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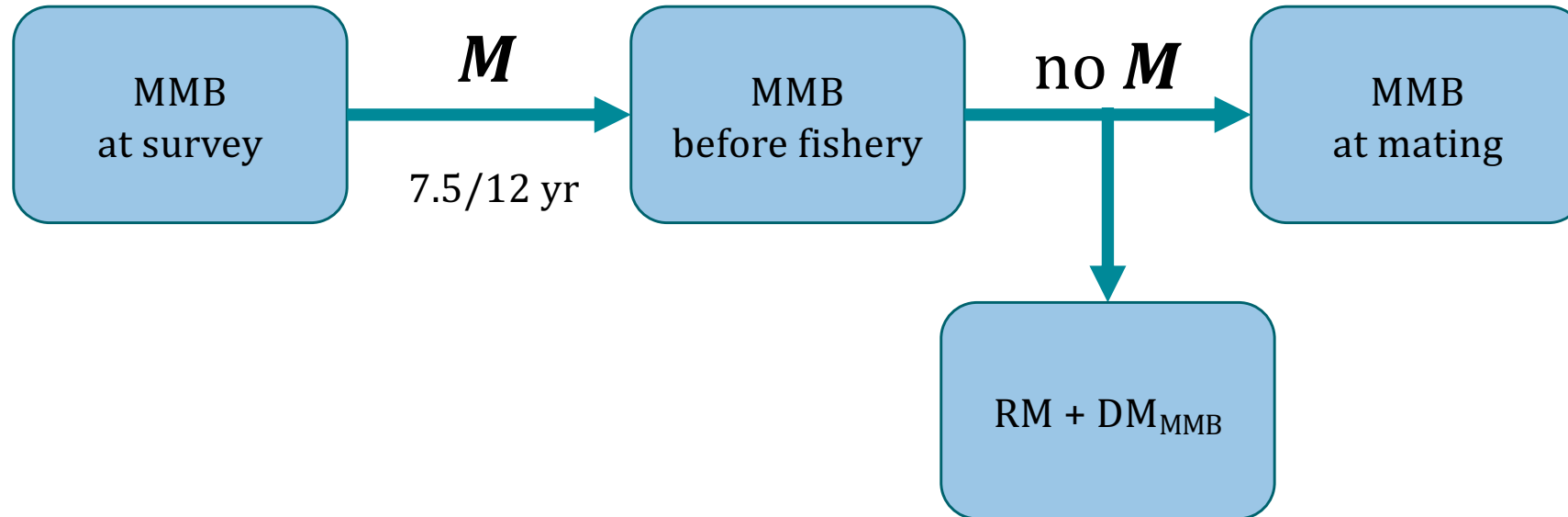
Pribilof Islands Blue King Crab Candidate Model for 2023 Assessment

