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Comparison of Performance of Models Fitting Organisms in Different Habitats

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1 Abstract

2 Accurate prediction of the impact of climate change on metabolic rate has overwhelm-
3 ingly practical significance. This study shows that the phenomenal model has stronger
4 predicting ability. And much more attention and patience are supposed to pay to mech-
5 anistic models. In order to get the best prediction, scholars should choose models and
6 model selection criteria with caution.

7 1 Introduction

8 Metabolism theory provides a foundation for using first principles of physics, chemistry,
9 and biology to link the performance from the molecular level(Allen et al. 2006) the ecology
10 level of population(Savage et al. 2004), community and ecosystem(Brown et al. 2004).
11 The theoretical basis is the core of metabolism consists of a small number of reactions
12 that are ultimately dependent on temperature(Gillooly et al. 2001).

13 Climate change is projected to have significant effects on public health(Woodward et al.
14 2014), agriculture(Howden et al. 2007), and fishing(Brander 2007). Prediction of the
15 impacts of climate change on freshwater, marine, and terrestrial environments can assist
16 the management of agriculture, fishery, and wildlife conservation(Prowse et al. 2009).

17 Here, a large dataset called BioTraits(Della et al. 2013), one phenomenological models,
18 Cubic model, one mechanistic models with three forms, Schoolfield model, Schoolfield
19 model for low temperature, and Schoolfield model for high temperature model, three se-
20 lection criteria, AIC, BIC, and AICc were used to study two main questions: (1) Is there
21 any difference in the performance of organisms in different habitats fitted by different
22 models? (2) If the answer of Question (1) is YES, so what is the best model in different
23 habitats?

24 In this study, the results show: (1) The differences in performances exist; (2) In most
25 cases, phenomenological model fits much better than mechanistic models. By differ-
26 ent model selection, the best mechanistic model varies under different environments.

27 Consequently, researchers should choose the most appropriate model selection criteria
28 according to the habitat where the research organism live in, the size of sample, and the
29 number of parameters, and choose the best model cautiously.

30 **2 Methods**

31 **2.1 Data**

32 BioTraits(Della et al. 2013) that consists of 25826 rows of data from 2165 researches from
33 small scale, like photosynthesis rate and respiration rate to large scale, like population
34 growth rate. The habitats included marine, terrestrial, and freshwater environments. At
35 kingdom level, the organisms were composed of archaea, bacteria, chromista, metazoa,
36 fungi, plantae, and protista.

37 But a large number of low quality data exist, NA, zero, and negetive value. If the body
38 temperature of consumer were missing, and ambient temperature replaced. Besides, the
39 researches with less than 8 data points were filtered out.

40 **2.2 Start values calculating for Schoolfield model**

41 Before fitting School model to the dataset, the starting value of the parameters were cal-
42 culated. First, the optimal temperature at which the trait value is highest. Before and after
43 this optimal temperature, linear regression analysis were conducted twice respectively.

44 E is the slope for the part after the optimal temperature, and its biological interpretation
45 is the activation energy. E_h is the slope before the optical temperature, and the biolog-
46 ical interpretation is the de-activation energy of enzyme at high temperature. T_h is the
47 at which the enzyme is 50% high-temperature deactivated, and was determined as the
48 mean value of the maximal and optical temperature. E_l is the slope for the part before
49 the optimal temperature, and the biological interpretation is the de-activation energy of
50 enzyme at low temperature. T_l is the at which the enzyme is 50% low-temperature deac-
51 tivated, and was determined as the mean value of the minimal and optical temperature.

Schoofield model would be discussed further in the following models section.

2.3 Models

2.3.1 Phenomenological model

Generally, the thermal performance curves have unimodal shape. Cubic model(Callaway 1959) are often used to study. This is Cubic model:

$$B = B_O + B_1 * T + B_2 * T^2 + B_3 * T^3 \quad (1)$$

The parameters: B_O , B_1 , B_2 , and B_3 don't have biological interpretations. This equation gives the trait value at the given T temperature.

