Title: A derivation error exists in the current implementation of the Johnson et al. (1942) modified

Arrhenius function that affects leaf carbon balance models

Running Title: Derivation error in modified Arrhenius model

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Abstract

Understanding biological temperature responses is crucial to predicting global carbon fluxes. The

current approach to modeling photosynthetic temperature responses in large scale modeling efforts

uses a modified Arrhenius equation. We rederived the modified Arrhenius equation from the source

publication and uncovered a missing term that was dropped between 1942 and 2002. We compare

fitted temperature response parameters between the new and old derivation of the modified

Arrhenius equation. We find that most parameters are minimally affected, though small errors still

exist. We then scaled the impact of these small errors to whole plant carbon balance and found that

the impact of the rederivation of the Arrhenius on modelled daily carbon gain causes a meaningful

deviation of ~1.8%. This suggests that the error in the derivation of the modified Arrhenius equation

has impacted predictions of carbon fluxes at larger scales. We argue that it is time to move beyond

the modified Arrhenius paradigm since the current implementation is categorically incorrect and to

use more thermodynamically-grounded temperature response equations going forward.

Keywords: Arrhenius, temperature, photosynthesis, gas exchange, modeling, carbon balance

Introduction

Globally, photosynthesis and autotrophic respiration are the largest biological carbon fluxes, with photosynthesis removing ~120 Gt C year^{-1} and autotrophic respiration releasing ~60 Gt C year^{-1} (Amthor, 2000; Ciais et al., 2013). Given the temperature sensitivity of these large carbon fluxes, understanding how photosynthesis and respiration respond on acute, acclimatory, and adaptive timescales is crucial for predicting vegetative and carbon cycle responses to future global climates (Rogers et al., 2017; Stinziano et al. 2018). Biological temperature responses including photosynthesis and respiration are typically assumed to be exponential or peaked (Way and Yamori, 2014; Smith & Dukes, 2017; Kumarathunge et al., 2019). Exponential responses are usually modelled based on an Arrhenius-type curve (Arrhenius, 1915):

$$f(T) = Ae^{\frac{-E_a}{RT}}$$
Equation 1

$$f(T) = k_{25}e^{\left[\frac{E_a(T-298.15)}{RT298.15}\right]}$$
Equation 2

where f(T) is the rate of the process at temperature, A is a pre-exponential factor, T in K, k_{25} is the rate of the process at 298.15 K, E_a is the activation energy in J mol^{-1}, R is the universal gas constant of 8.314 J mol^{-1} K^{-1}, and 298.15 is the reference temperature in K. As for peaked responses, while a few options are available (Kruse et al., 2008; Hobbs et al., 2013; Heskel et al., 2016), the most commonly implemented version is the modified Arrhenius model of Johnson et al. (1942) as presented in Medlyn et al. (2002):

$$f(T) = k_{25}e^{\left[\frac{E_a(T-298.15)}{RT298.15}\right]} \left[\frac{1+e^{\frac{298.15\Delta S - H_d}{298.15R}}}{1+e^{\frac{T\Delta S - H_d}{TR}}}\right]$$
Equation 3

where H_d is the deactivation energy in J mol^{-1}, and ΔS is the entropy of the process in J mol^{-1}.

Equation 3 is used for modeling the temperature responses of photosynthetic capacity (based on Farquhar et al., 1980): maximum carboxylation capacity of rubisco, V_{cmax}, maximum electron transport capacity, J_{max}, and related kinetics as well as Sharkey (1985) and Harley & Sharkey (1991) in their

calculation of triose phosphate utilization capacity (TPU). These parameters are then used in

ecophysiological studies to understand thermal acclimation of photosynthesis (see Kattge & Knorr,

2007; Smith & Dukes, 2017; and Kumarathunge et al., 2019 for examples). Furthermore, this equation

is also used in terrestrial biosphere models to predict the future state of the Earth system (e.g. Rogers

et al., 2017).

