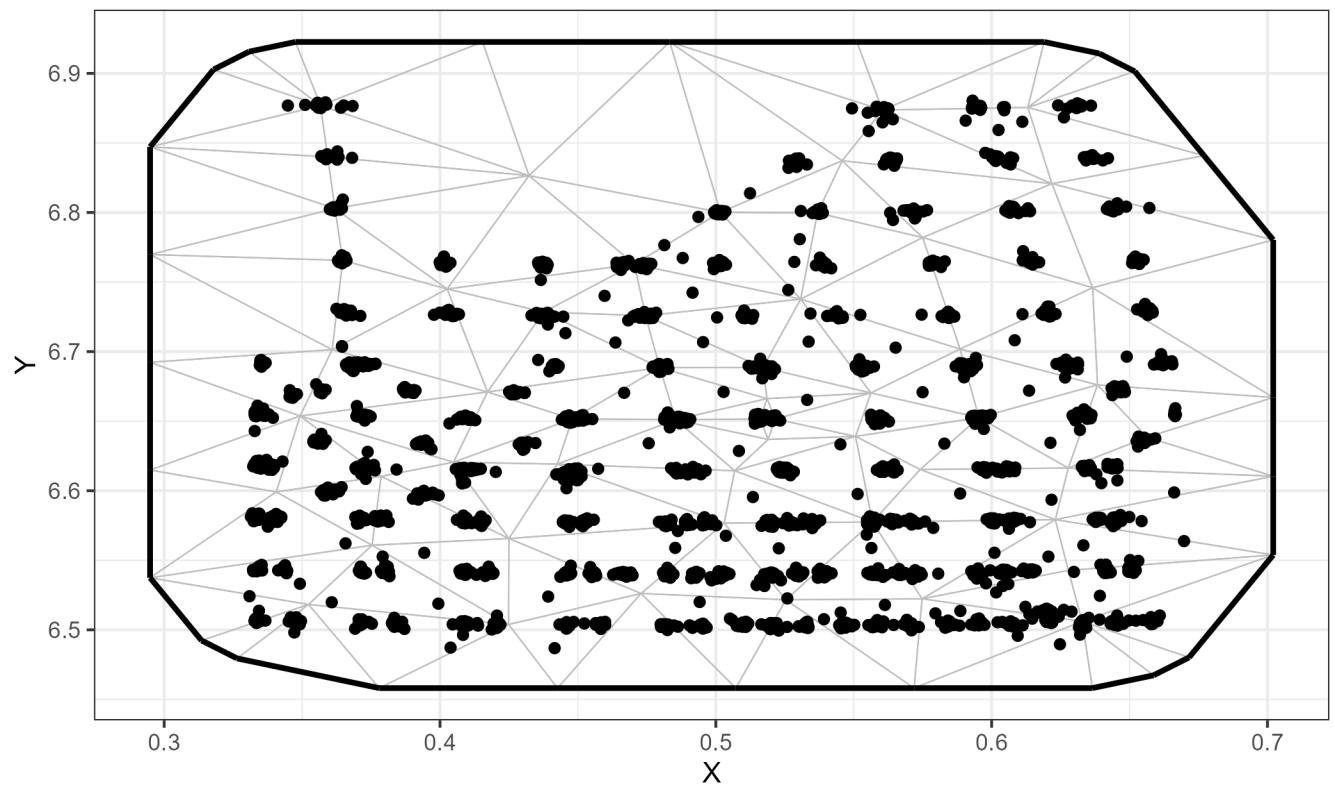


Figure 20: Spatial mesh with 50 knots used for fitting St. Matthew Island blue king crab spatial models. Points represent observations and vertices represent knot locations.