2.3.2 Mechanistic models

A biological understanding of the response of metabolic rate to temperature is critical. Consequently, Schoolfiel model is used. This is full Schoolfiel model(Schoolfield et al. 1981):

$$B = \frac{B_0 e^{\frac{-E}{k}(\frac{1}{T} - \frac{1}{283.15})}}{1 + e^{\frac{E_l}{k}(\frac{1}{T_l} - \frac{1}{T})} + e^{\frac{E_h}{k}(\frac{1}{T_h} - \frac{1}{T})}} \quad (2)$$

Where k is the Boltzmann constant ($8.617 * 10^{-5} eV K^{-1}$), B the value of the trait at a given temperature T (K) ($K = ^\circ C + 273.15$), while B_0 is the trait value at 283.15 K ($10^\circ C$) which stands for the value of the growth rate at low temperature. E_l and E_h are the enzyme's low-temperature de-activation energy (eV) and high-temperature de-activation energy (eV). T_l and T_h are the at which the enzyme is 50% low-temperature deactivated and 50% high-temperature deactivated. E is the activation energy (eV).

In some cases, low temperature inactivation and high temperature are not available, the simplified models for low temperature and high temperature are more appropriate. This is the Schoolfield model for low temperature:

$$B = \frac{B_0 e^{\frac{-E}{k}(\frac{1}{T} - \frac{1}{283.15})}}{1 + e^{\frac{E_L}{k}(\frac{1}{T_L} - \frac{1}{T})}} \quad (3)$$

72 This is the Schoolfield model for high temperature:

$$B = \frac{B_0 e^{\frac{-E}{k}(\frac{1}{T} - \frac{1}{283.15})}}{1 + e^{\frac{E_h}{k}(\frac{1}{T_h} - \frac{1}{T})}} \quad (4)$$

73 **2.4 Model selection criteria**

74 Candidate models are compared to one another by evaluating the relative support in the
75 observed data for every model(Johnson & Omland 2004). Akaike information criteria
76 (AIC)(Burnham & Anderson 2004), Bayesian information criterion (BIC)(Schwarz et al.
77 1978), and Small sample unbiased AIC(AICC)(Burnham & Anderson 2004).

78 AS for AIC criteria, after models were fitted within each research, the minimal AIC was
79 determined, and the delta AIC was calculated. Finally, the relative weight of every model
80 was calculated. The best model is the model with most support(Burnham & Anderson
81 2004).

82 **2.5 Statistics analysis**

83 P values were calculated using Kruskal-Wallis test(Vargha & Delaney 1998) to compare
84 the significant difference in multiple groups. In two groups, p values were calculated
85 using Mann-Whitney test(Ruxton 2006). The reason is that the normal distribution and
86 consistent variance can't be guaranteed.

87 **2.6 Visualizaton**

88 The histogram suing point geom with the size of samples within one research, the boxplot
89 with weight supported by AIC, BIC, and AICC, and model fitting figure with one specific
90 research were plotted using ggplot2 (Wickham 2016) in R.

2.7 Computing languages

Bash 4.3.48: It was used to combine the pipeline together and compile the report into pdf format.

Python 3.5.2: It was used to model fitting. Some helpful packages were also used, *lmfit*(Newville et al. 2016), *pandas* (McKinney 2011), *numpy*(Van Der Walt et al. 2011), and *scipy* (Millman & Aivazis 2011).

R 3.2.3: It was used to pre-process data, calculate the starting value, plot the results and calculate the p values. Some helpful packages were used, *dplyr* (Wickham et al. 2015), *plyr*(Wickham 2009), *ggplot2* (Wickham 2016), and *reshape2*(Wickham 2012).

3 Results

3.1 Data

After removing the low quality data, the modified Biotraits contains 17525 rows of data from 971 researches. Here are the summaries of the modified Biotraits. In this dataset, Terrestrial and freshwater environments dominate over marine environments.

Table 1: Summary of researches included in different habitats. Terrestrial and freshwater environments dominate.

Habitats	Number of researches
Terrestrial	443
Marine	51
Freshwater	274

In this dataset, Bateria, fungi, metazoa, and plantae dominate over archaea, chromista, and protista.