Due to its ubiquity, we revisited the original Johnson et al. (1942) modified Arrhenius function to

rederive Equation 3. In the process of this rederivation, we uncovered a term that was dropped

sometime between Johnson et al. (1942) and Medlyn et al. (2002) which causes a systematic error in

the application of Equation 3 in calculations covering individual species (e.g. Medlyn et al., 2002) to

global scale processes (e.g. Rogers et al., 2017). We then refit a freely available dataset

(Kumarathunge et al., 2019), with both versions of the modified Arrhenius model, and fed the

temperature response fits through a carbon balance model to estimate the impact of the derivation

error on modelled plant carbon balance. We predicted that the derivation error would cause

substantial variation in fitted temperature response parameters, and that these differences would

propagate through to modelled daily carbon balance.

Materials & methods

Rederivation of the modified Arrhenius equation

Johnson et al. (1942, equation 24) describe the temperature response of the light intensity of a

luciferase reaction as:

85
$$I = \frac{c'' T e^{\frac{-\Delta H\ddagger}{RT}}}{1 + e^{\frac{\Delta S}{R}} e^{\frac{-\Delta H}{RT}}}$$
     Equation 4

where I is the intensity of the luciferase reaction, c'' is not explicitly defined in Johnson et al. (1942),

but presumably represents a second derivative of the rate, T is the temperature in K, R is the universal

gas constant of 8.314 J mol^{-1} K^{-1}, $\Delta H\ddagger$ is the activation energy in J mol^{-1}, ΔH is the deactivation energy

in J mol^{-1}, and ΔS is the entropy in J mol^{-1}. We can relativize the equation to a reference temperature:

$$\frac{I}{I_{25}} = \frac{\dfrac{c''Te^{\frac{-\Delta H\ddagger}{RT}}}{1+e^{\frac{\Delta S}{R}}e^{\frac{-\Delta H}{RT}}}}{\dfrac{c''298.15e^{\frac{-\Delta H\ddagger}{R298.15}}}{1+e^{\frac{\Delta S}{R}}e^{\frac{-\Delta H}{R298.15}}}}$$ Equation 5

$$\frac{I}{I_{25}} = \frac{c''Te^{\frac{-\Delta H\ddagger}{RT}}}{c''298.15e^{\frac{-\Delta H\ddagger}{R298.15}}}\frac{1+e^{\frac{\Delta S}{R}}e^{\frac{-\Delta H}{R298.15}}}{1+e^{\frac{\Delta S}{R}}e^{\frac{-\Delta H}{RT}}}$$ Equation 6

$$\frac{I}{I_{25}} = \frac{T}{298.15}e^{\frac{-\Delta H\ddagger}{RT}-\frac{-\Delta H\ddagger}{R298.15}}\frac{1+e^{\frac{\Delta S}{R}+\frac{-\Delta H}{R298.15}}}{1+e^{\frac{\Delta S}{R}+\frac{-\Delta H}{RT}}}$$ Equation 7

$$I = I_{25}\frac{T}{298.15}e^{\left[\frac{-\Delta H\ddagger 298.15 - -\Delta H\ddagger T}{RT298.15}\right]}\left[\frac{1+e^{\frac{298.15\Delta S+-\Delta H}{298.15R}}}{1+e^{\frac{T\Delta S+-\Delta H}{TR}}}\right]$$ Equation 8

$$I = I_{25}\frac{T}{298.15}e^{\left[\frac{\Delta H\ddagger(T-298.15)}{RT298.15}\right]}\left[\frac{1+e^{\frac{298.15\Delta S-\Delta H}{298.15R}}}{1+e^{\frac{T\Delta S-\Delta H}{TR}}}\right]$$ Equation 9

Harmonizing the notation scheme to that typically used in plant ecophysiology:

$$f(T) = k_{25}\frac{T}{298.15}e^{\left[\frac{E_a(T-298.15)}{RT298.15}\right]}\left[\frac{1+e^{\frac{298.15\Delta S-H_d}{298.15R}}}{1+e^{\frac{T\Delta S-H_d}{TR}}}\right]$$ Equation 10

Note the difference between Equations 3 and 10: the term T / 298.15 is missing from Equation 3. This

introduces multiple systematic errors:

1) fitted parameters E_a, H_d, and ΔS;

2) scaling the rate variable using the wrong equation;

3) acclimation of E_a, H_d, and/or ΔS as errors due to 1) at each acclimation temperature.

