

A continued investigation into the emergence of a parasite in sea scallops

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Affected scallops

- Typical gross appearance and intensity of affected scallops.
- Reports from industry concerning infected scallops began in May of 2015 and have continued throughout 2016.
- Reports of infected scallops began in the DMV and have extended into the ET.



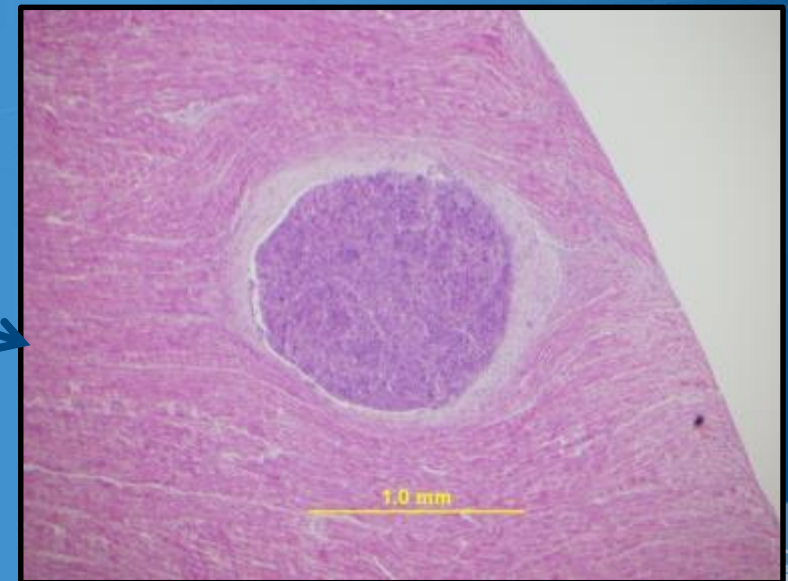
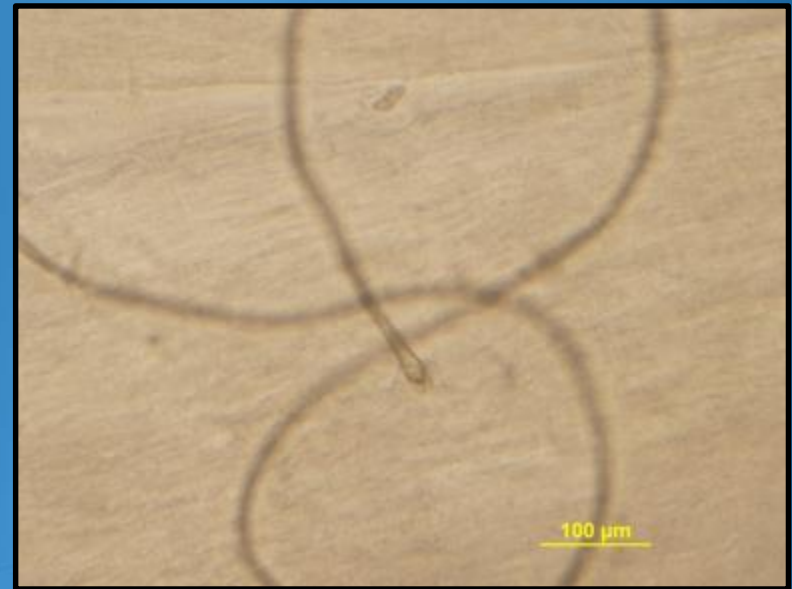
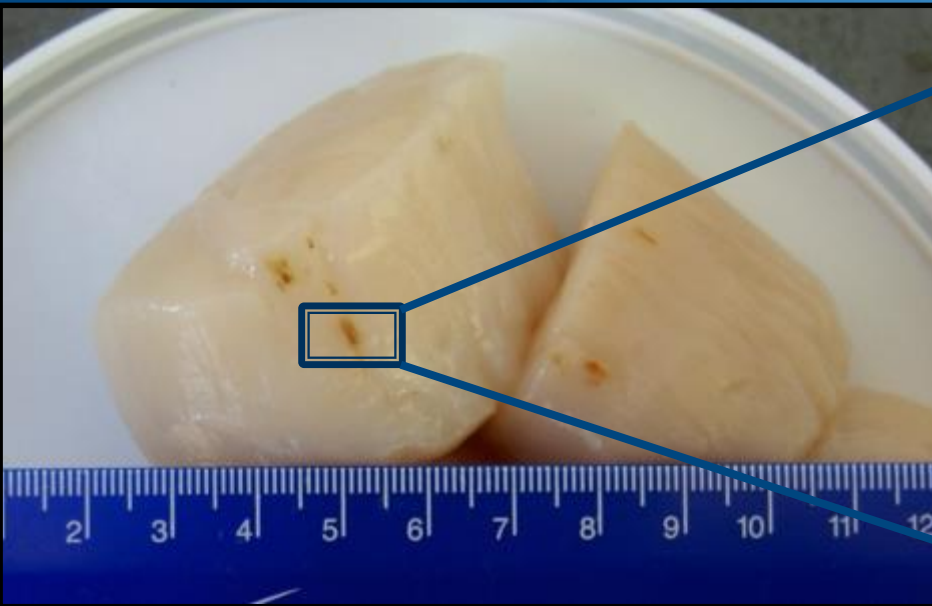
Appearance of affected scallops



