

# Empirical dynamic modelling summary

## EDM group

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Government  
of Canada

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du Canada

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TESA ecosystem approach workshop, Nanaimo.  
Friday 25<sup>th</sup> November 2016

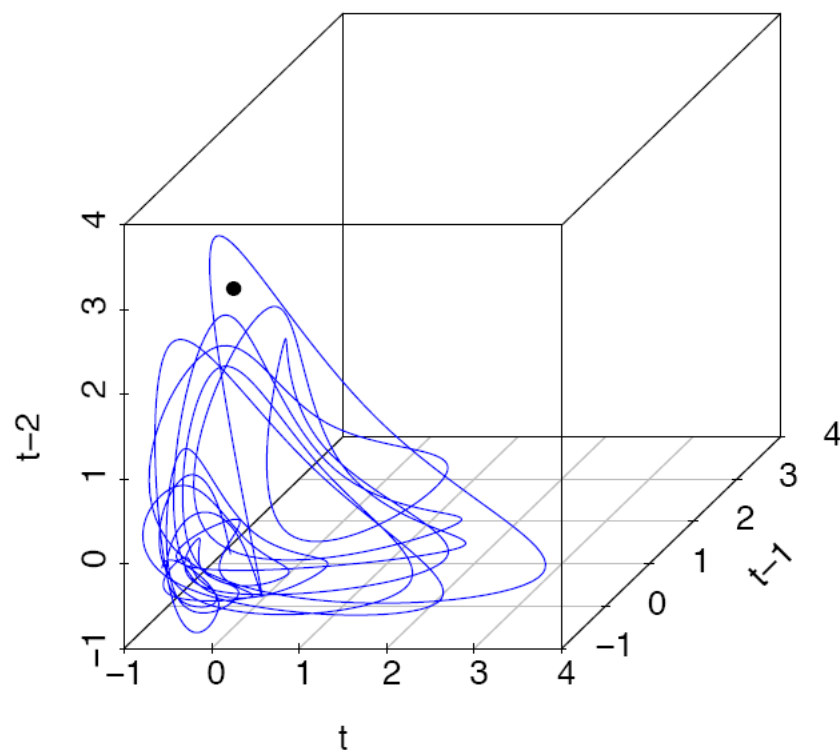
# Background

“Incorporating an ecosystem approach into single-species stock assessments”

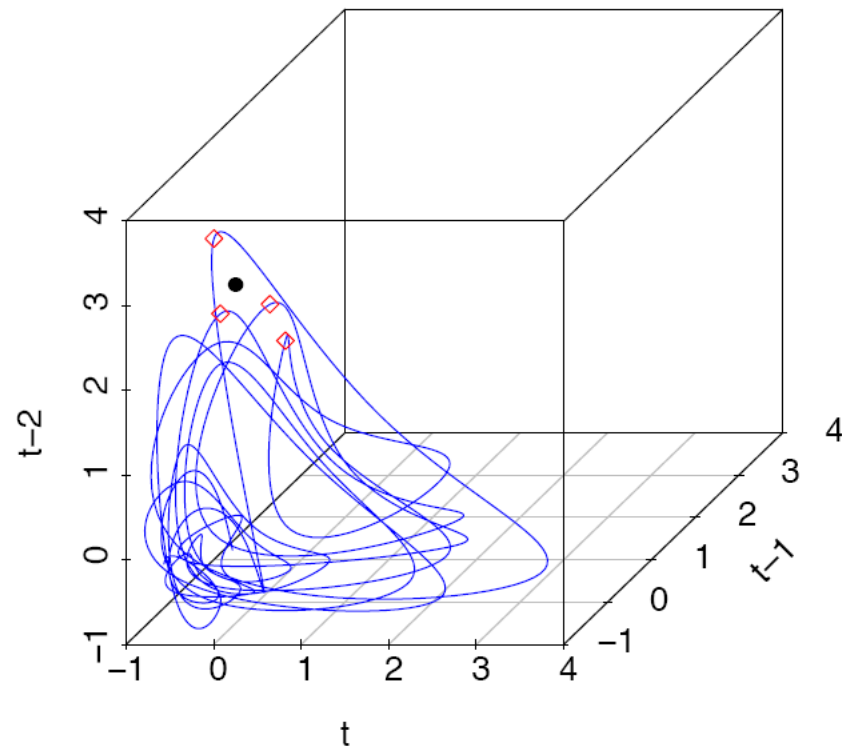
This is a different approach, more like:

“An approach for analysing ecosystem influences on single (or multiple) species.”