Here we focus on the impact of 1 and 2 on modeling whole-plant carbon balance.

Data analysis

Using data from Kumarathunge et al. (2019), available from Kumarathunge et al. (2018), we fit

Equations 3 and 10 to the data, both setting H_d to 200 kJ mol^{-1} and allowing H_d to be fit. We used the

R package {minpack.lm} (Elzhov et al., 2016) with starting parameters of E_a = 40 kJ mol^{-1}, ΔS = 0.650 kJ

mol^{-1}, k_{25} = mean(parameter), and H_d (when fitted) varying from 1 to 500 kJ mol^{-1}, followed by the

{BIC} function to select the best model based on Bayesian Information Criteria. We obtained 92

successful curve fits for the fixed H_d case, and 84 successful curve fits for the fitted H_d case. This

allows us to explore the impact of the missing term on the output data under a typical fitting scenario

(H_d is fixed) and under the full fitting scenario.

Modeling

We modelled the impact of the equations 3 and 10 on daily net plant carbon balance under

conditions of H_d = 200 kJ mol^{-1} and fitted H_d. Data for leaf area, root and shoot masses, as well as leaf

dark respiration at 25 °C were taken for white spruce (*Picea glauca*) from Stinziano & Way (2017),

while stomatal conductance model parameters were calculated with the gas exchange data reported

in Stinziano & Way (2017). Mean data were taken from the control treatment at weeks 0 and 12 to

provide contrasting biomass allocation patterns such that week 0 is a low respiration scenario and

week 12 is a high respiration scenario. These different respiration scenarios were used to reduce bias

in any conclusions regarding the impact of Equations 3 and 10 on carbon balance, as the ratio of

photosynthesis to respiration may alter the sensitivity of carbon balance to the Arrhenius equation

used. Root respiration for white spruce was taken from Weger and Guy (1991) and we assumed that

stem respiration was equal to root respiration (Table 1).

Table 1. Parameters used in modeling daily carbon gain.

Parameter	Group	Value	Reference
Respiration	R_{dark}	2.78 μmol m^{-2} s^{-1}	Stinziano & Way, 2017
	R_{day}	0.7 * R_{dark}	Ayub et al., 2011
	R_{root}	0.0095 μmol g^{-1} s^{-1}	Weger & Guy, 1991
	R_{stem}	0.0095 μmol g^{-1} s^{-1}	Assumed
	Q_{10}	2.015	Atkin & Tjoelker, 2003
Γ^*	25 °C	42.75 μmol mol^{-1}	Bernacchi et al., 2001
	E_a	37.83 kJ mol^{-1}	Bernacchi et al., 2001
K_m	25 °C	718.4 μmol mol^{-1}	Bernacchi et al., 2001
	E_a	65.51 kJ mol^{-1}	Bernacchi et al., 2001
α		0.8	Norman & Campbell, 1998
ϕ		0.08	Norman & Campbell, 1998
Leaf Area	Low R	0.015 m^2	Stinziano & Way, 2017
	High R	0.025 m^2	Stinziano & Way, 2017

Stem Mass	Low R	0.496 g	Stinziano & Way, 2017
	High R	2.523 g	Stinziano & Way, 2017
Root Mass	Low R	0.498 g	Stinziano & Way, 2017
	High R	5.072 g	Stinziano & Way, 2017

Q_{10}: thermal sensitivity coefficient; Γ^*: photorespiratory CO_2 compensation point; K_m: Michaelis-Menten constant for rubisco; α: absorbance of photosynthetically activation radiation; ϕ: maximum quantum efficiency of photosynthetic electron transport; E_a: activation energy; R_{dark}: leaf respiration in the dark; R_{day}: leaf respiration in the light; R_{root}: root respiration; R_{stem}: stem respiration; Low R: low respiration scenario; High R: high respiration scenario.