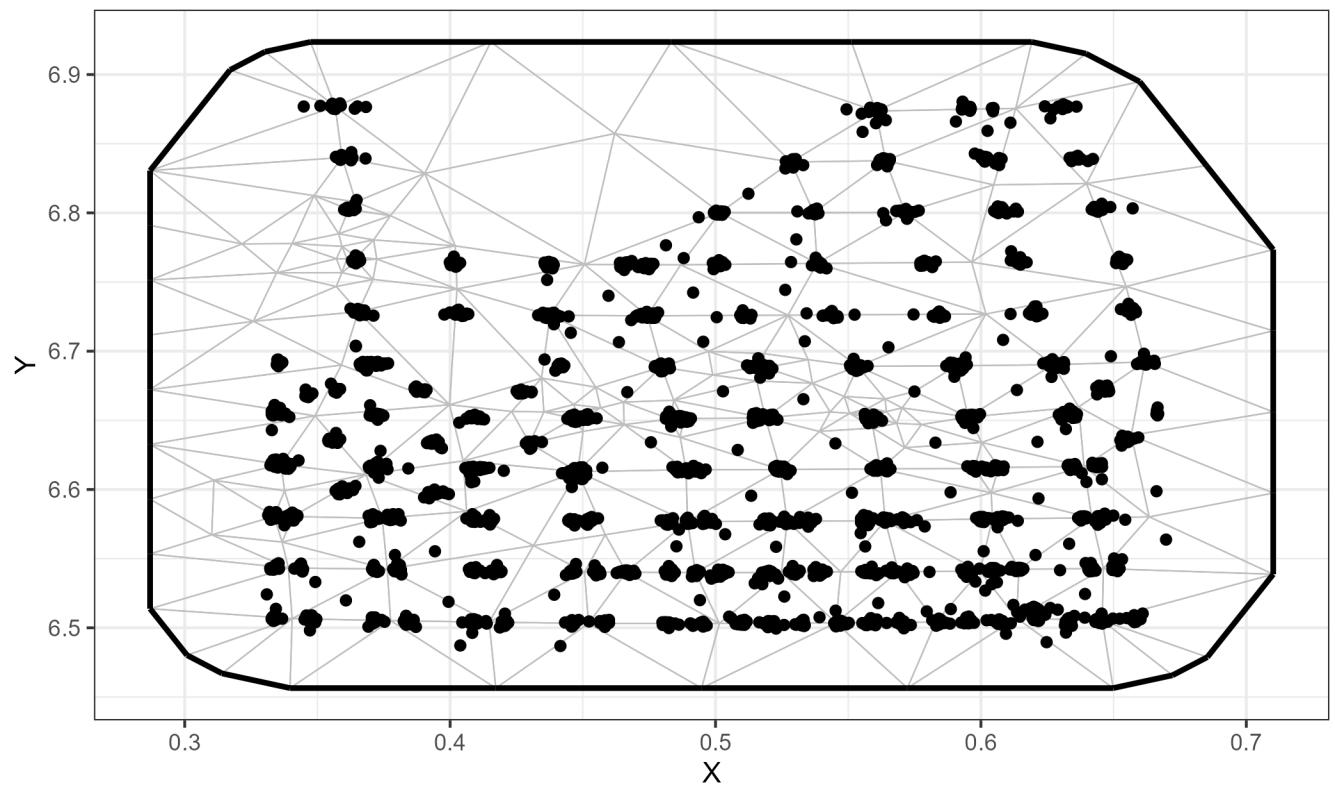


Figure 21: Spatial mesh with 90 knots used for fitting St. Matthew Island blue king crab spatial models. Points represent observations and vertices represent knot locations.