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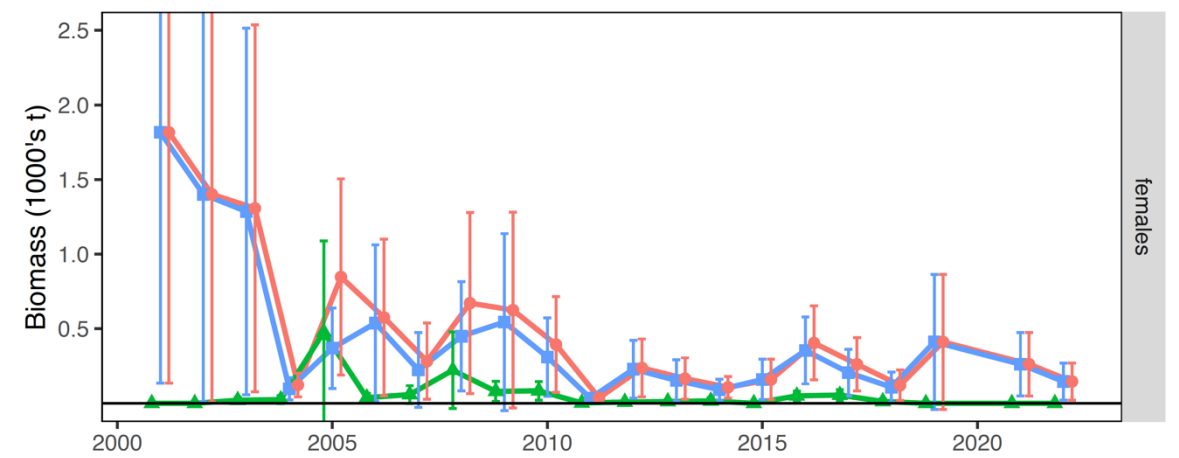
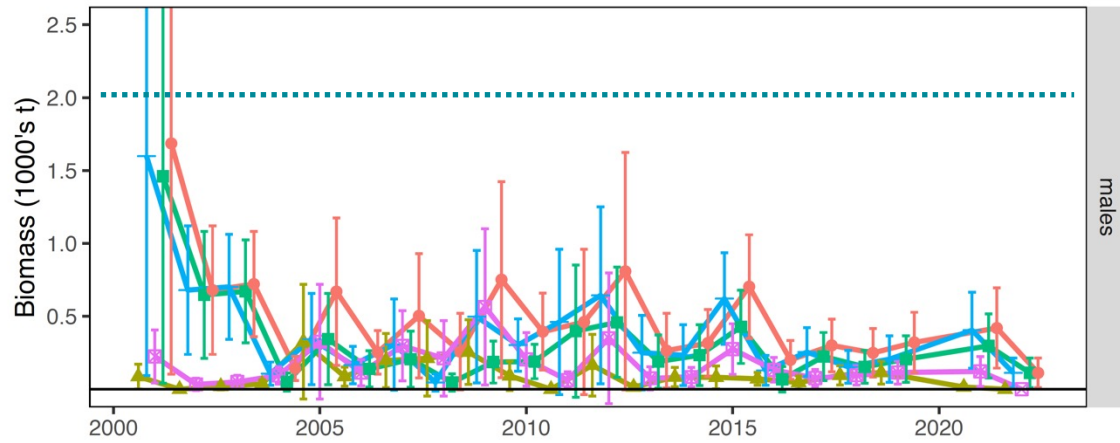
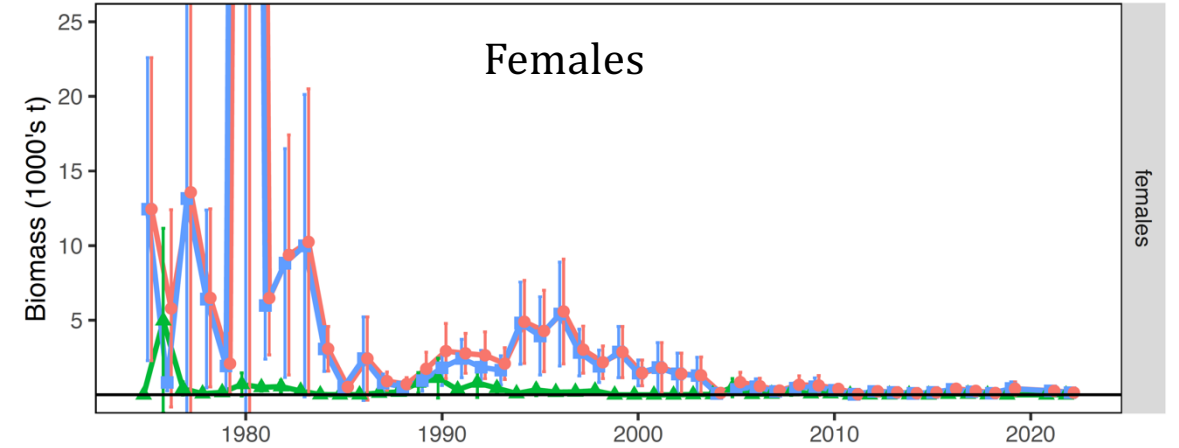
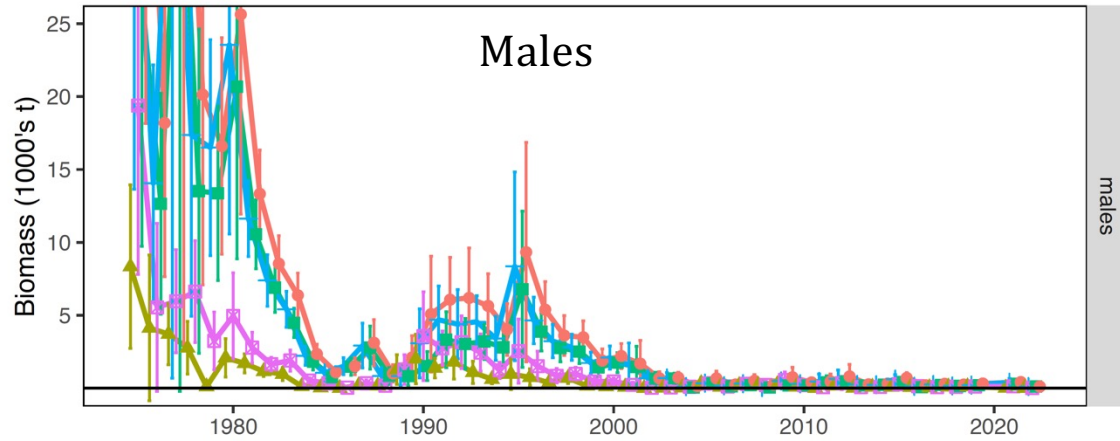
May 2023