Table 2: Summary of researches on organisms at Kindom level. Bateria, fungi, metazoa, and plantae dominate.

Kingdoms	Number of researches
Archaea	24
Bacteria	205
Chromista	16
Fungi	122
Metazoa	236
Plantae	366
Protista	1

107 Also, the size of samples within one research was summarised. The minimal is 8, be-
 108 cause researches with less than 8 data point were filtered out. And the maximal is 637.
 109 Most researches consists of less than 50 samples.

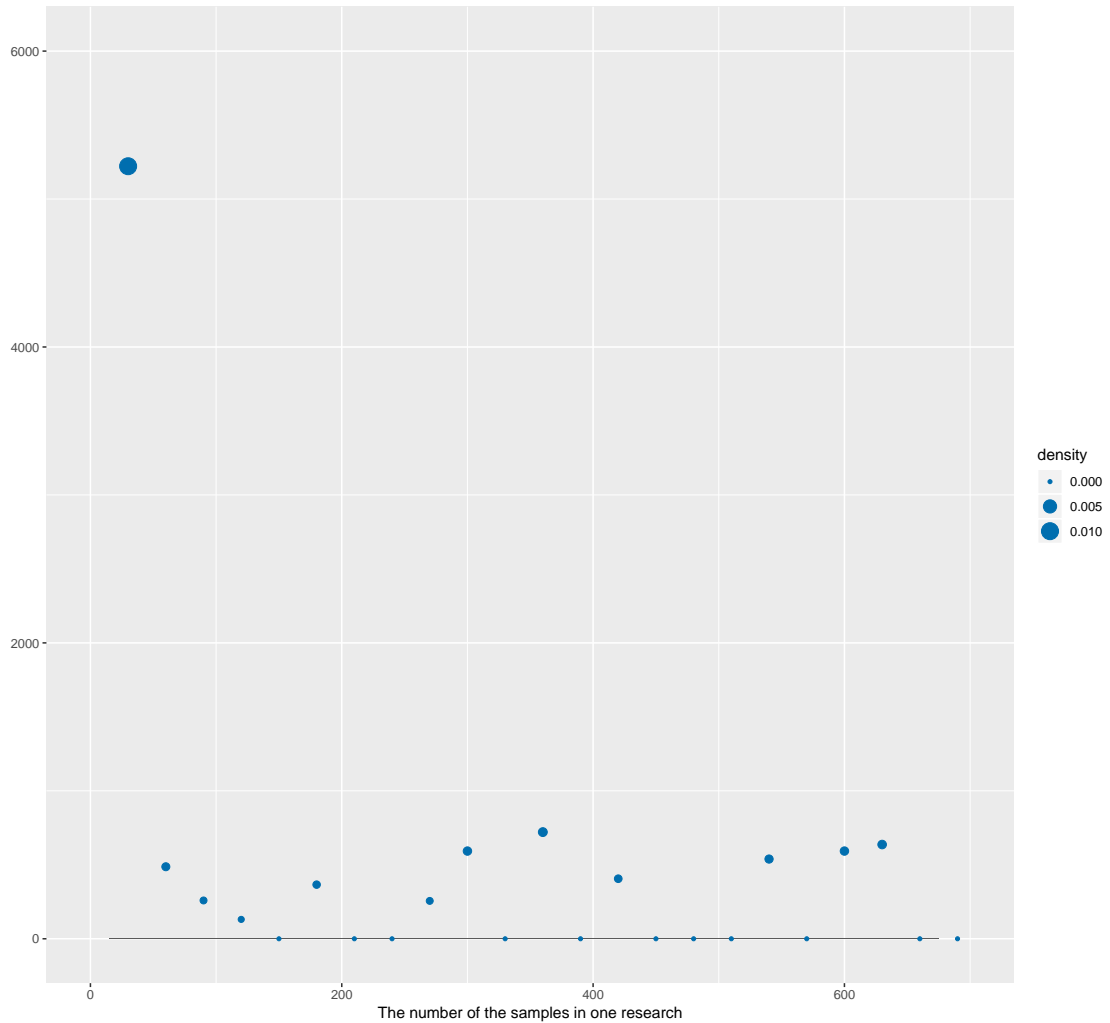


Figure 1: The size of samples within one research, most researches contains less than 50 samples.