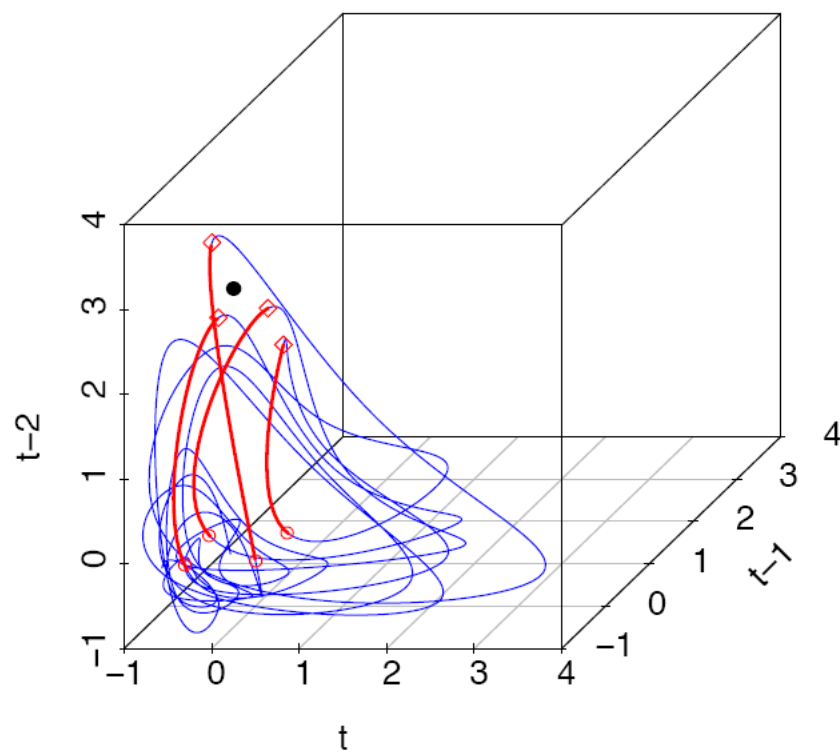
# Simplex Projection



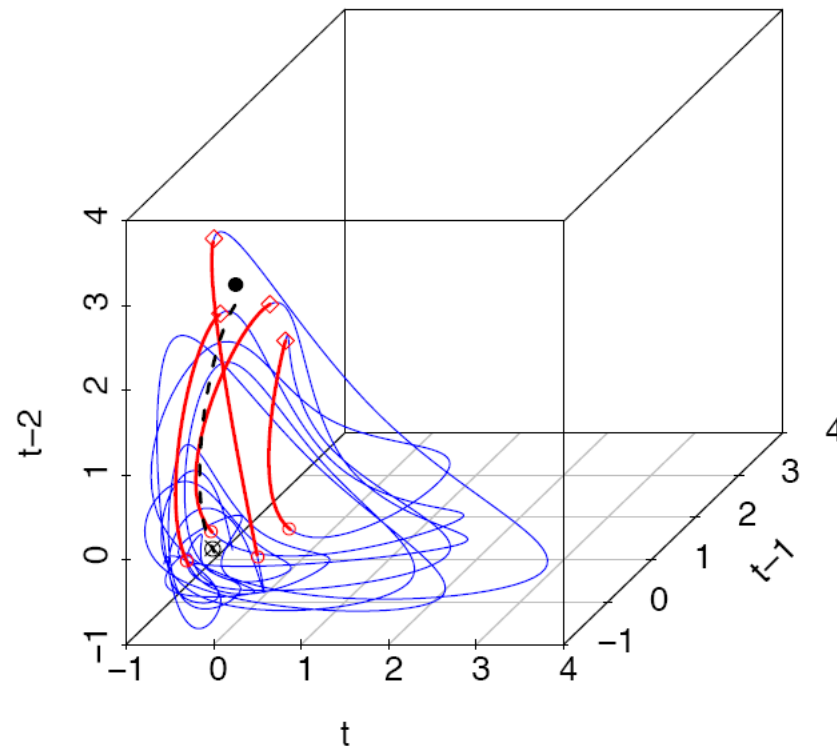
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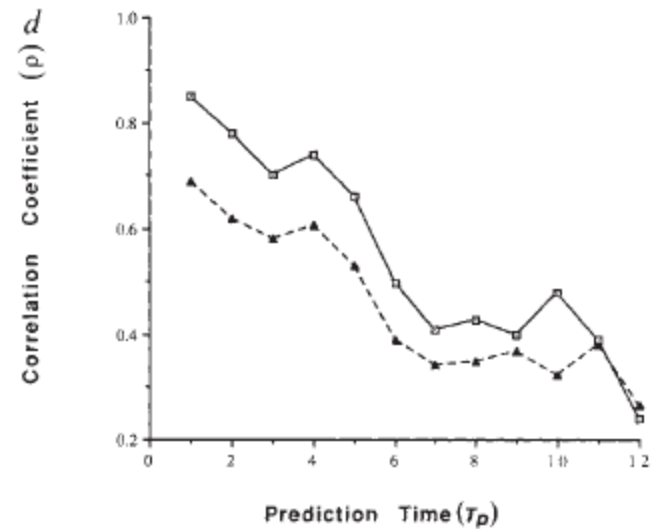
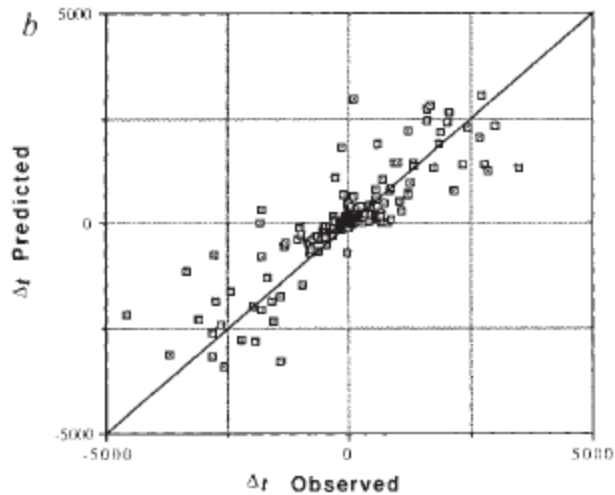
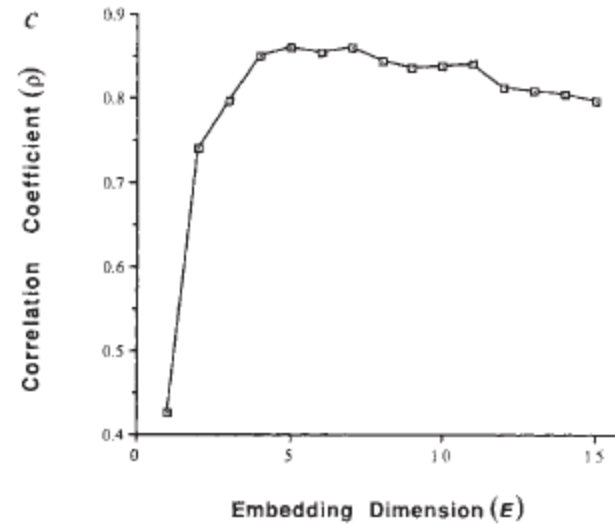
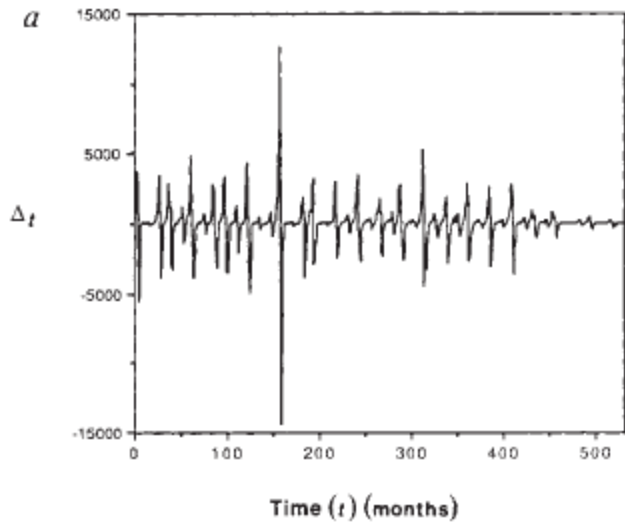
# Simplex Projection



# What we did

1. rEDM package – works well, we understand some of it (worked through some of vignette). Data input challenging.
2. Random data – gives nonsense. Deterministic model – does well.
3. Read paper – Sugihara and May (1990).
4. Salmon – tried to replicate some results.
5. Salmon – updated data set. Predictability similar, best models different.
6. Real data – crab data: R1, R2, R3.
7. Simulated competition model.

# Helpful paper



Sugihara and May, 1990, Nature, 344:734.