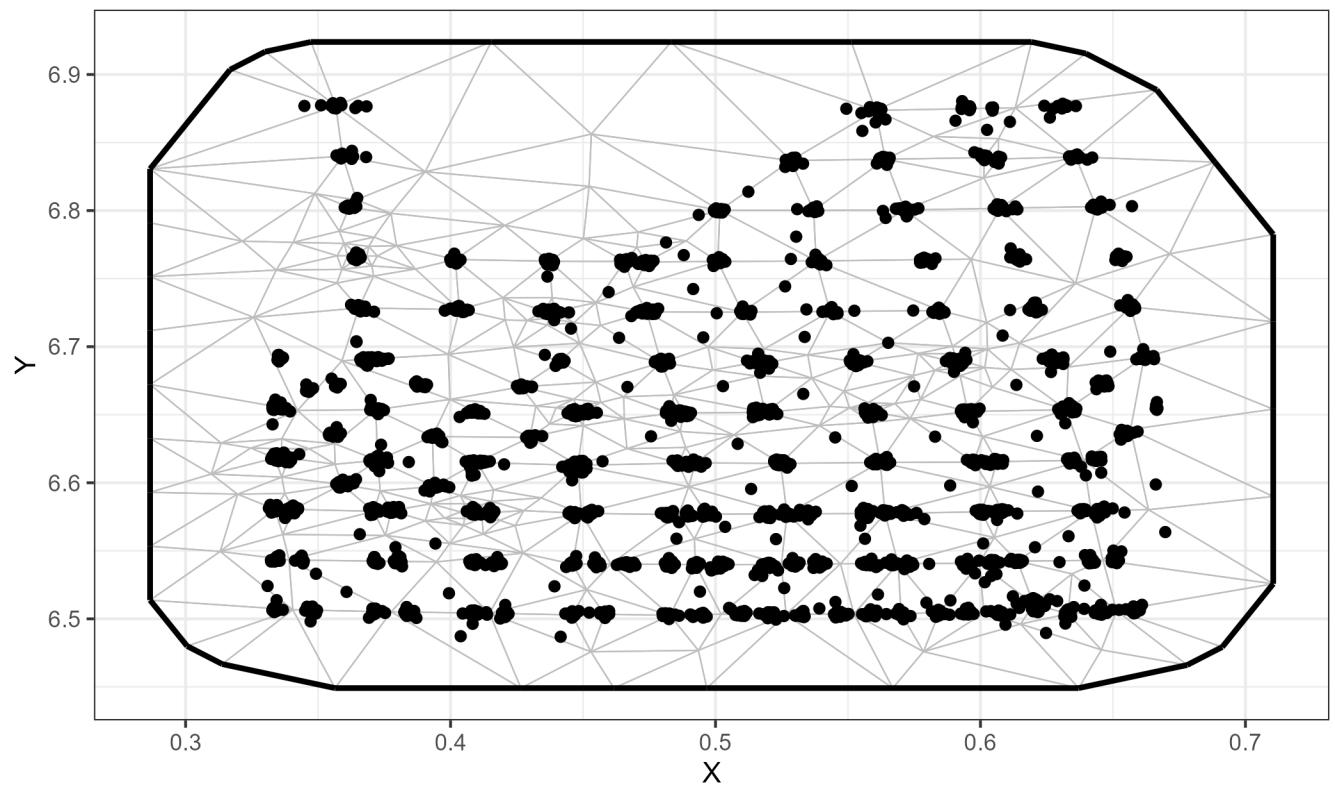


Figure 22: Spatial mesh with 120 knots used for fitting St. Matthew Island blue king crab spatial models. Points represent observations and vertices represent knot locations.

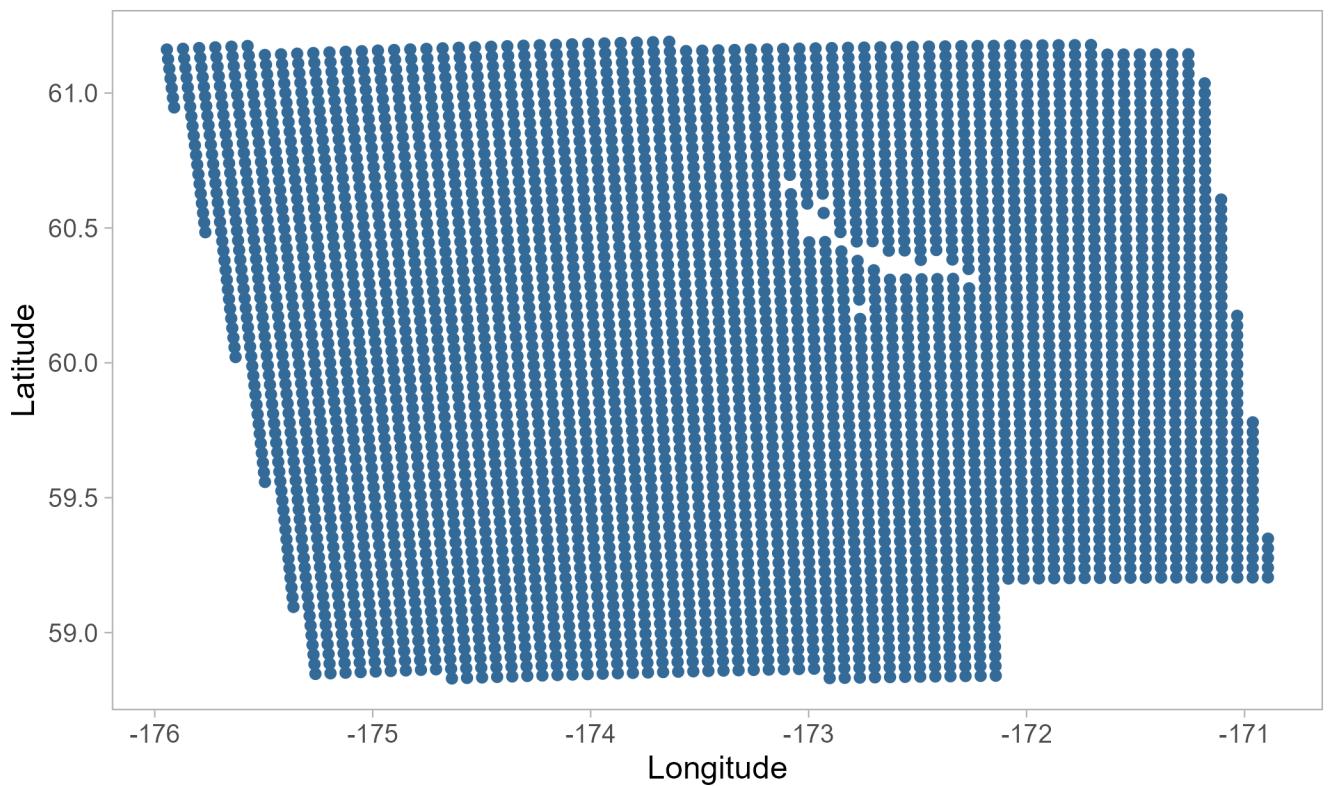


Figure 23: Prediction grid used for St. Matthew Island blue king crab spatial abundance predictions. Spatial resolution is 4 km<sup>2</sup>.