3.2 Comparisons in performances of models fitting in habitats

Overall, the performances of all the models fitting in different habitats have significant difference, the p value $< 1.534e-07$. Cubic model has the best fitting performance over other three Schoolfield models. Within the three forms of Schoolfield models, the Schoolfield model for high temperature has the best performance.

In freshwater environments, under AIC, BIC, and AICC, the four models have different performances. All the pairwise p value < 0.01 , only one p value between Schoolfield model and Schoolfield model under AICC is 0.01296. And Cubic model fits best, especially under AICC.

119 In marine environments, only under AICC, the Cubid model has significant difference
120 with other three Schoolfield models. Under AIC and BIC, there isn't significant difference
121 with the four models. The reason could be penalty of the number of parameters from
122 AICC(Johnson & Omland 2004).

123 In terrestrial environments, under AICC, the Cubid model has significant difference with
124 other three Schoolfield models. Under AIC, the Cubic model has significant difference
125 with the Schoolfield model for high temperature and the Schoolfield model for low tem-
126 perature, the p values are < 0.009082 , and $< 6.75e-08$, respectively. And the Schoolfield
127 model has significant difference with the Schoolfield model for high temperature and the
128 Schoolfield model for low temperature, as well. The p values are < 0.03602 , and $< 1.13e-$
129 04 , respectively. Under BIC, except p value between Cubic model and the Schoolfield
130 model is > 0.05 , other p values are significant(p values are not shown).

131 In a word, phenomenological model has better performance than mechanistic models.
132 Under AICC, the weight value of three forms of the Schoolfield models are exceedingly
133 low. It is likely that there are too many parameters(Johnson & Omland 2004). With the
134 three forms, the Schoolfield model for high temperature and the Schoolfield model for low
135 temperature have relatively good performance in marine environments.