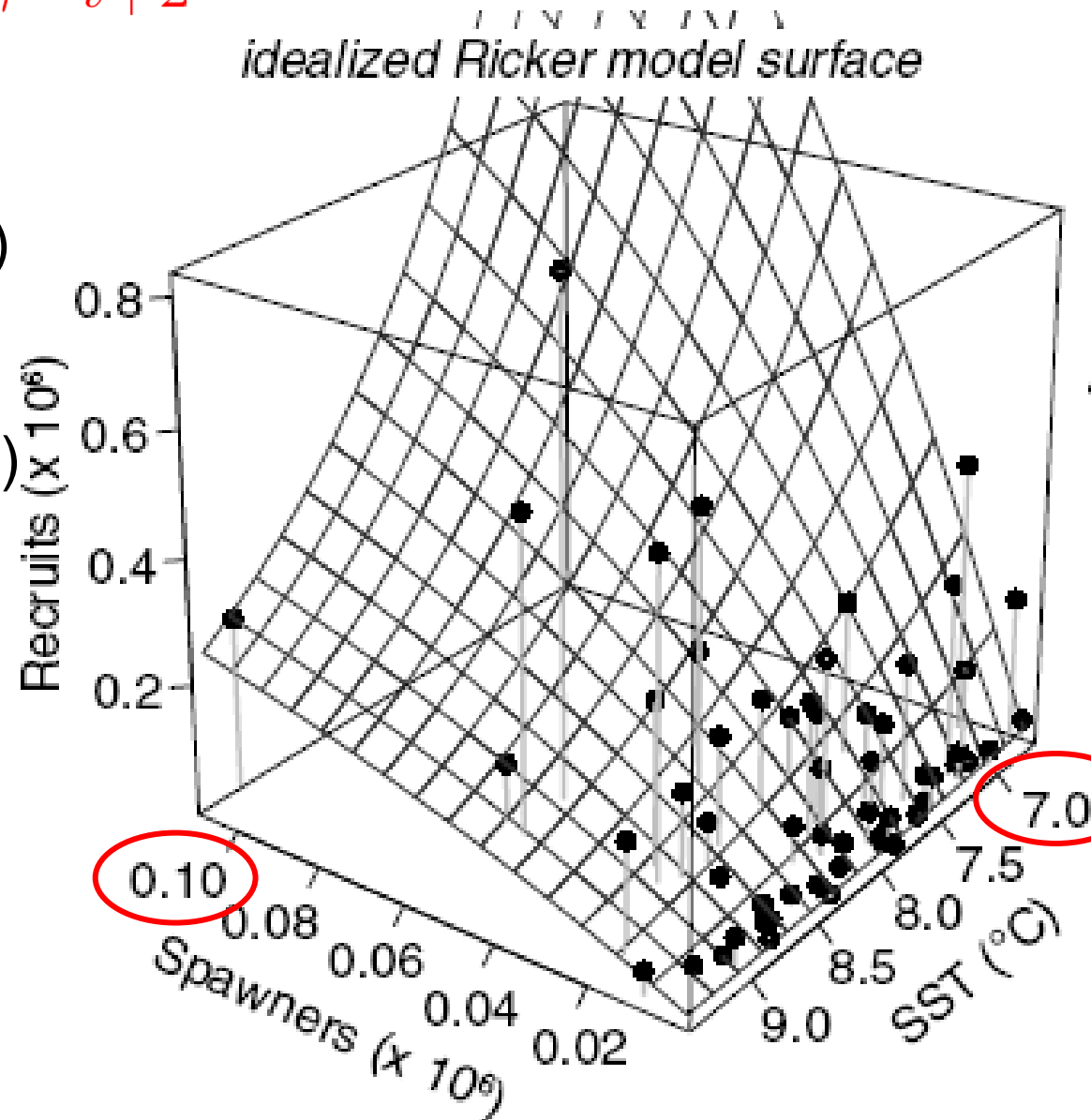


# Salmon – incorporate temperature

$$\hat{R}_t = S_t e^{\alpha - \beta S_t + \gamma U_{t+2}}$$

Low (but attainable) SST, with high (but attainable) spawners, predicts ridiculously high (historically unattainable) recruits.

Prescribing above equation gives only one hypothesis for how temperature can affect recruitment.