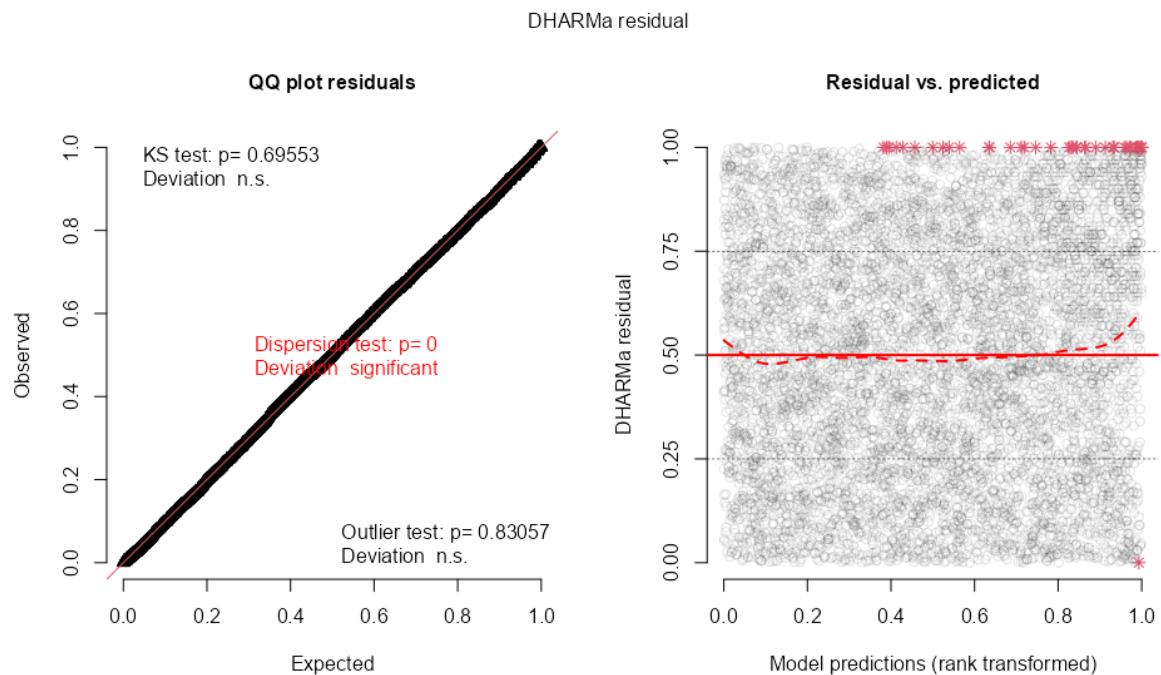


Figure 24: Model diagnostic plots using DHARMA residuals for the model with 50 knots.

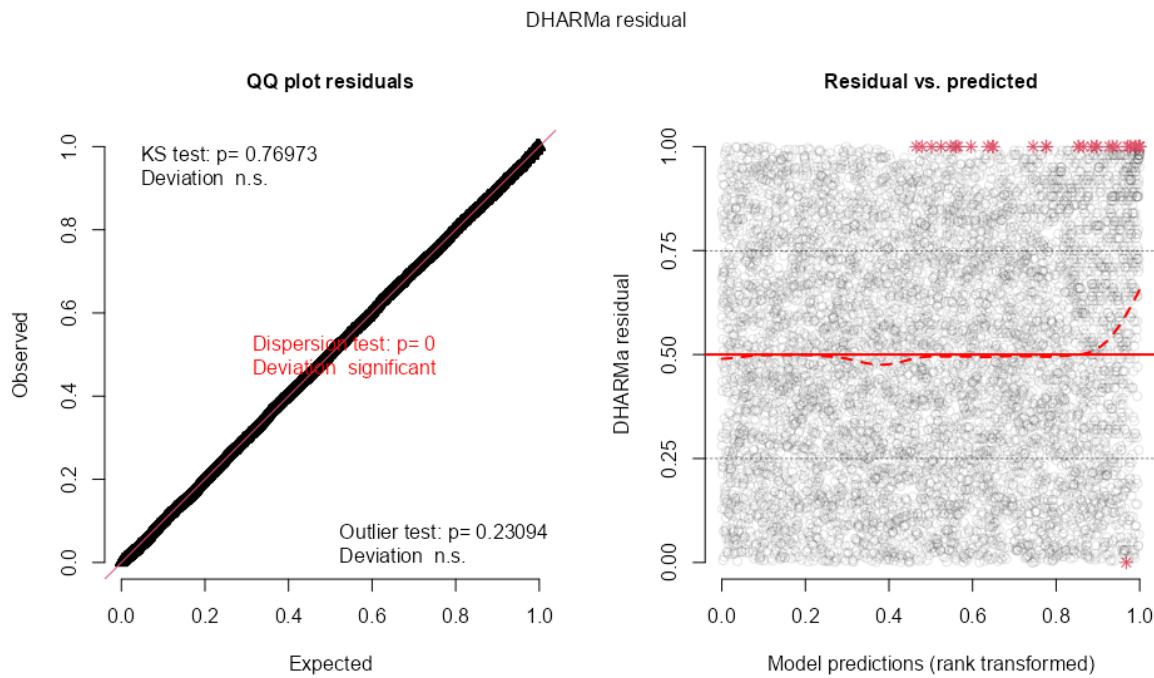


Figure 25: Model diagnostic plots using DHARMA residuals for the model with 90 knots.