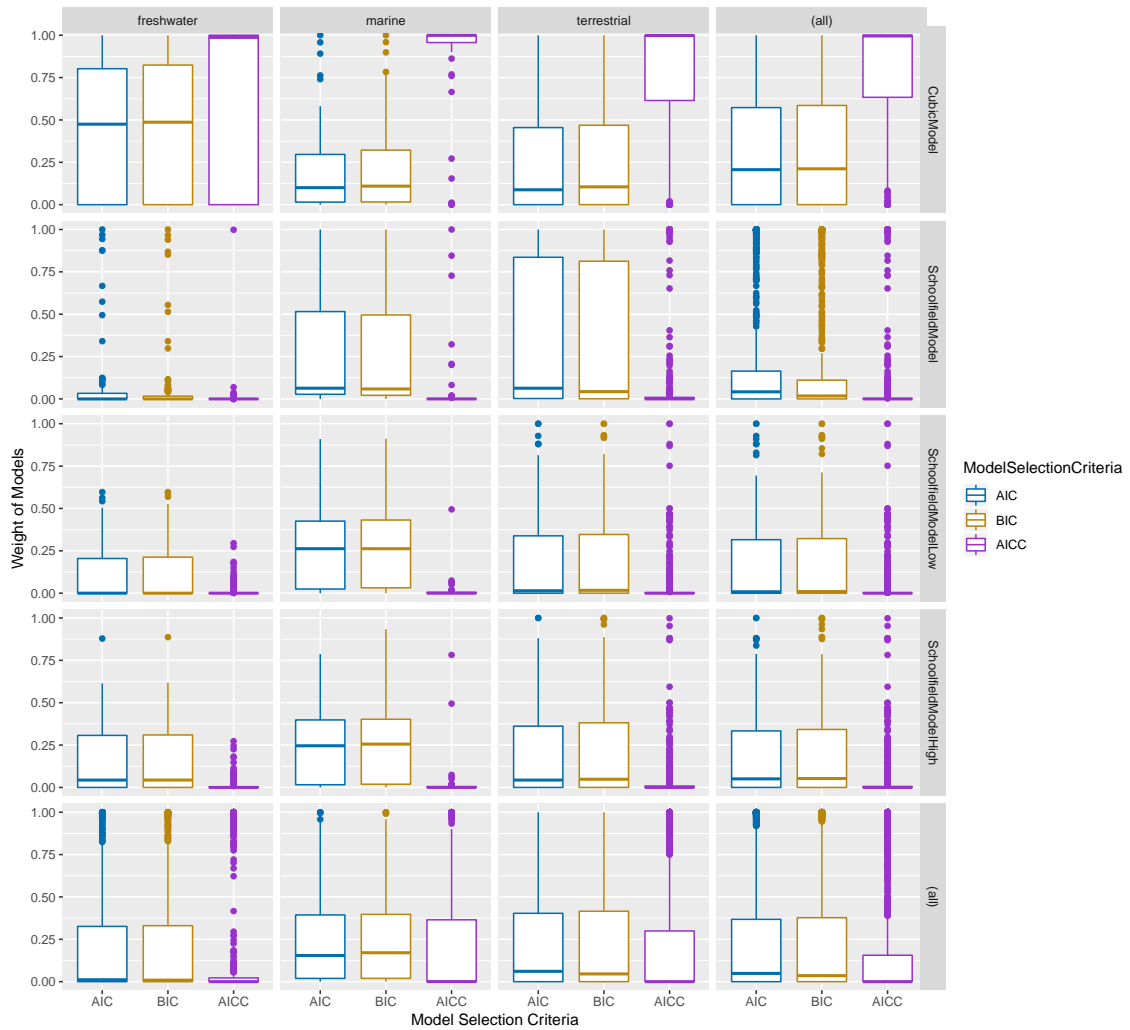


Figure 2: Comparisons in performances of models fitting in habitats. Cubic has the best performance. Under AICC, the weight value of three forms of Schoolfield model is extremely low.

3.3 Four models fitting one research

The FinalID of this research is MTD5347(Ratkowsky et al. 2005), the species is bacterium, *Escherichia.coli*, the original trait name is Square Root (Maximum Specific Growth Rate). The reason why this plot can only show three models is that the Schoolfield model for high temperature and the Schoolfield model for low temperature are quite similar and are overlapped.