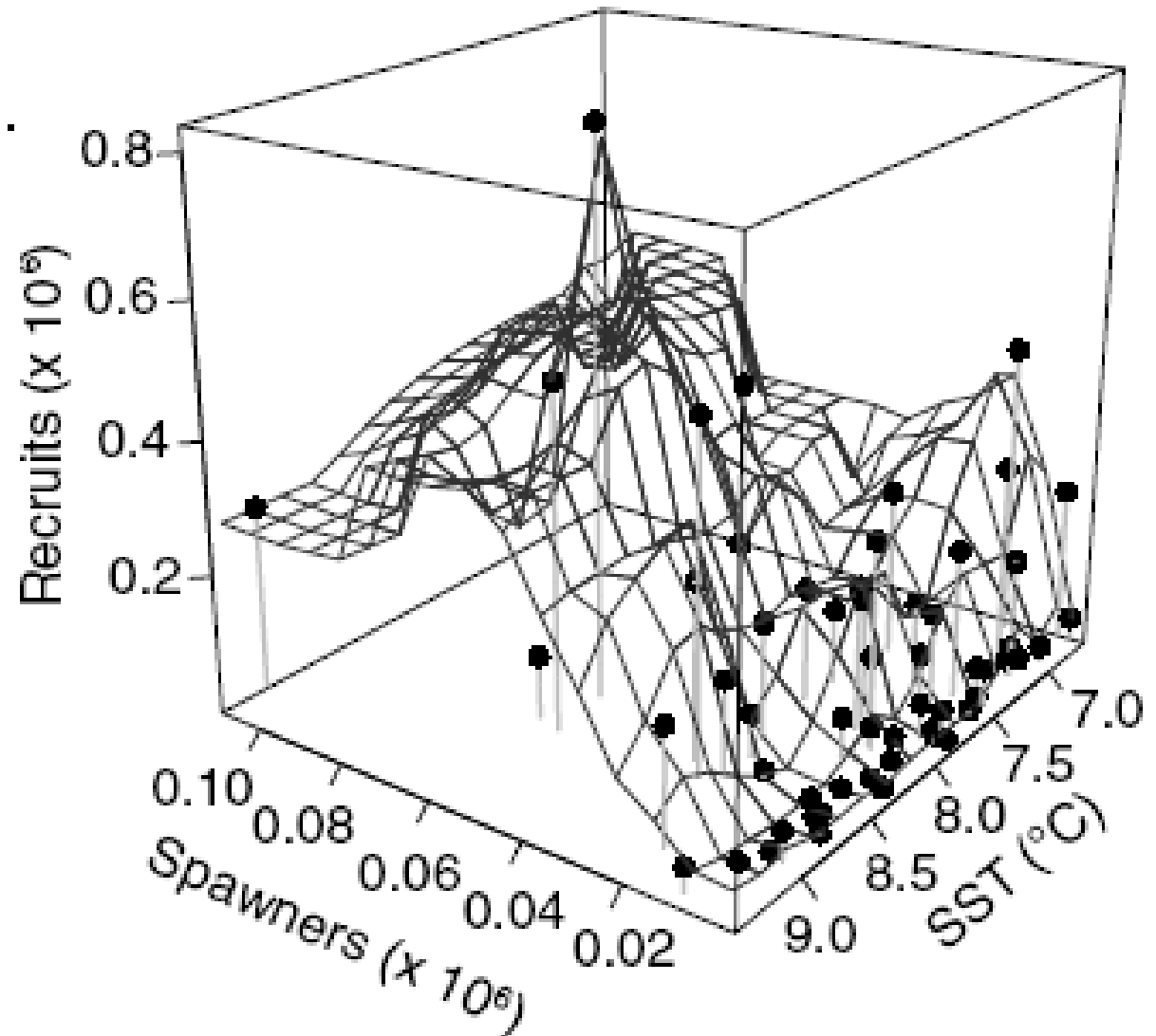
# EDM approach

*empirical (non-parametric) model surface*

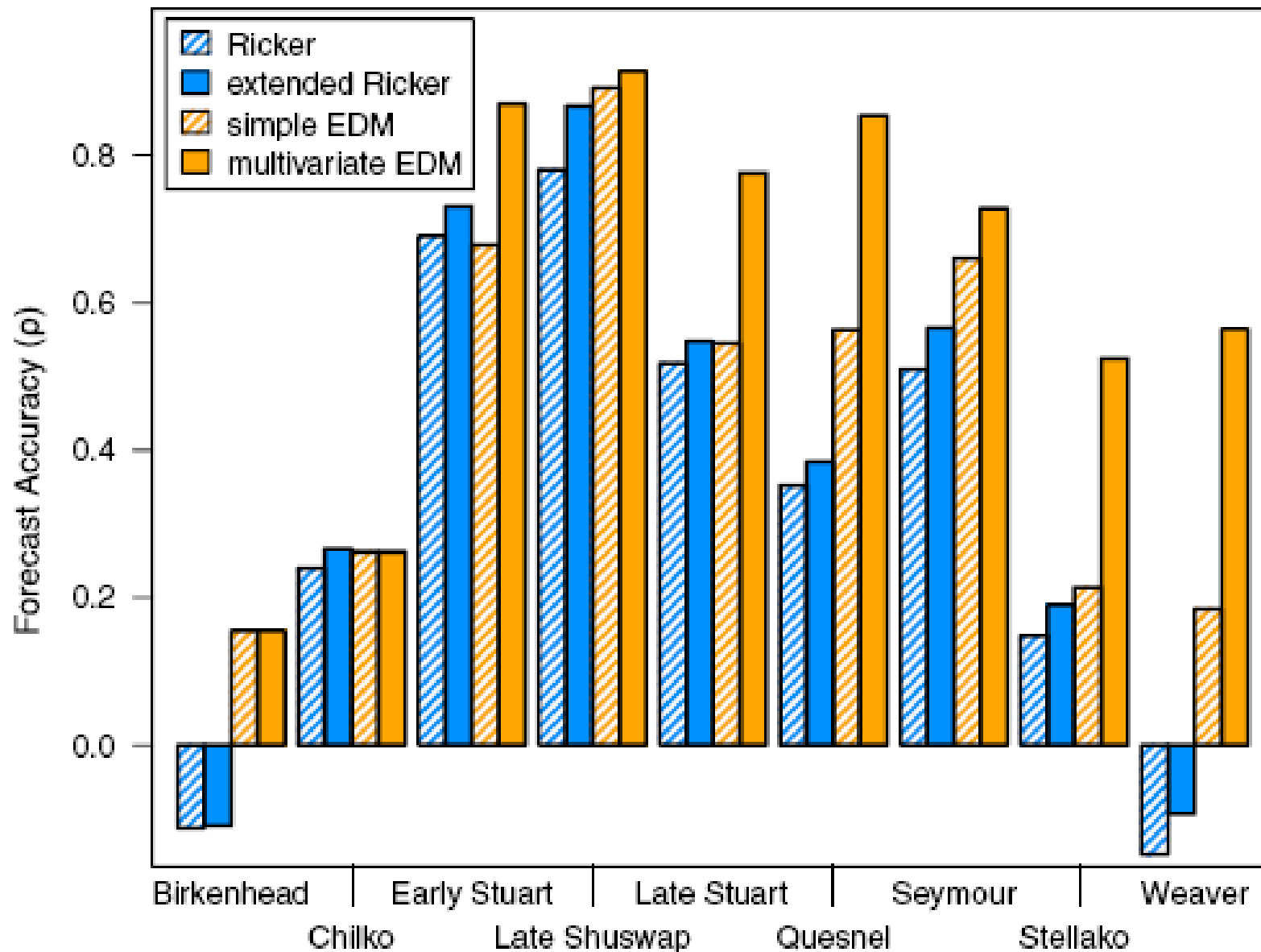
No equation specified.

No hypothesised  
relationship.

Just uses data.