- Typical lesion size with number per scallop meat ranging from 1-6.
- The lesions presented on the exterior of the adductor muscle, typically opposite the sweet meat.
- Visible to the naked eye against the white meat. (~2-5mm)

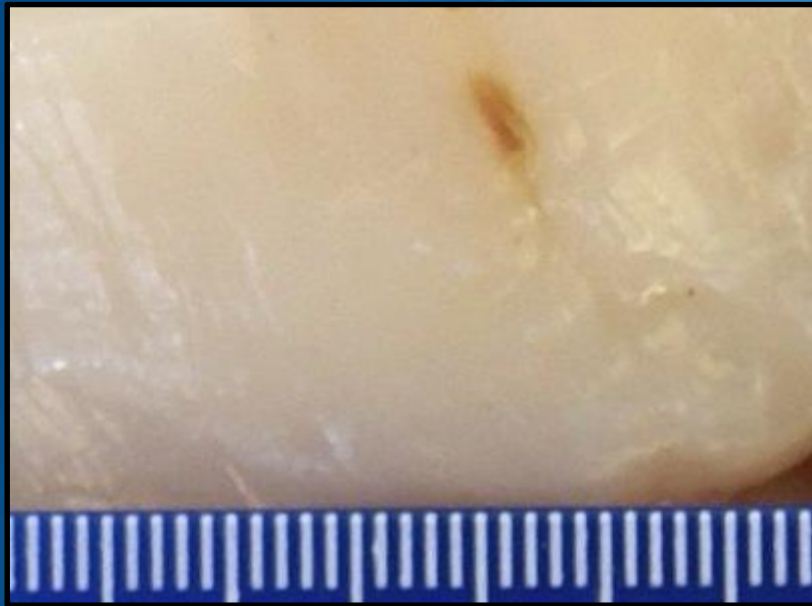
Preliminary histology

Fresh squash mount



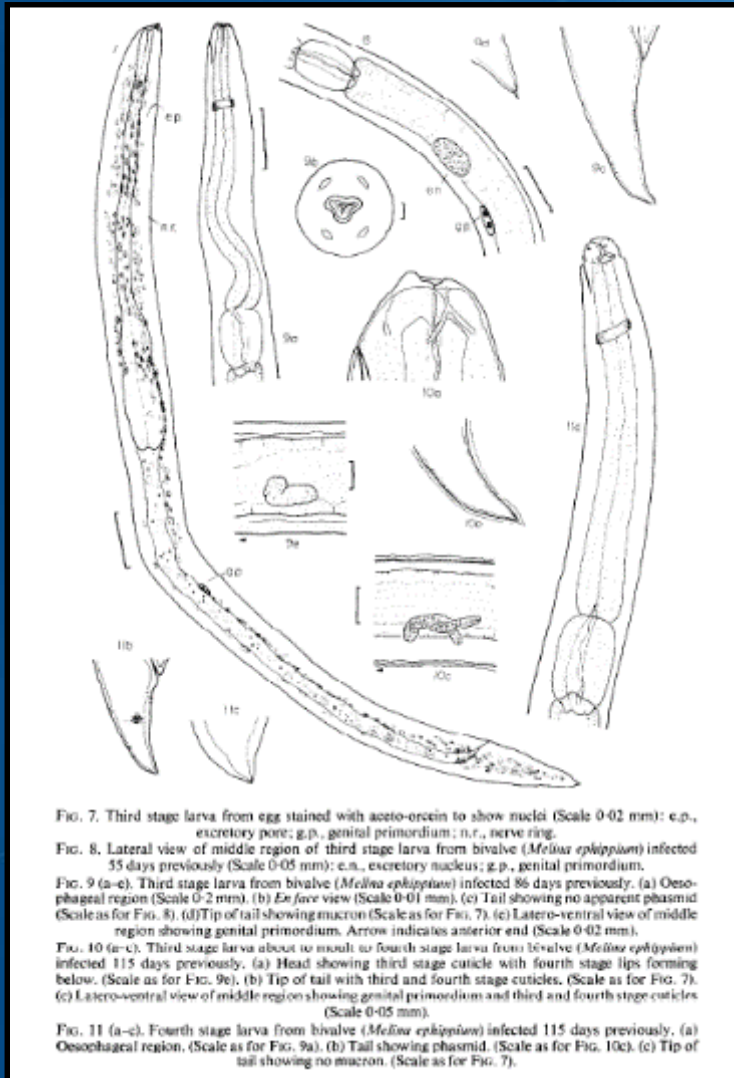
Histologically processed: pink=muscle,
blue=hemocytes surrounding foreign object
(host response)

Preliminary histology



Fourth stage nematode larvae coiled within brownish lesion in sea scallop adductor muscle. Some of the brown coloration may be the result of a *Haplosporidium* spp. hyperparasite