For the full model structure and equations, please see the accompanying R package {arrhenius.comparison} ("arrhenius.comparison_1.0.0.tar.gz"; Stinziano & Murphy, 2020) (see Table 2 for equations). Briefly, we linked the Medlyn et al. (2011) stomatal conductance model with the Farquhar et al. (1980) C₃ photosynthesis model, assuming infinite mesophyll conductance to CO_2 as these assumptions were used in fitting the data from Kumarathunge et al. (2018). Photosynthetic capacity, both maximum rubisco carboxylation capacity, V_{cmax}, and maximum electron transport rate, J_{max} were scaled to temperature using either Equation 3 or 10, while respiration was scaled according to (Atkin & Tjoelker, 2003). Photosynthesis and respiration were summed across each modelled day to calculate daily plant carbon assimilation.

Table 2. Equations used in modeling daily carbon uptake.

Equation	Reference
$$f(T) = k_{25} Q_{10}^{\frac{T-298.15}{10}}$$	Atkin & Tjoelker, 2003
$$f(T) = k_{25} e^{\left[\frac{E_a(T-298.15)}{RT298.15}\right]}$$	Arrhenius, 1915
$$f(T) = k_{25} e^{\left[\frac{E_a(T-298.15)}{RT298.15}\right]} \left[\frac{1 + e^{\frac{298.15\Delta S - H_d}{298.15R}}}{1 + e^{\frac{T\Delta S - H_d}{TR}}}\right]$$	Johnson et al., 1942; Medlyn et al., 2002
$$f(T) = k_{25} \frac{T}{298.15} e^{\left[\frac{E_a(T-298.15)}{RT298.15}\right]} \left[\frac{1 + e^{\frac{298.15\Delta S - H_d}{298.15R}}}{1 + e^{\frac{T\Delta S - H_d}{TR}}}\right]$$	Johnson et al., 1942; this study
$$W_c = V_{cmax} \frac{C_i - \Gamma^*}{C_i + K_m}$$	Farquhar et al., 1980

$$W_j = min\left(J_{max}, \alpha\phi Q_{in}\frac{C_i - \Gamma*}{C_i + 2\Gamma*}\right)$$

Farquhar et al., 1980; Way et al., 2011

$$A_{gross} = min(W_c, W_j)$$

$$A_{net} = A_{gross} - R_{day}$$

$$g_s = g_0 + 1.6\left(1 + \frac{g_1}{\sqrt{VPD}}\right)\left(\frac{A_{net}}{C_a}\right)$$

Medlyn et al., 2011

$$C_i = C_a - 1.6\frac{A_{net}}{g_s}$$

$$A_{plant} = A_{net} \times LA - R_{dark} \times LA - R_{stem} \times SM$$
$$- R_{root} \times RM$$

$$C_{balance} = \sum A_{plant} \times 3600 \times \frac{12.01}{1,000,000}$$

f(T): rate of a process at a given temperature; T: temperature in K; k_{25}: rate of a process at 25 °C;

Q_{10}: thermal sensitivity coefficient; E_a: activation energy in kJ mol^{-1}; ΔS: entropy parameter in kJ

mol^{-1}; H_d: deactivation energy in kJ mol^{-1}; R: universal gas constant in 0.008314 kJ mol^{-1} K^{-1}; W_c: rate

of CO_2-limited carboxylation in μmol m^{-2} s^{-1}; W_j: rate of RuBP regeneration-limited carboxylation in

μmol m^{-2} s^{-1}; V_{cmax}: maximum rate of rubisco carboxylation capacity in μmol m^{-2} s^{-1}; J_{max}: maximum

rate of electron transport in μmol m^{-2} s^{-1}; C_i: intercellular CO_2 concentration in μmol mol^{-1}; $\Gamma*$:

photorespiratory CO_2 compensation point in μmol mol^{-1}; K_m: Michaelis-Menten constant for rubisco

in μmol mol^{-1}; α: absorbance of photosynthetically active radiation; ϕ: maximum quantum

efficiency of electron transport; Q_{in}: incident photosynthetically active radiation; A_{gross}: gross CO_2

assimilation in μmol m^{-2} s^{-1}; A_{net}: net CO_2 assimilation in μmol m^{-2} s^{-1}; R_{day}: leaf day respiration in

μmol m^{-2} s^{-1}; g_s: stomatal conductance to water in mol m^{-2} s^{-1}; g_o: intercept for the Medlyn et al.