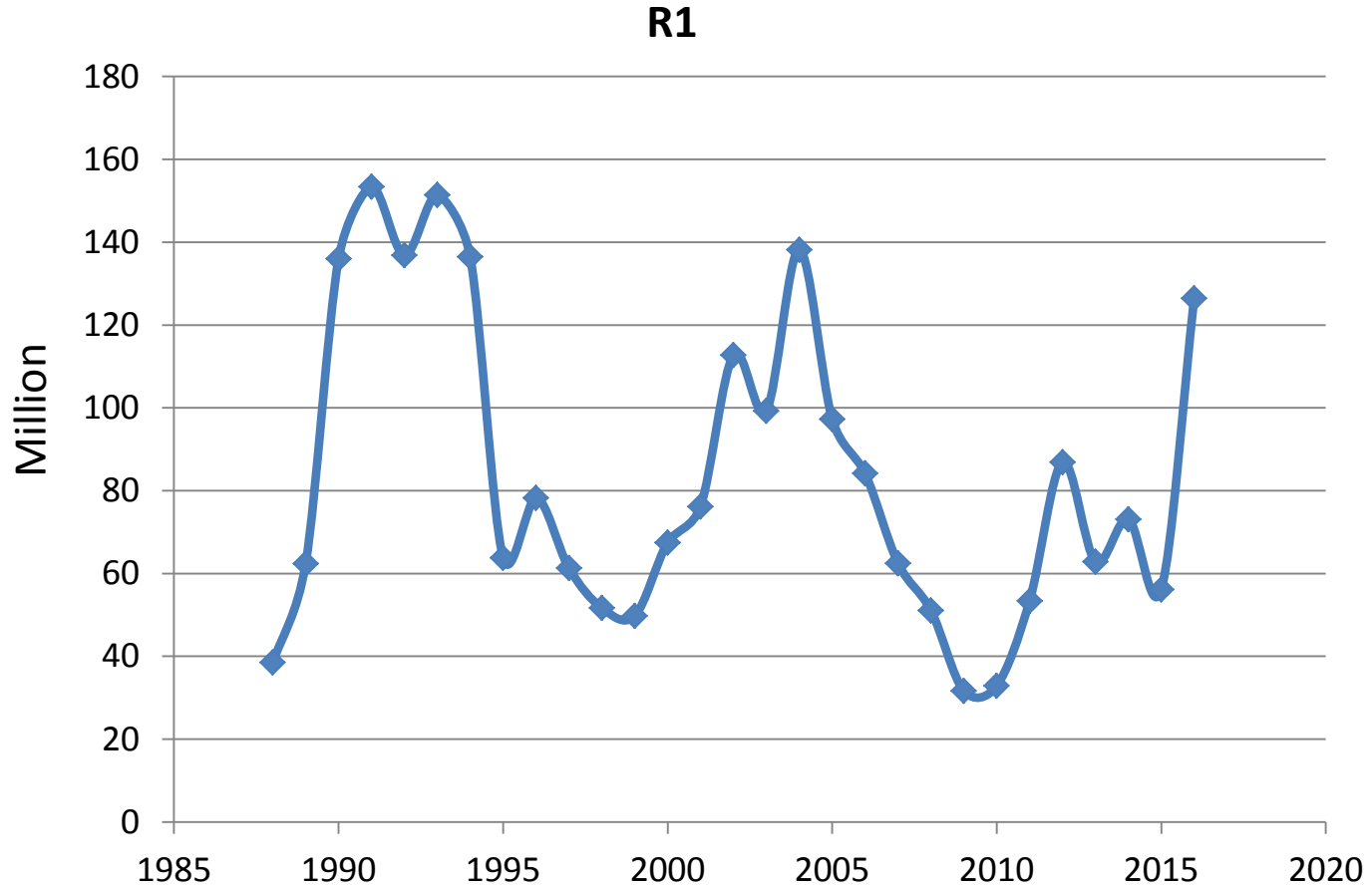


# Forecast accuracy

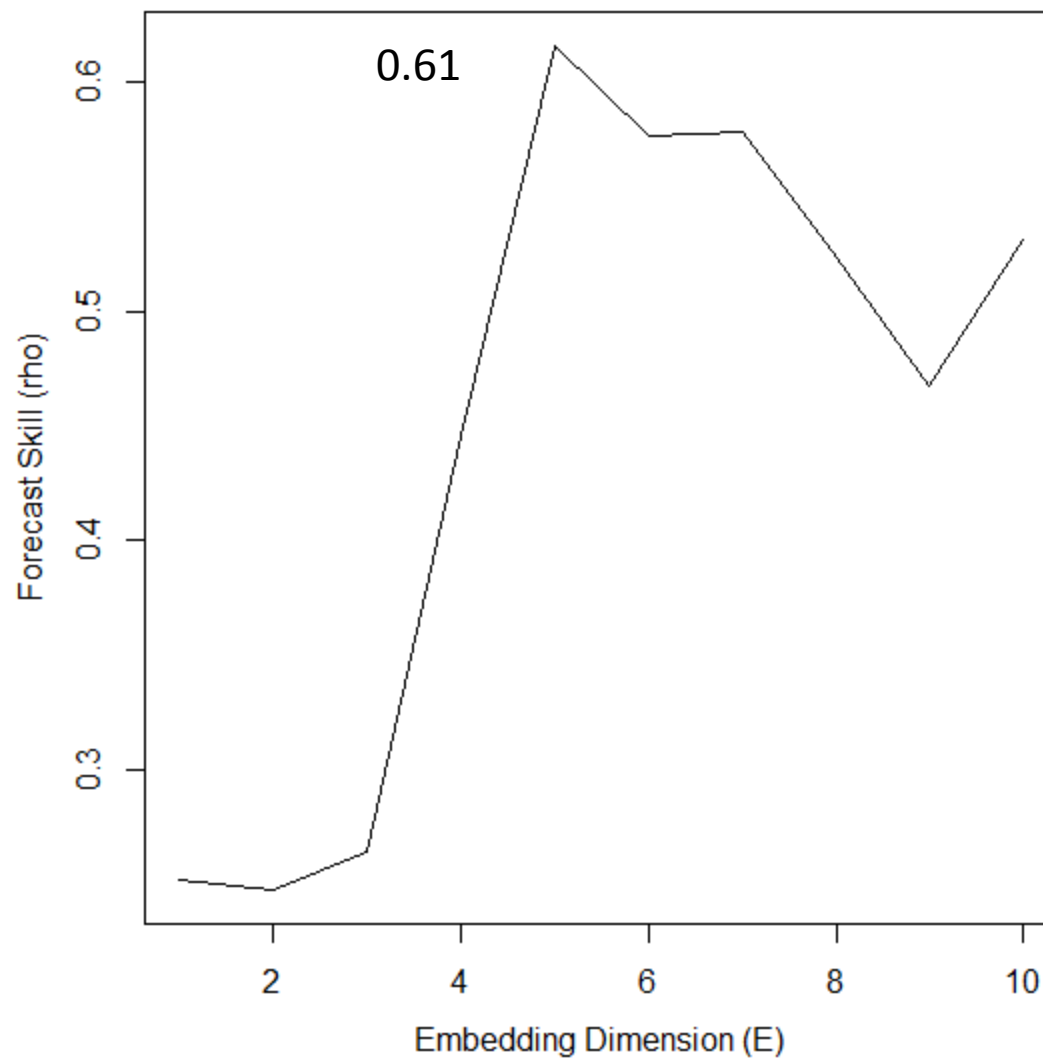


# Crab data

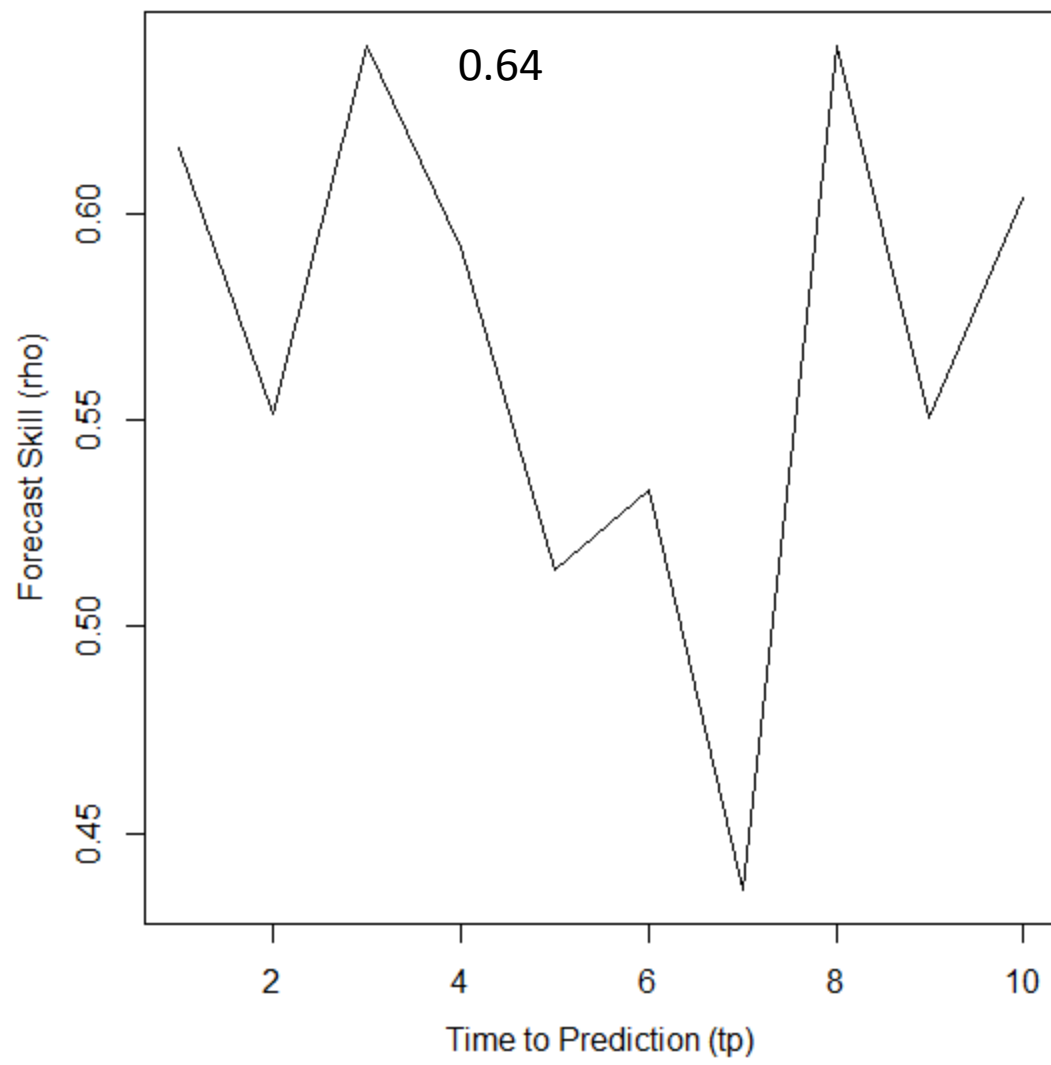