Preliminary identification

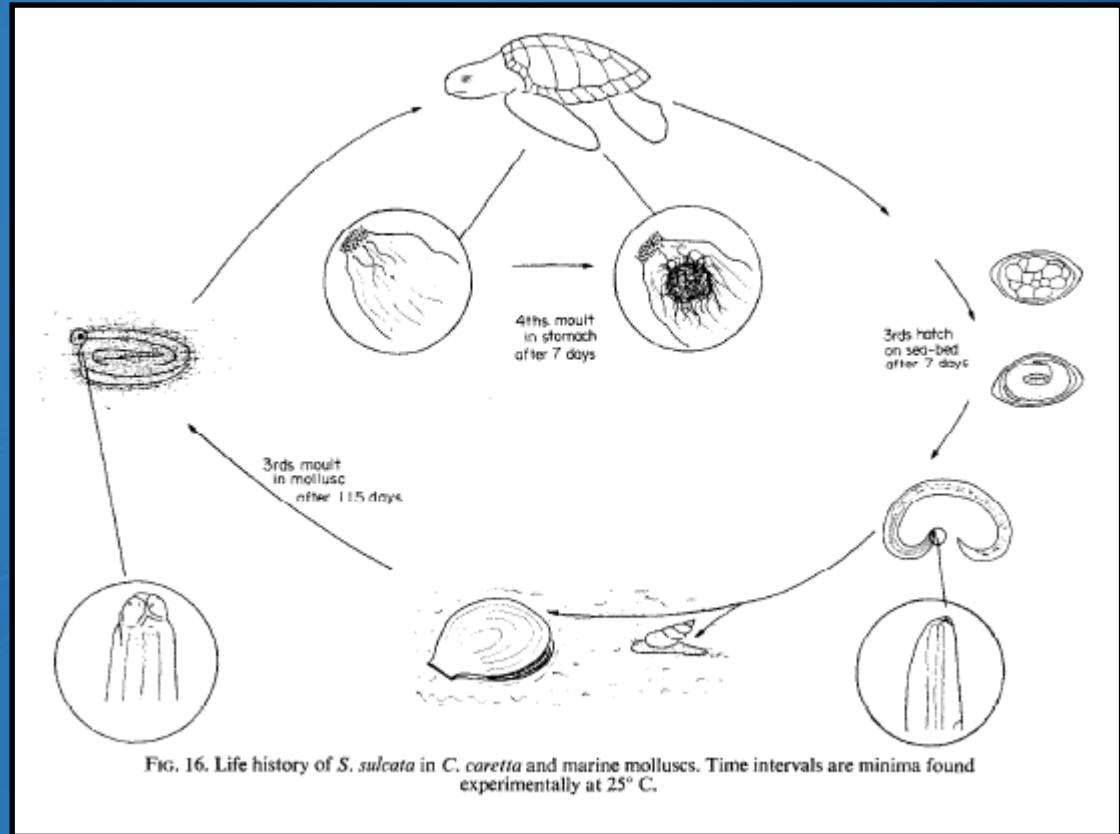


- Preliminary histology and molecular analysis has been completed on samples taken from the 2015 MAB survey.
- DNA results concluded that the sequences analyzed have a 99% identity with *Sulcascaris sulcata* .
- This species is cosmopolitan and has been identified in many genera of bivalve molluscs.
- Saucer scallop (Aus.), Calico scallop (US), Surf clams (US).
- Similar ephemeral observation of similar affected sea scallops was reported in May 2003.