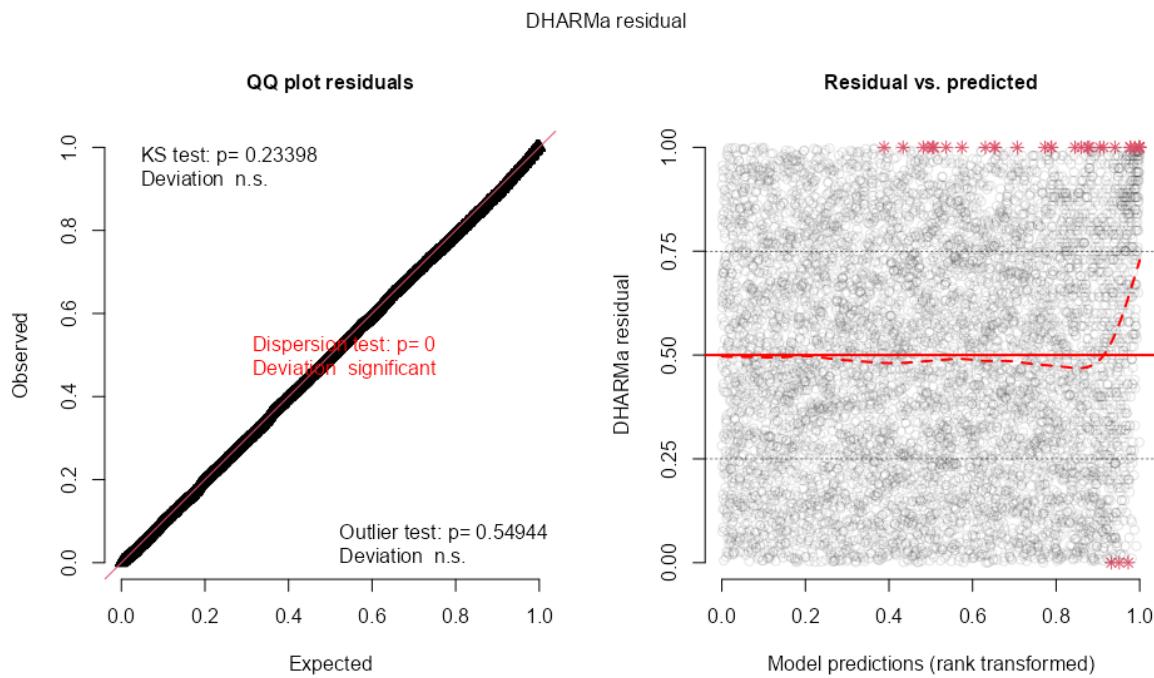


Figure 26: Model diagnostic plots using DHARMA residuals for the model with 120 knots.