R1 (recruitment one year prior to entering fishery- Male mature gt 95mm)



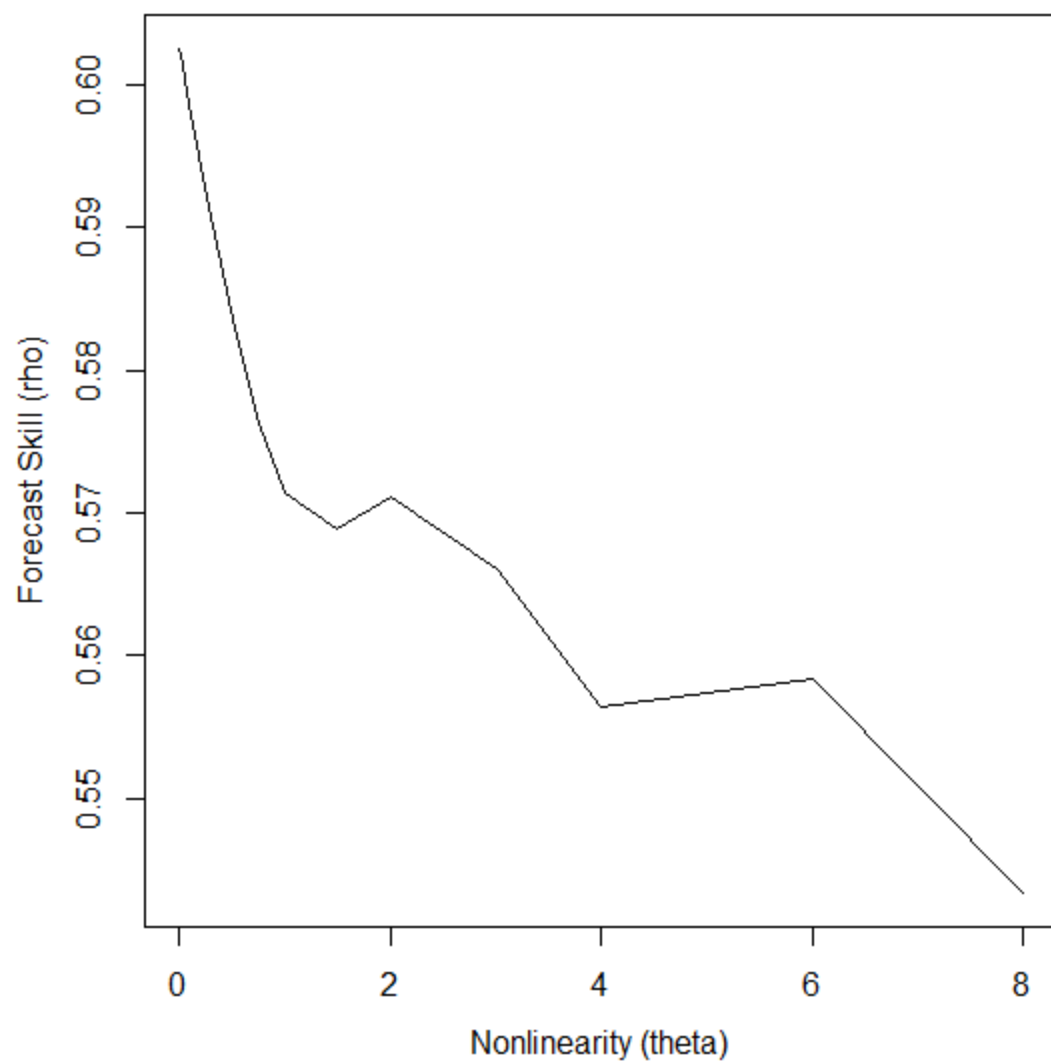
R1 (Male mature gt 95mm)