From Berry and Cannon, 1981

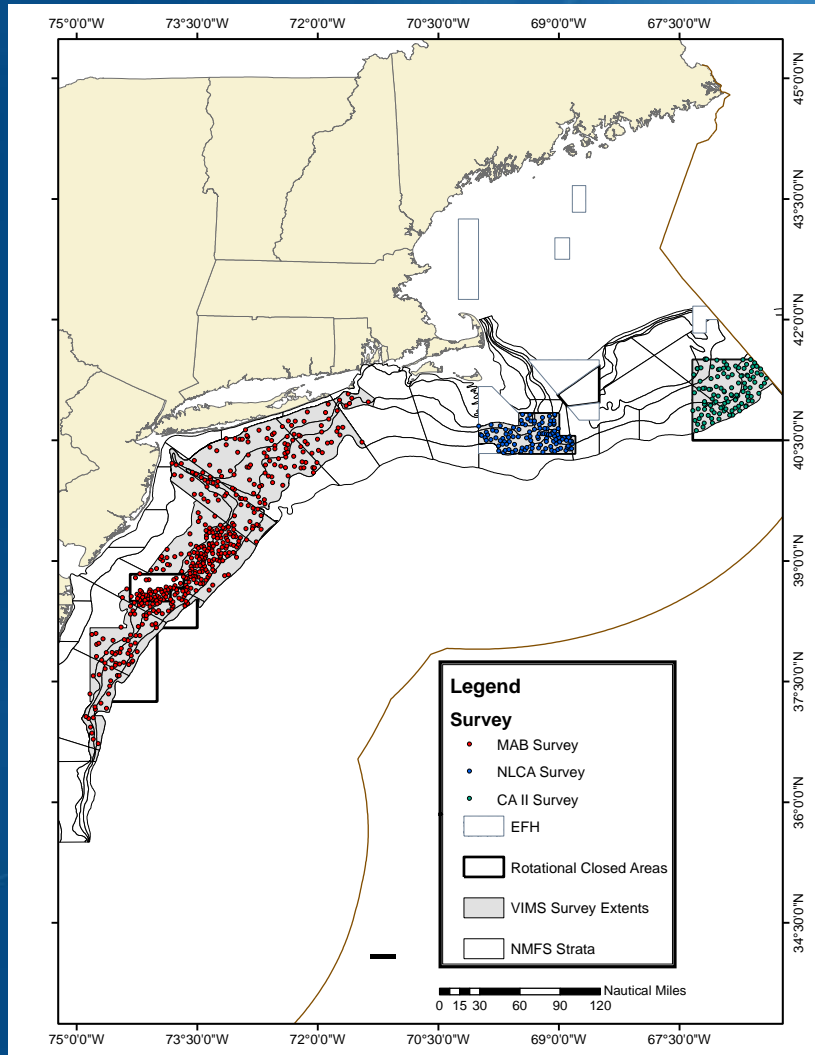
Sulcascaris sulcata life cycle

- The life cycle of *Sulcascaris sulcata* involves two hosts.
- Adult nematodes attach to the esophagus of Loggerhead and Green sea turtles.
- Eggs pass through the GI tract and enter the benthos via the feces.
- Eggs are filtered by benthic molluscs and the larval stages (1-4) develop.
- Fourth stage larvae are ingested by turtles.