### BKC predicted abundance, 50 knots

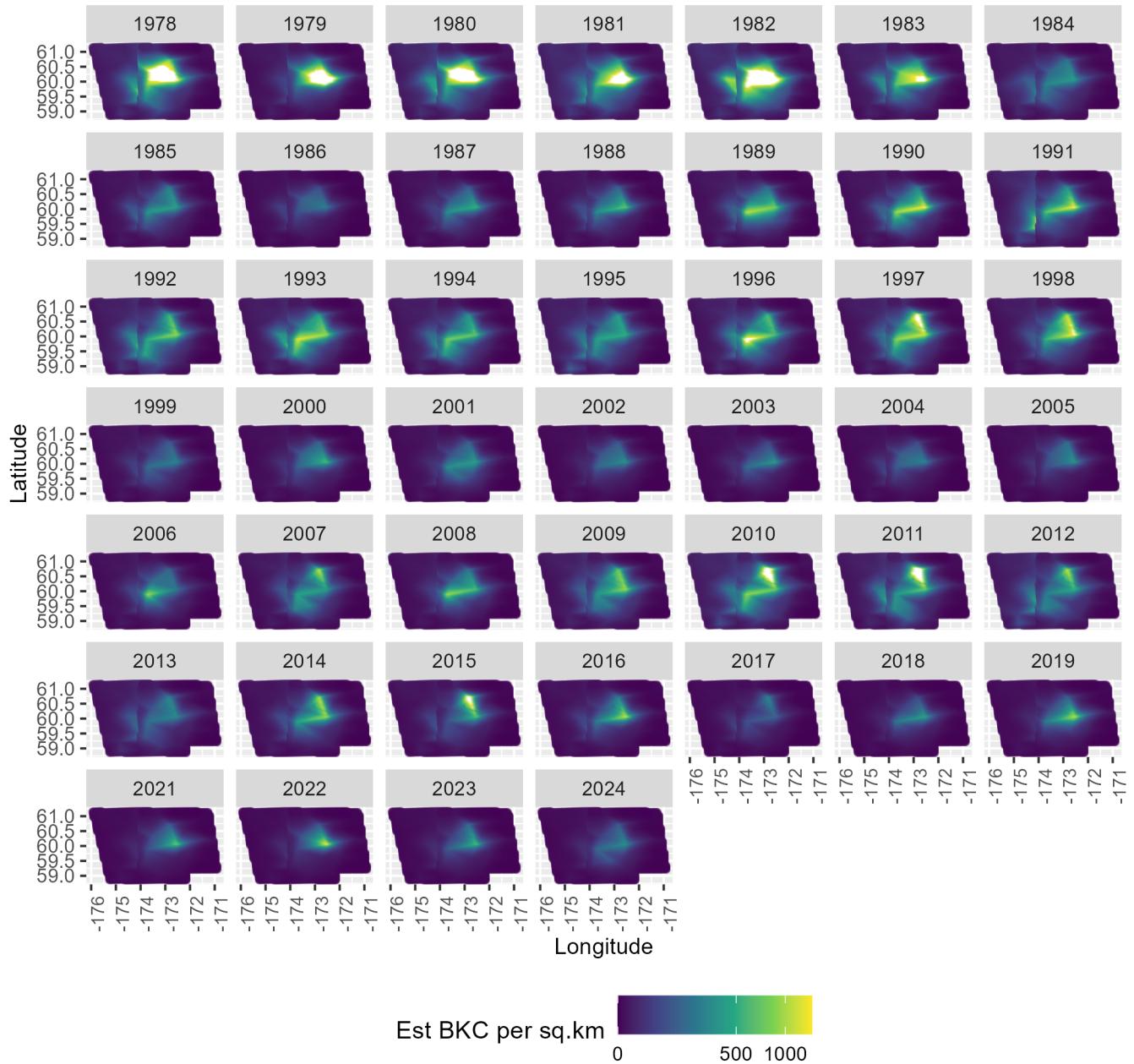


Figure 27: Heat map of St. Matthew Island blue king crab predicted abundance generated using the model with 50 knots.