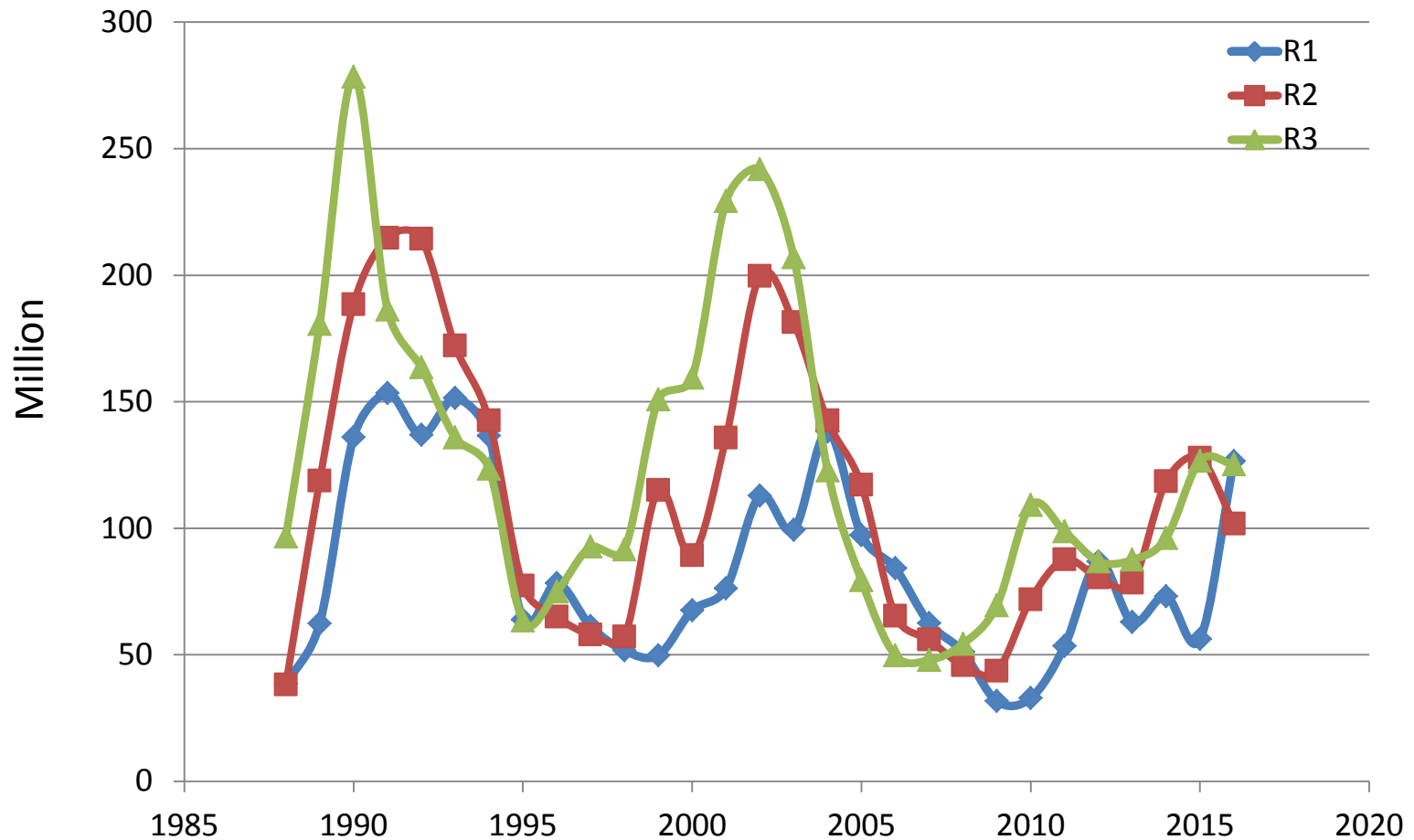
Embedding dimension  $E = 5$



S-map



R1 (recruitment 1 year prior to entering fishery- Male mature gt 95mm)  
R2 (recruitment 2 years prior to entering fishery- Male immature gt 83mm)  
R3 (recruitment 3 year prior to entering fishery- Male immature gt 63mm)