From Berry and Cannon, 1981

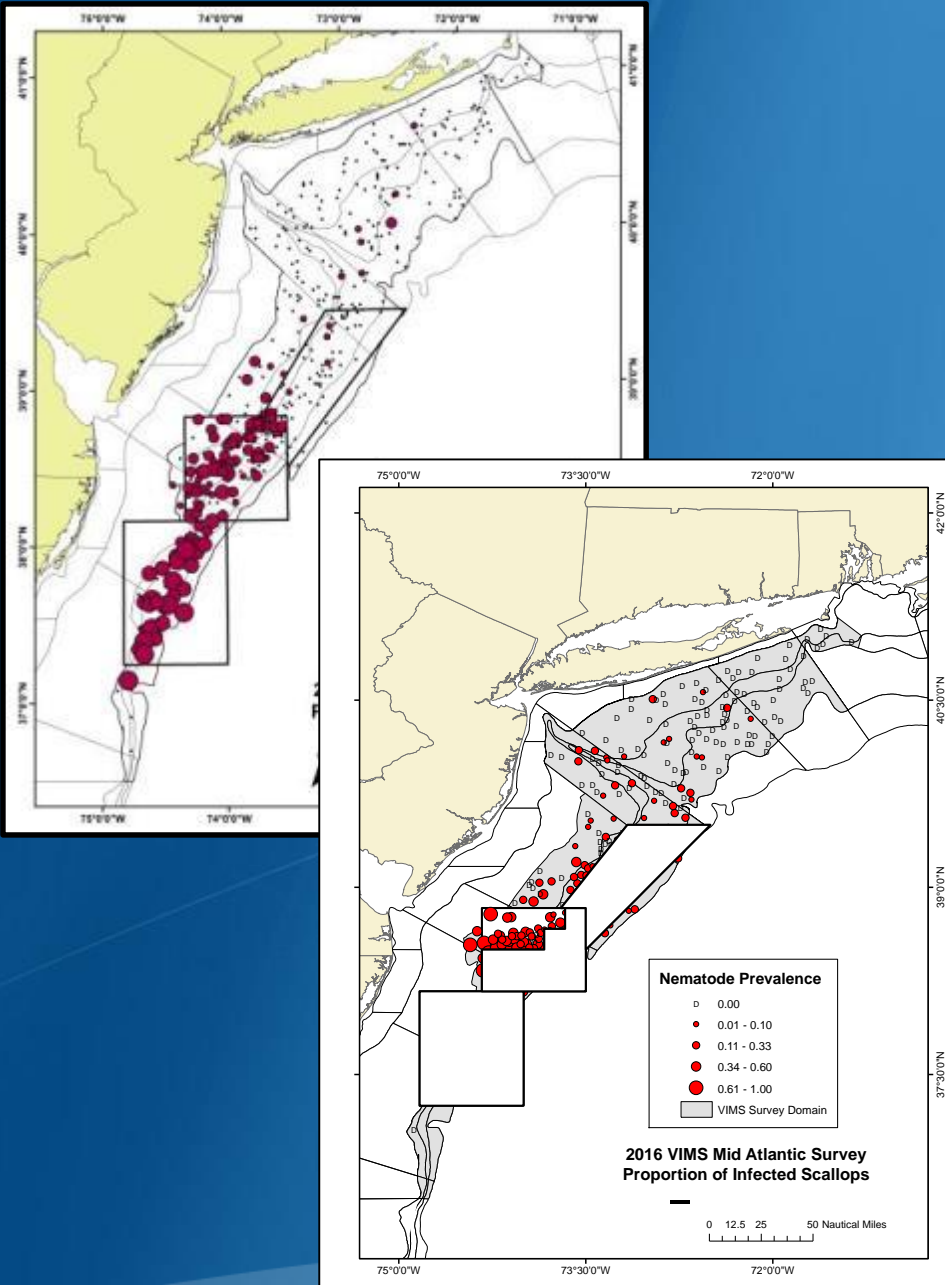
Parasite surveillance



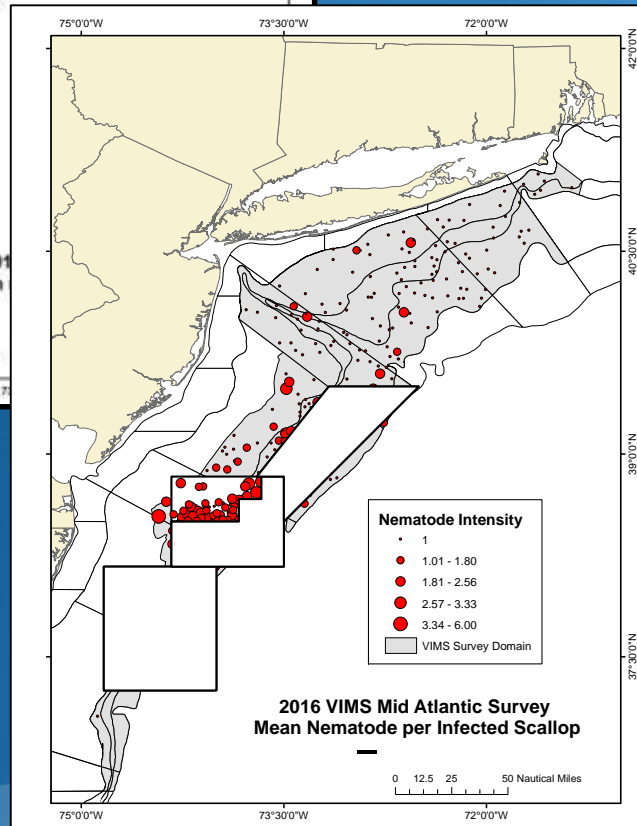
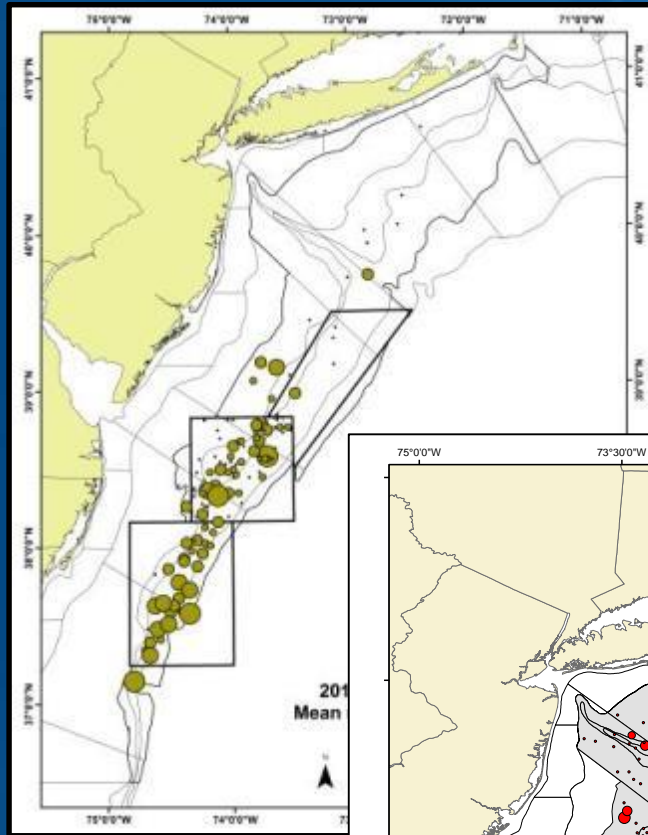
- For the 2016 surveys, VIMS continued an expanded biological sampling protocol to capture the spatial extent of the parasite as well as the prevalence and intensity of infected scallops.
- Sampled 15 animals at every station that had scallops .
 - Histological and genetic samples.
 - Gross observation of the number of infected scallops/sample (prevalence).
 - Gross observation of the number of nematodes/scallop (intensity).

Nematode Prevalence

- Spatial distribution of the prevalence of the parasite in the sampled scallops.
- No infected scallops observed in the NLCA or the CA II survey areas.
- For each station with sampled scallops, a proportion of the sample that contained at least one nematode was calculated.
- Intensity appears to increase as a function of decreasing latitude.
- Prevalence appears to be increasing in the ET and Hudson Canyon compared to 2015.