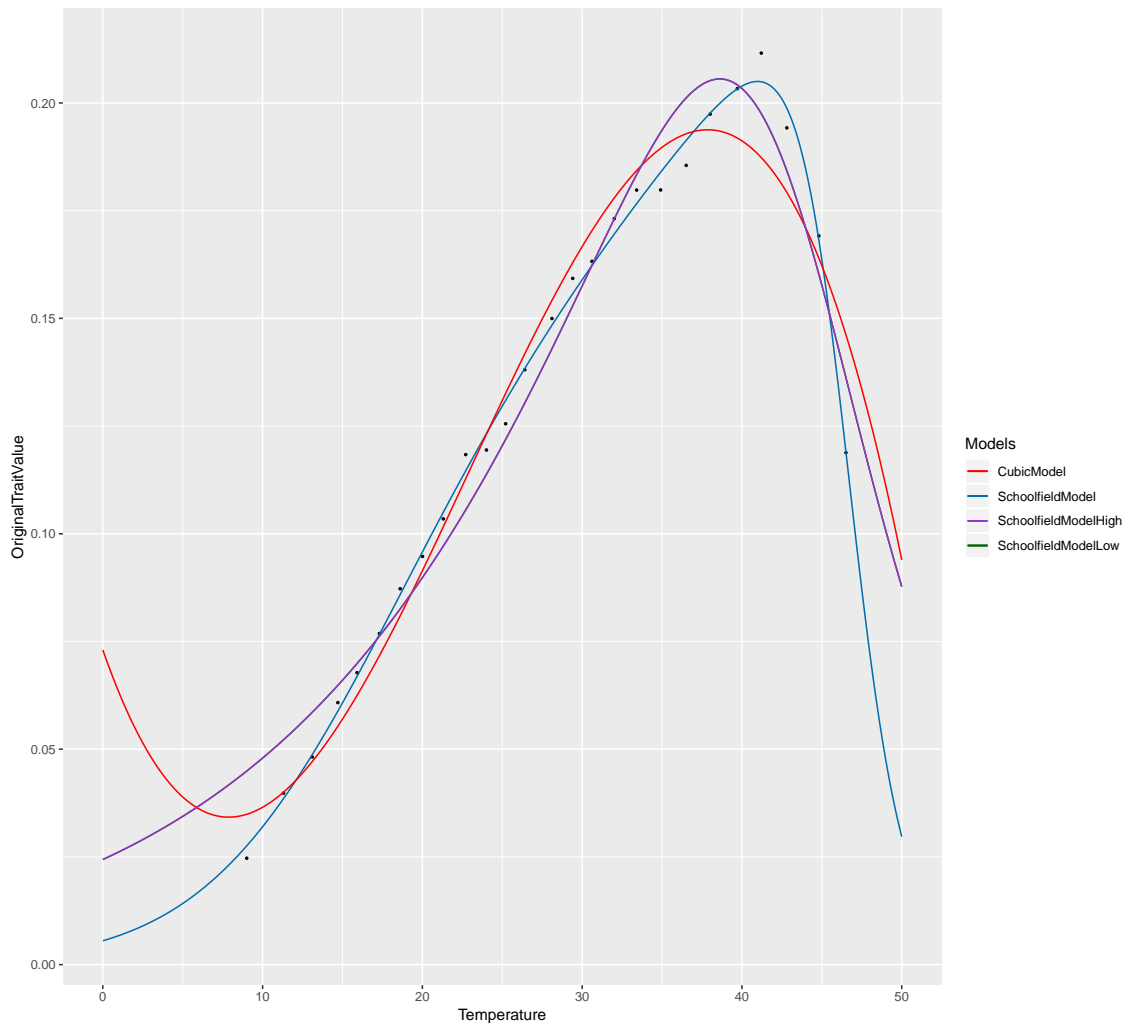


Figure 3: The four models fitted on one research. This curve shows the growth rate of *Escherichia.coli*.

4 Discussion

4.1 The reason behind the different performance

To start with, the distinct temperature ranges in different habitats is considered. For instance, under water, most organisms do not experience temperatures above 25°C – 30 °C, but in terrestrial environments, high temperatures of 36°C - 40 °C are common in summer.

However, later analysis shows the assumption is not supported. If the assumption is

149 true, the Schoolfield model for low temperature should have much better performance
150 than others. As shown above, that results are not drawn. Accordingly, researches on the
151 reason are needed.

152 **4.2 Limitations**

153 **4.2.1 Without further mining**

154 The large dataset contains many interesting contents. But this report is without further
155 mining. For instance, many Kingdom are included. Overall, the four models have sig-
156 nificant difference at Kingdom level, the p value is $1.55e-10$. But the pairwise significant
157 difference were not calculated. The reason has not been considered, either.



Figure 4: Comparisons in performances of models fitting in different organisms at Kingdom level.

158 4.2.2 Without further understanding of the model selection criteria

159 The extremely low weight value of Schoolfield models is due to the penalty of the num-
 160 bers of parameters from AICC. But the difference between AIC and BIC has not been
 161 discussed.

162 4.2.3 Visual flaw

163 Some of the figures are too complicated to add p values, and all the p values can't be
 164 shown at the figures.

165 **5 Conclusion**

166 In conclusion, phenomenological model has better performance than mechanistic models
167 in different habitats. As for mechanistic models, in different habitats and under different
168 model selection criteria, there is better model to choose. Researchers are supposed to
169 choose the better models and model selection criteria carefully in terms of the specific
170 background, habitat and so forth.

171 We need more researches to study to mechanisms behind the metabolic reactions and
172 propose mechanistic models with precise prediction, which is especially essential at the
173 time when extreme climate frequent and climate change has a critical influence on agri-
174 culture, fisheries, and health.

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