Tier 4 Model for PIBKC Status Determination



- B_{MSY} = average MMB-at-mating
- status ratio = current MMB-at-mating/ B_{MSY}

Survey Trends



● all males
 ▲ immature males
 ■ legal males
 + mature males
 ◆ sublegal males

● all females
 ▲ immature females
 ■ mature females

Issues

- Design-based time series of survey MMB has large CVs
- Issues with VAST estimates given limited spatial size and island effects?
- Current approach: apply state-space random walk model to design-based estimates of MMB to capture time series trend with smaller CV's
 - “bespoke” ADMB code
- Proposed modification: use *rema** R package to fit same model
 - developed for groundfish Tier 5 assessments
 - uses TMB rather than ADMB
 - larger user community for future development

*<https://github.com/JaneSullivan-NOAA/rema>
<https://afscassessments.github.io/rema/>



State-Space Random Walk Model

Observation model

$$\ln(MMB_{s_y}) = \langle \ln(MMB_s) \rangle_y + \eta_y, \text{ where } \eta_y \sim N(0, \sigma_{s_y}^2)$$

Process model for unobserved state ()

$$p(\langle \ln(MMB_s) \rangle_y | \langle \ln(MMB_s) \rangle_{y-1}) \sim N(0, \phi^2) \quad (\text{random walk})$$

Joint likelihood

$$\Lambda = \sum_y \left[\ln(2\pi\phi) + \left(\frac{\langle \ln(MMB_s) \rangle_y - \langle \ln(MMB_s) \rangle_{y-1}}{\phi} \right)^2 \right] + \sum_y \left(\frac{\ln(MMB_{s_y}) - \langle \ln(MMB_s) \rangle_y}{\sigma_{s_y}} \right)^2$$