### BKC predicted abundance, 90 knots

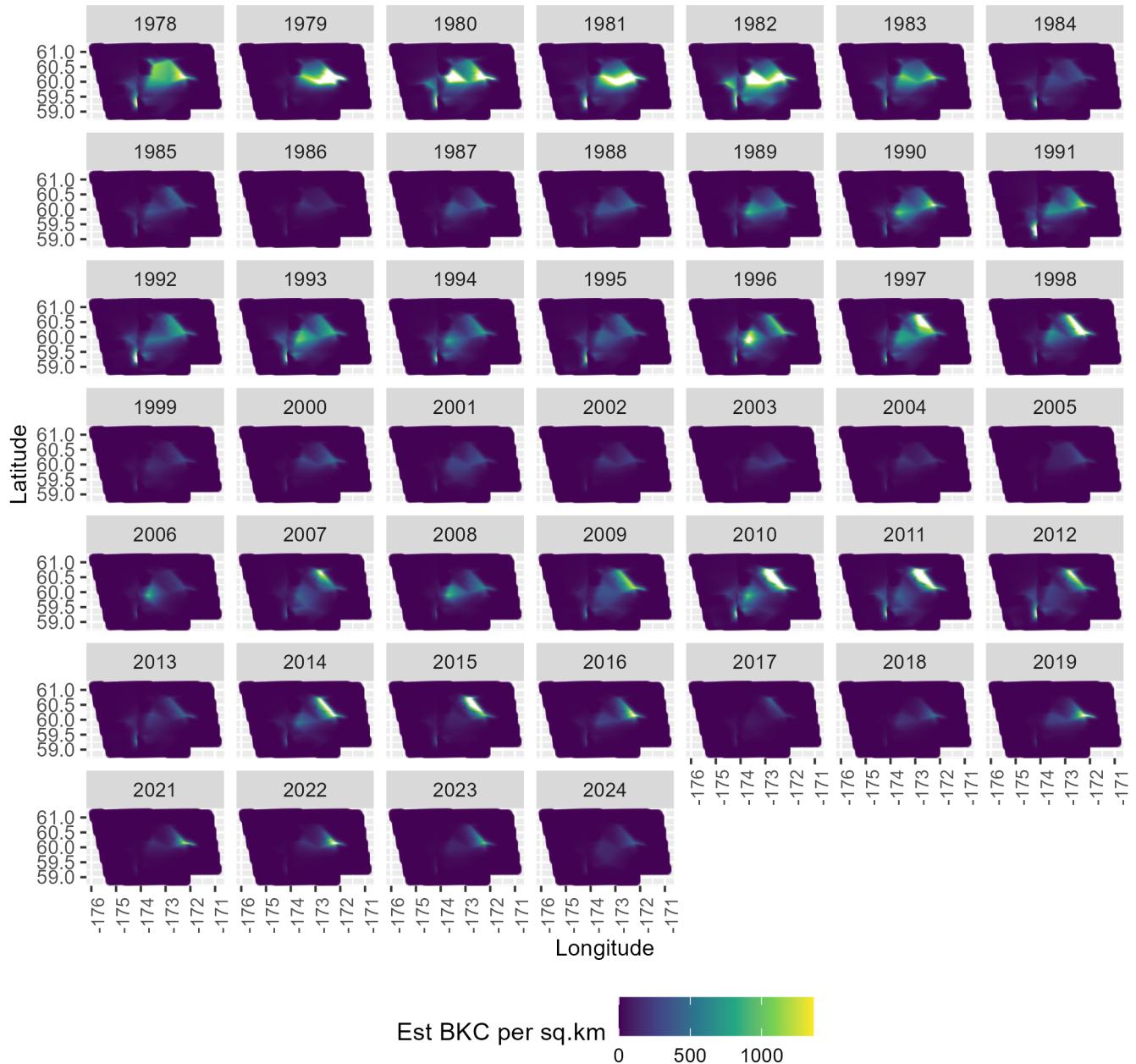


Figure 28: Heat map of St. Matthew Island blue king crab predicted abundance generated using the model with 90 knots.

### BKC predicted abundance, 120 knots

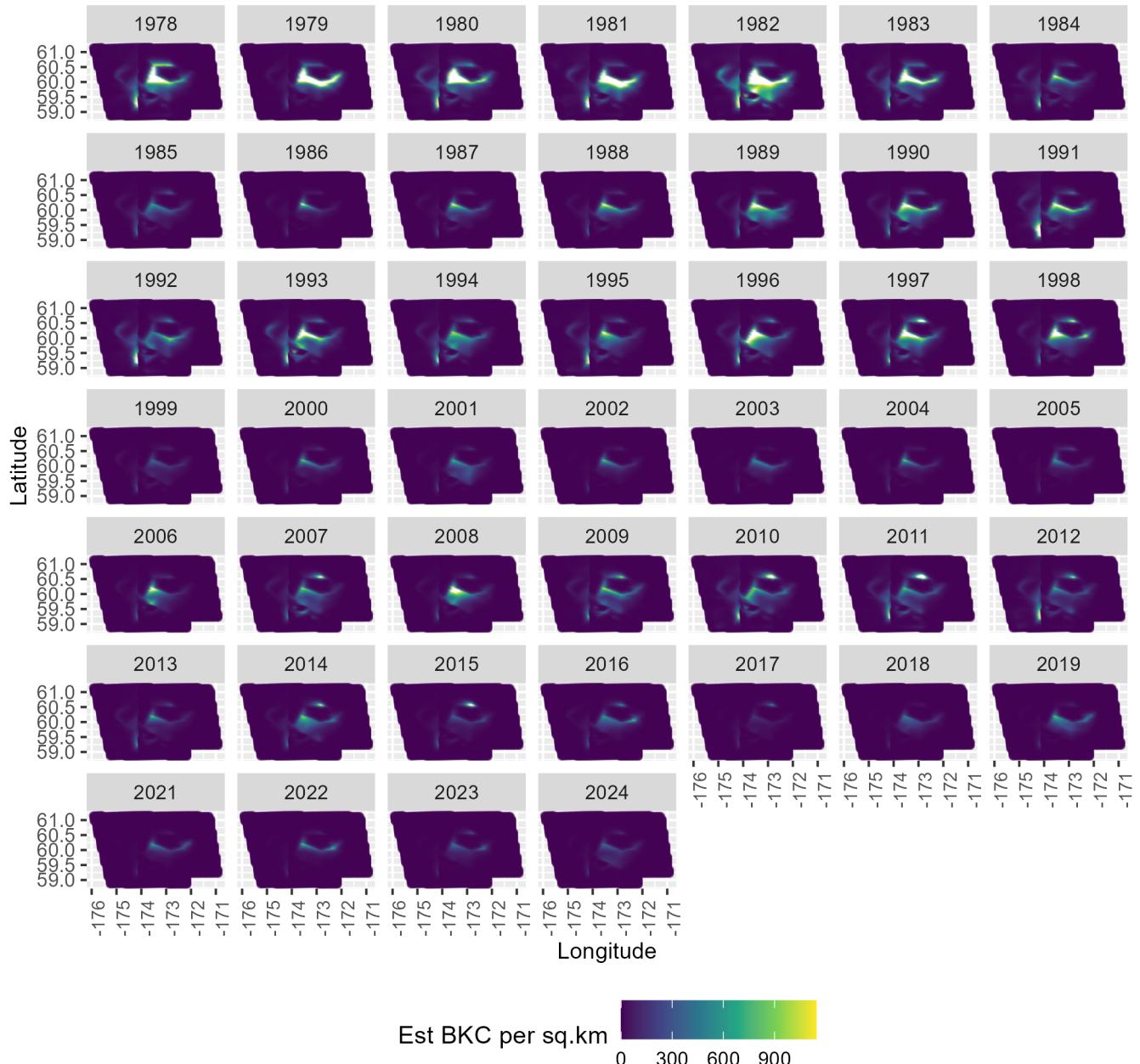


Figure 29: Heat map of St. Matthew Island blue king crab predicted abundance generated using the model with 120 knots.