2011 model; g_1: slope for the Medlyn et al. 2011 model; VPD: vapor pressure deficit in kPa; C_a: CO_2

concentration at the leaf surface in μmol mol^{-1}; A_{plant}: whole plant net CO_2 assimilation; LA: leaf

area in m^{-2}; R_{dark}: leaf dark respiration in μmol m^{-2} s^{-1}; R_{stem}: stem respiration in μmol m^{-2} s^{-1}; R_{root};

root respiration in μmol m^{-2} s^{-1}; SM: stem mass in g; RM: root mass in g; $C_{balance}$: whole plant daily

carbon balance in g $plant^{-1}$ day^{-1}.

Modeling was performed on 18 total days of environmental data, with three days of data from three

months (17th – 19th of May, August, and October, 2019) obtained from external irradiance sensors at

the Biotron Experimental Climate Change Research Centre at the University of Western Ontario and

the remaining environmental data from Environment Canada historical climate data for South London

(43.01°N, 81.27°W, altitude: 251 m) and the rooftop greenhouse at the University of New Mexico

(35.08°N, 106.62°W, altitude: 1587 m) to capture different levels of environmental variability (Fig. 1).

Figure 1. Environmental data used to drive the model in Table 1 covering 3 days (17th, 18th, and 19th)

of 3 months. (a,c,e) Albuquerque, NM, USA; (b,d,e) London, ON, Canada.

Overall, the modeling approach allows us to assess the relative differences between Equations 3 and

10 under a low- and high- respiration scenario, fixing versus fitting H_d, and across different ranges of

seasonal variability.

Statistical analysis

Data were analyzed using the {lm} function in R v.3.6.1 (R Core Team, 2019), regressing the data

obtained from Equation 10 against the data obtained from Equation 3. All code and data will be made

freely available on GitHub upon publication in the {arrhenius.comparison} R package (Stinziano &

Murphy, 2020).

Results

Fixed H_d of 200 kJ mol^{-1}

Incorporating the missing term into Equation 3 to produce Equation 10 caused small differences in

the fitting of E_a for V_{cmax}, with a slope of 0.997 ± 0.001 and intercept of -2.162 ± 1.19 · 10^{-1} kJ mol^{-1}

($F_{1,90}$ = 4.56 · 10^5, R^2 = 0.9998, P < 2.2 · 10^{-16}) (Fig. 2b). For ΔS, the slope was 0.991 ± 3 · 10^{-3} and an

intercept of 0.005 ± 2 · 10^{-3} kJ mol^{-1} (intercept approximately 1% of fitted ΔS) ($F_{1,90}$ = 1.23 · 10^5, R^2 =

0.9993, P < 2.2 · 10^{-16}) (Fig. 2f). Meanwhile V_{cmax25} was identical between both approaches with a

slope of 1.000 ± 1.1 · 10^{-5} (intercept not significant; $F_{1,91}$ = 8.2 · 10^9, R^2 = 1.000, P < 2.2 · 10^{-16}) (Fig. 3b).

For J_{max}, there were small differences in the fitting of E_a for J_{max}, with a slope of 0.998 ± 2 · 10^{-3} and

intercept of -2.230 ± 1.08 · 10^{-1} kJ mol^{-1} ($F_{1,91}$ = 3.3 · 10^5, R^2 = 0.9997, P < 2.2 · 10^{-16}) (Fig. 2d). For ΔS,

the slope was 0.962 ± 3 · 10^{-3} and an intercept of 0.025 ± 2 · 10^{-3} kJ mol^{-1} (intercept approximately 5%

of fitted ΔS) ($F_{1,90}$ = 1.1 · 10^5, R^2 = 0.9992, P < 2.2 · 10^{-16}) (Fig. 2h). Meanwhile J_{max25} was identical

between both approaches with a slope of 1.000 ± 1.2 · 10^{-5} (intercept not significant; $F_{1,92}$ = 6.7 · 10^9,

R^2 = 1.000, P < 2.2 · 10^{-16}) (Fig. 3d).