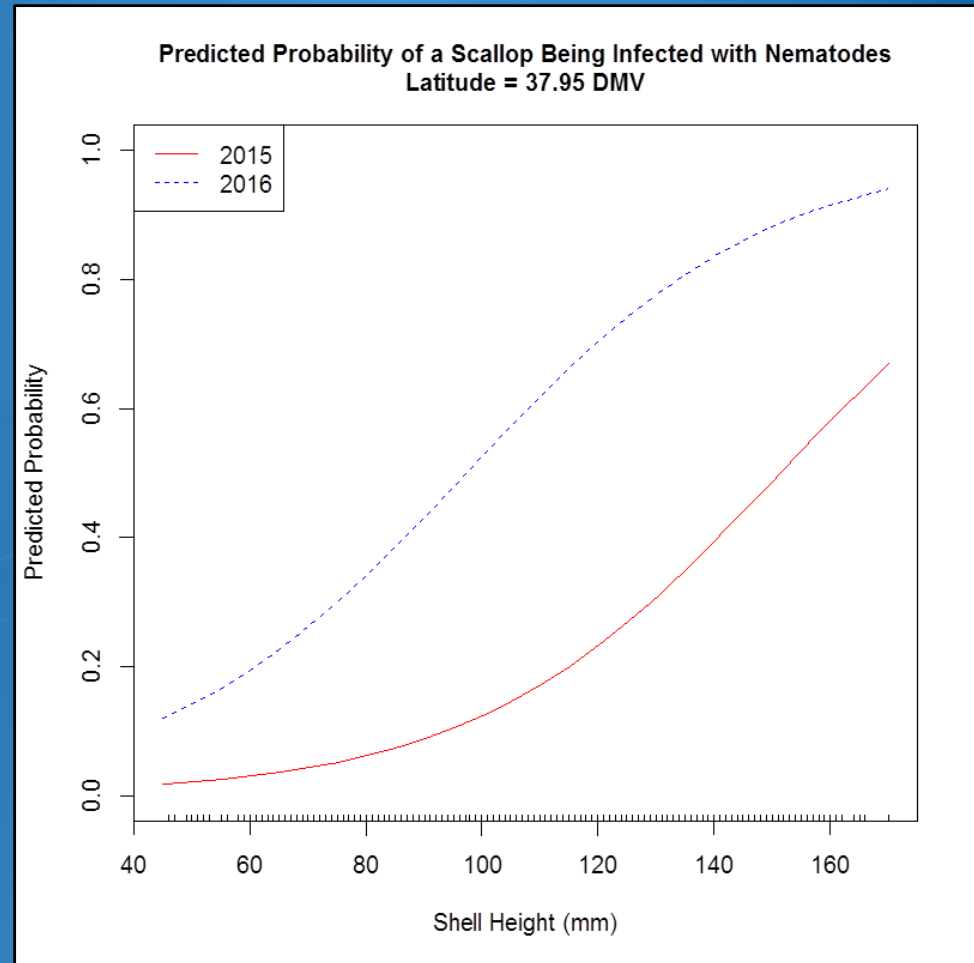
Nematode Intensity



- Spatial distribution of the intensity of the parasite in the sampled scallops.
- For each positive identification at a given station, the mean number of nematodes per scallop was calculated.
- Intensity appears to increase as a function of decreasing latitude.
- Intensity in northern areas is also increasing compared to 2015.

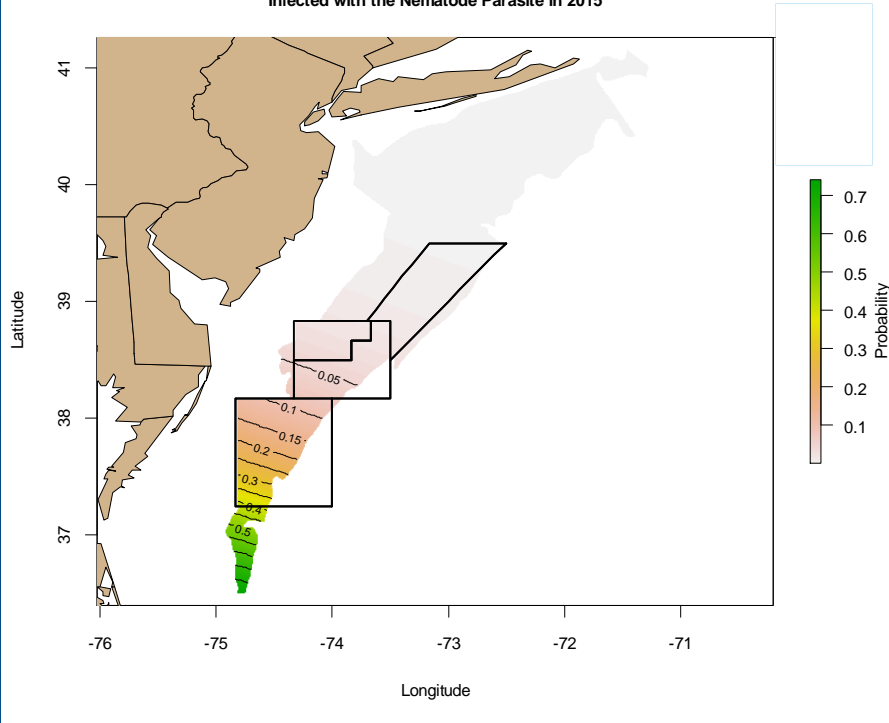
Logistic Regression for Nematode Presence