Parameters

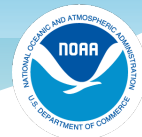
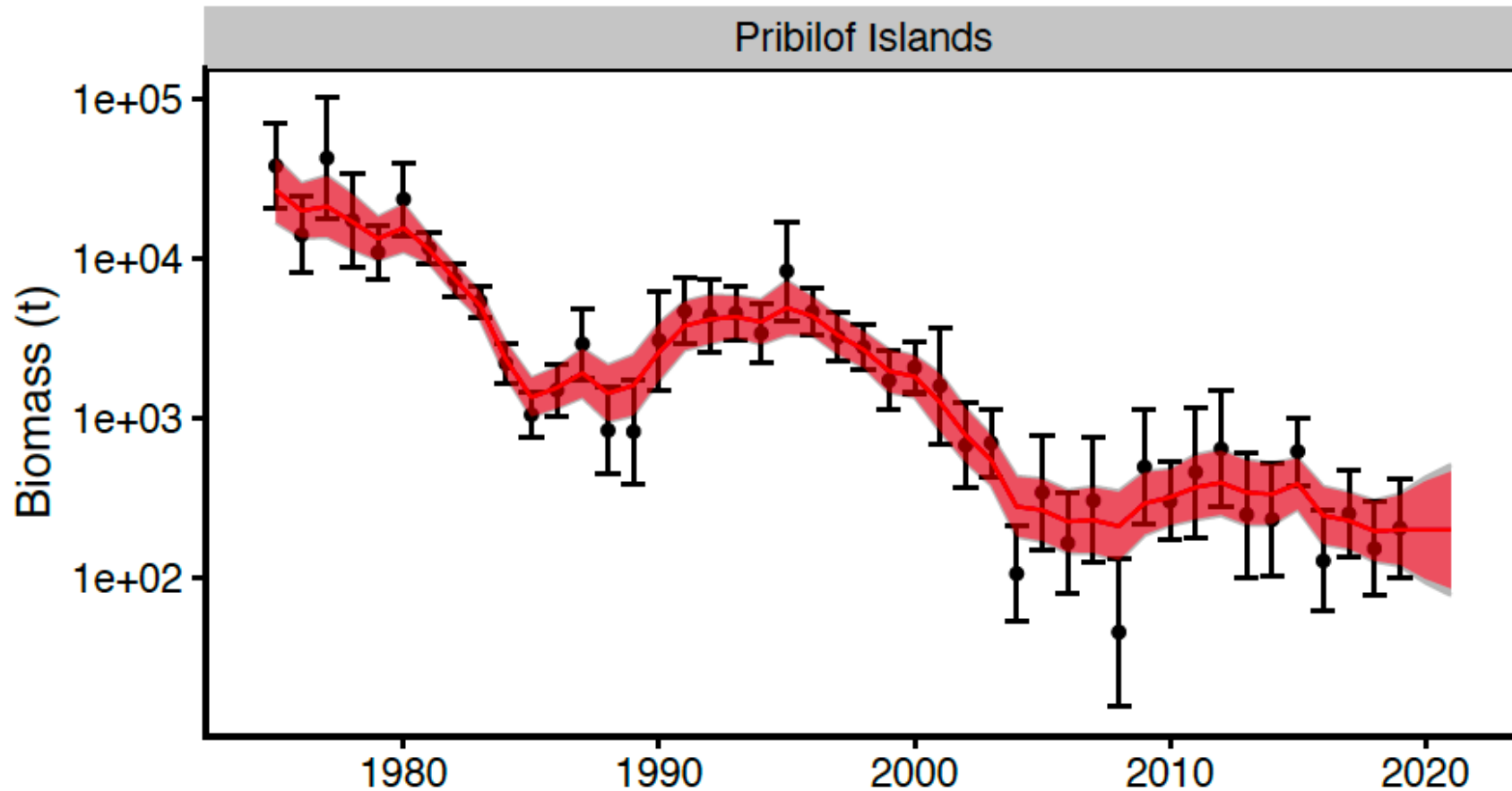
λ , where $\phi^2 = \exp(2 \cdot \lambda)$: fixed parameter characterizing scale of process error

$\langle \ln(MMB_s) \rangle_y$: random effects representing the state



Comparison between ADMB and rema

model	objective function	max gradient	estimate	std err
ADMB	28.3803	4.5486e-08	0.4332	0.0779
rema	28.3492	7.2188e-15	0.4331	0.0779