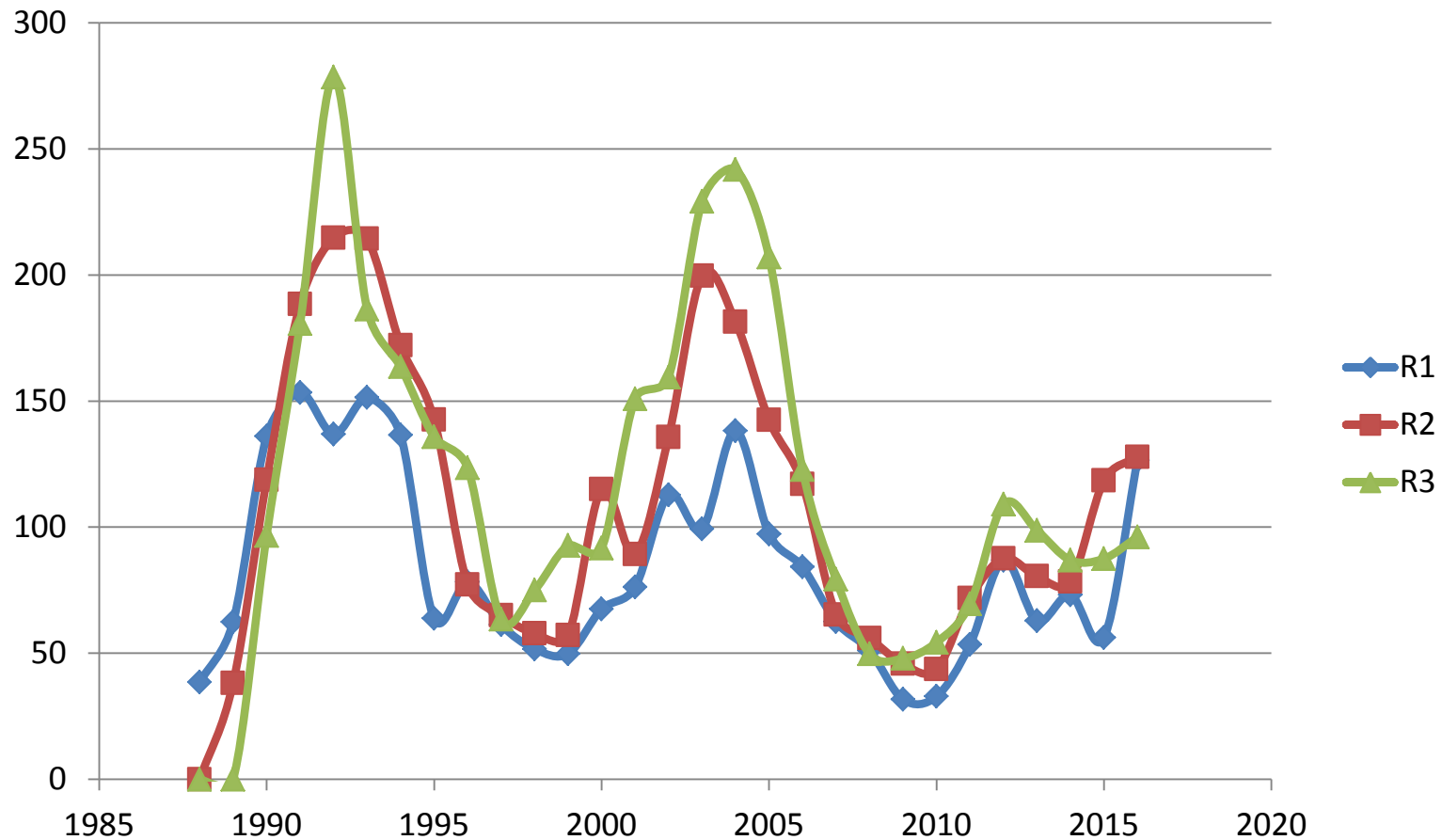




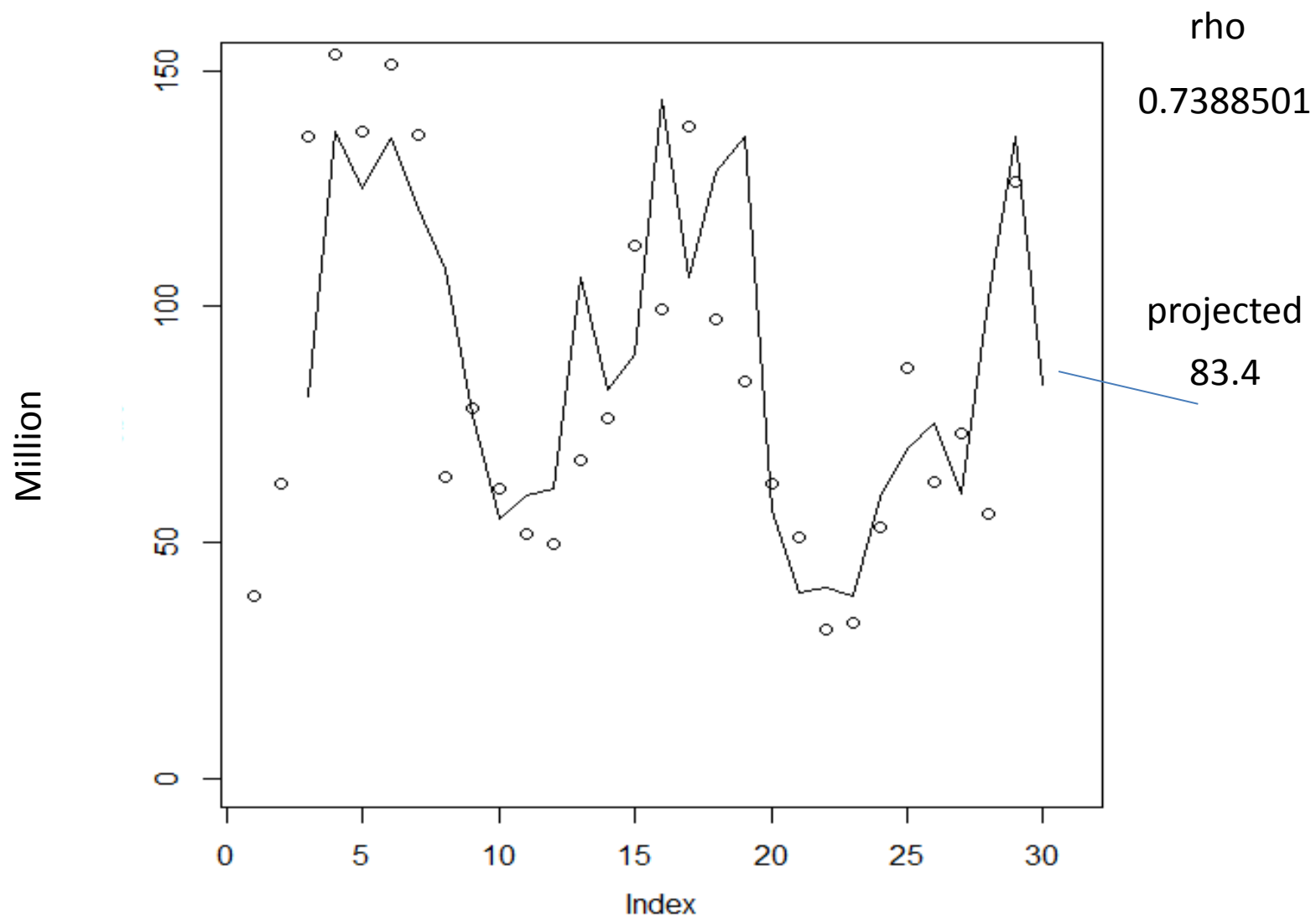
R1 from survey at year Y

R2 from survey at year Y-1 .... shown offset to match with year Y

R3 from survey at year Y-2 .... shown offset to match with year Y



R1 o observed — predicted



# Example code

```
> data("two_species_model")
> ts <- two_species_model$x[1:200]
> out = simplex(ts, lib = c(1, 100),
                pred=c(101, 200))

> head(round(out, dig=4))
```

	E	tau	tp	nn	num_pred	rho	mae
1	1	1	1	2	99	0.9977	0.0084
2	2	1	1	3	98	0.9989	0.0051
3	3	1	1	4	97	0.9991	0.0050
4	4	1	1	5	96	0.9972	0.0081
5	5	1	1	6	95	0.9965	0.0095
6	6	1	1	7	94	0.9949	0.0127

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4	4	1	1	5	96	0.9972	0.0081
5	5	1	1	6	95	0.9965	0.0095
6	6	1	1	7	94	0.9949	0.0127

Easy to know 'enough to be dangerous'.

# Recommendations

1. Worth continuing to understand – do more with simulated data.
2. Seems to be capturing some underlying dynamics in salmon (with updated data).
3. Does require a firm understanding of environmental mechanisms (don't jump to conclusions).