- Logistic regression was developed to predict the probability of a scallop being infected with nematodes.
- Significant predictor variables included year (2015 & 2016), latitude & shell height.
- The overall probability of a scallop being infected increased as a function of shell height and decreased as a function of latitude.
- All else being equal, $P(2016) > P(2015)$
- Predictions are sensitive to the latitude value used.

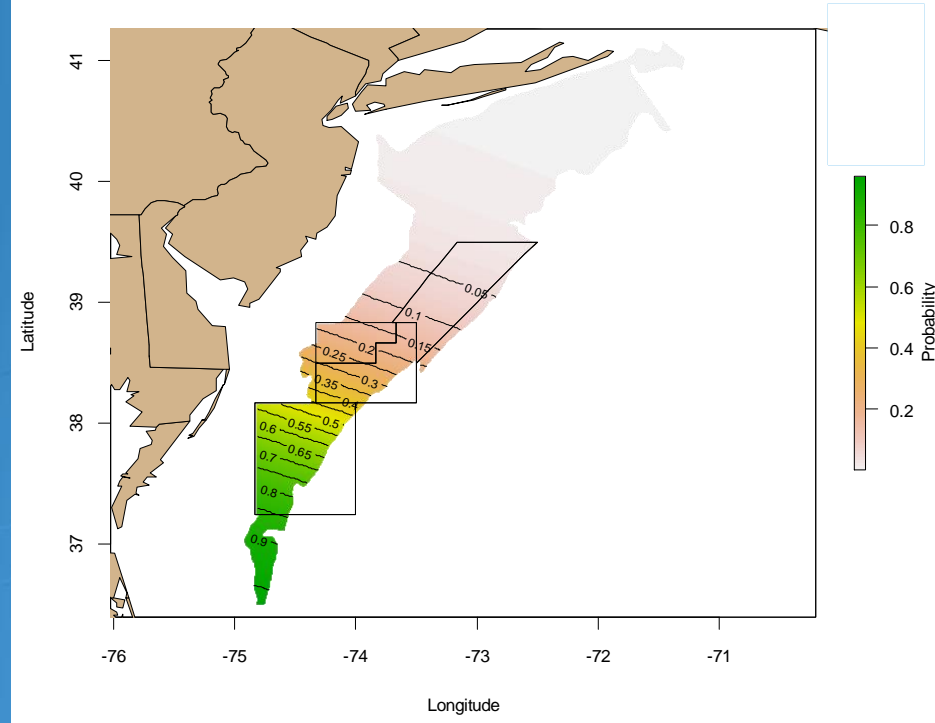


GAMM for Nematode Presence

Predicted Probability of a 100 mm Scallop Being Infected with the Nematode Parasite in 2015



Predicted Probability of a 100 mm Scallop Being Infected with the Nematode Parasite in 2016



- GAMM was developed to predict the probability of a scallop being infected with nematodes.
- Significant predictor variables included year (2015 & 2016), tensor product of latitude & longitude & shell height.

Summary

- One parasite species (*S. sulcata*) has been positively identified using taxonomic and genetic techniques.
- Samples collected during the 2016 MAB survey will be further analyzed using histology and molecular approaches.
 - Same species of nematode
 - Population structure of nematode.
- Additional work being done to understand transmission as well as spatio-temporal dynamics of the nematode.
- Impact on fishery.
 - Clear overlap with the core of the current scallop biomass and the highest prevalence and intensity of the parasite.
 - In May of 2003, reports waned over time and there were no additional reported sightings until 2015.