Discussion

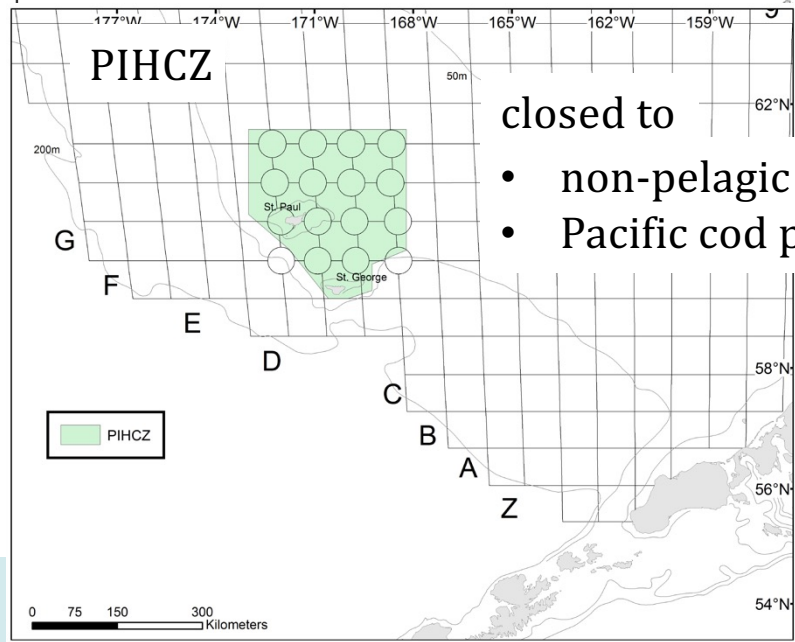
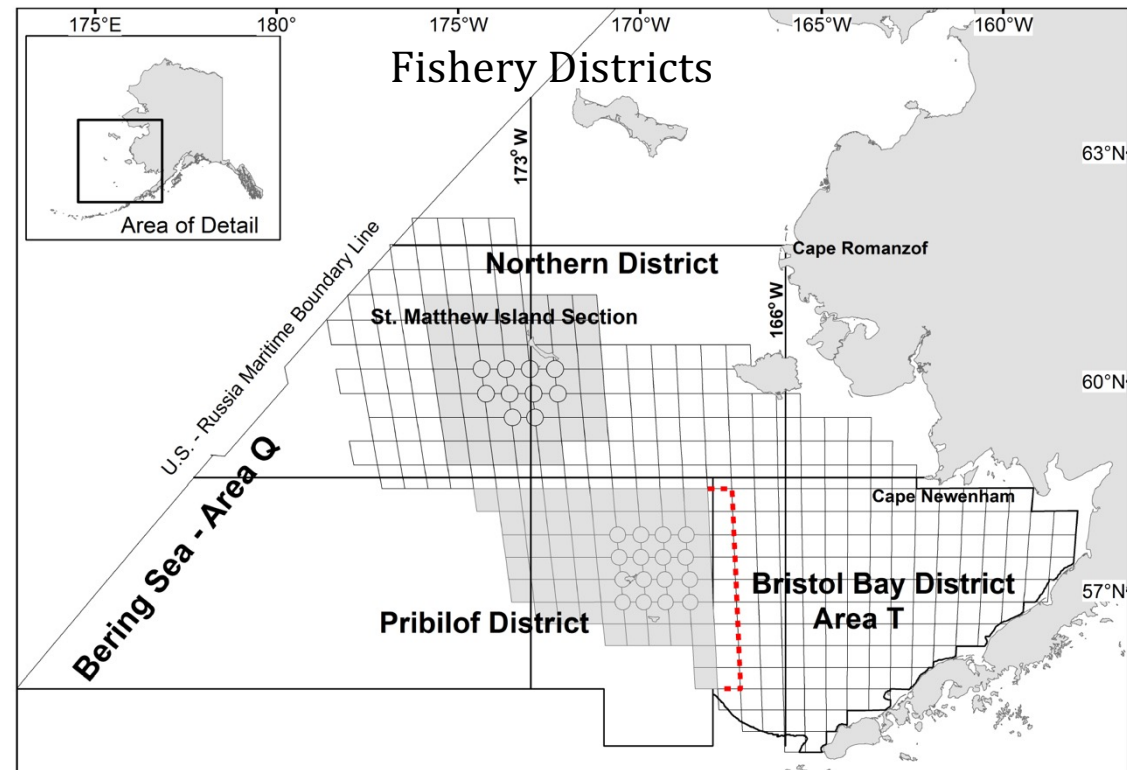
- Differences between ADMB and *rema* results are negligible
- *rema* package is standard model framework adopted for groundfish Tier 5 stock
 - TMB has better performance than ADMB for random effects models
 - larger user community
 - support for future development (diagnostics, etc.)
- Recommendation: adopt *rema* package in favor of ADMB code for PIBKC assessment
 - proposed is NOT a change in model structure





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Stock Distribution



- closed to
- non-pelagic trawl gear
 - Pacific cod pot gear