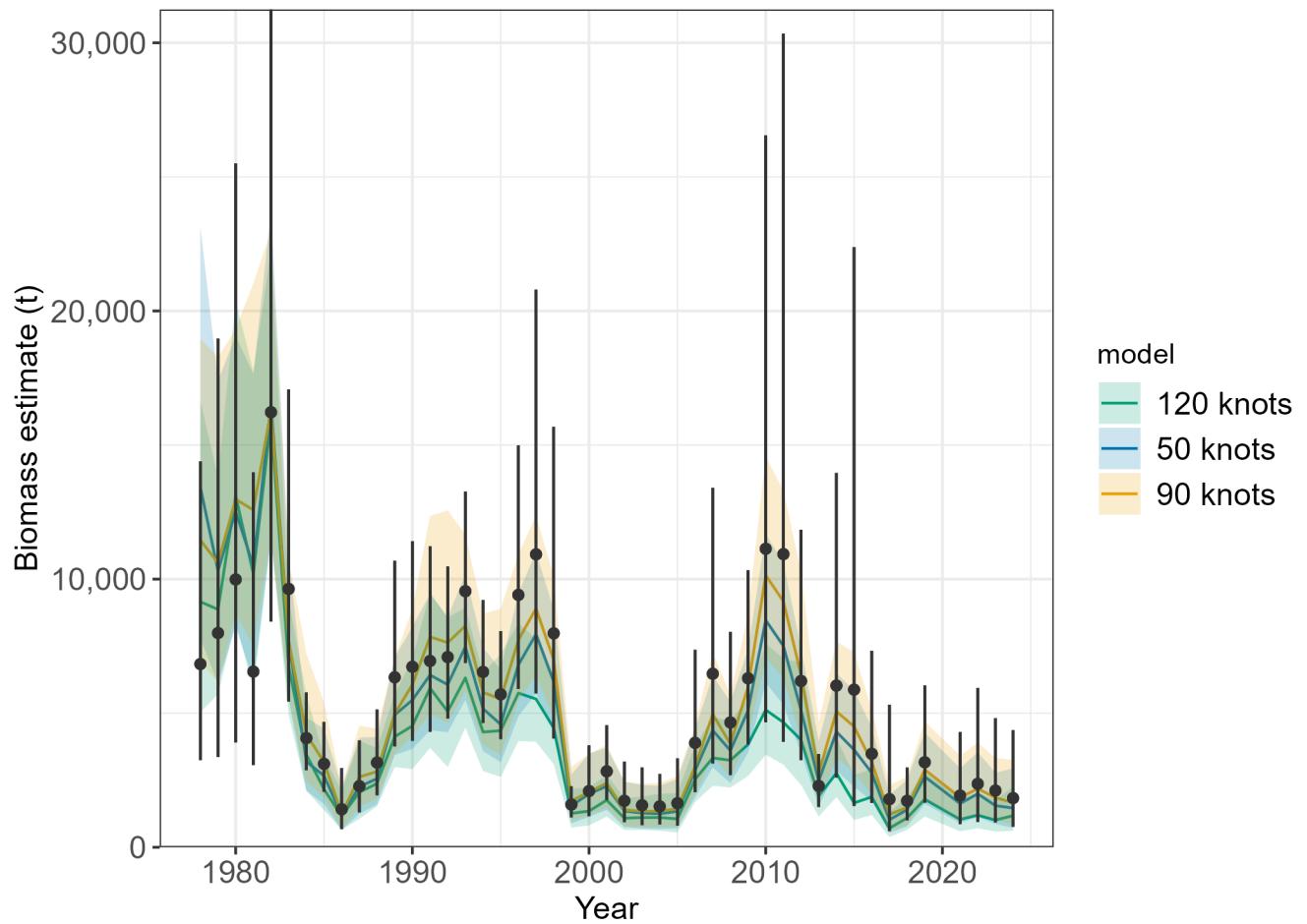


Figure 30: Estimated biomass (t) for St. Matthew Island blue king crab. Colored lines represent biomass ( $\pm 95\%$  CI) estimated using sdmTMB. Black points represent biomass ( $\pm 95\%$  CI) estimated by the NMFS EBS bottom trawl survey.