Figure 2 – The modified Arrhenius equation with the missing term (Equation 10) gives similar temperature response parameters as Equation 3 for E_a (a-d) and ΔS (e-h) for both V_{cmax} (a, b, e, f) and J_{max} (c, d, g, h) under scenarios where H_d is allowed to vary (a, c, e, g) and is fixed to 200 kJ mol^{-1} (b, d, f, h). "New" indicates Equation 10, "Old" indicates Equation 3. E_a: activation energy, ΔS: entropy parameter, H_d: deactivation energy, V_{cmax}: maximum capacity of rubisco carboxylation, J_{max}: maximum rate of electron transport. Black line indicates 1:1 line and grey dashed line indicates respective modeled slopes and intercepts.

Figure 3 – The modified Arrhenius equation with the missing term (Equation 10) gives similar temperature response parameters as Equation 3 for k_{25} (a-d) and H_d (e, f) for both V_{cmax} (a, b, e) and J_{max} (c, d, f) under scenarios where H_d is allowed to vary (a, c, e, f) and is fixed to 200 kJ mol^{-1} (b, d). "New" indicates Equation 10, "Old" indicates Equation 3. k_{25}: rate of the process at 25 °C, H_d: deactivation energy, V_{cmax}: maximum capacity of rubisco carboxylation, J_{max}: maximum rate of electron transport. Black line indicates 1:1 line and grey dashed line indicates respective modeled slopes and intercepts.

Variable H$_d$

Incorporating the missing term into Equation 3 to produce Equation 10 caused small differences in the fitting of E$_a$ for V$_{cmax}$, with a slope of 0.966 ± 2.6 · 10^{-2} (intercept not significant; $F_{1,74}$ = 3005, R^2 = 0.976, P < 2.2 · 10^{-16}) (Fig. 2a). For ΔS, the slope was 0.989 ± 1.7 · 10^{-2} (intercept not significant; $F_{1,74}$ = 3339, R^2 = 0.9783, P < 2.2 · 10^{-16}) (Fig. 2e), while H$_d$ had a slope of 0.992 ± 1.9 · 10^{-2} (intercept not significant; $F_{1,74}$ = 2812, R^2 = 0.9744; P < 2.2 · 10^{-16}) (Fig. 3e). Meanwhile V$_{cmax25}$ was nearly identical between both approaches with a slope of 1.000 ± 4 · 10^{-3} (intercept not significant; $F_{1,74}$ = 7.98 · 10^6, R^2 = 1.000, P < 2.2 · 10^{-16}) (Fig. 3a). For J$_{max}$, there were small differences in the fitting of E$_a$ for J$_{max}$, with a slope of 1.000 ± 5 · 10^{-3} and an intercept of -3.119 ± 1.089 kJ mol^{-1} ($F_{1,80}$ = 3.6 · 10^4, R^2 = 0.9978, P < 2.2 · 10^{-16}) (Fig. 2c). ΔS was identical, the slope was 1.000 ± 4 · 10^{-3} (intercept not significant; $F_{1,81}$ = 6.4 · 10^4, R^2 = 0.9987, P < 2.2 · 10^{-16}) (Fig. 2g), while H$_d$ was identical with a slope of 1.000 ± 4 · 10^{-3} (intercept not significant; $F_{1,81}$ = 5.2 · 10^4, R^2 = 0.9985; P < 2.2 · 10^{-16}) (Fig. 3f). J$_{max25}$ was identical between both approaches with a slope of 1.000 ± 1 · 10^{-3} (intercept not significant; $F_{1,81}$ = 5.8 · 10^7, R^2 = 1.000, P < 2.2 · 10^{-16}) (Fig. 3c).

Impacts on modelled net carbon balance

In general, the differences in thermal response parameters were amplified when integrated at the whole-plant level. For daily photosynthesis (A) measured at a fixed H$_d$ of 200 kJ mol^{-1}, the slope for the low respiration model was 0.977 ± 7.666 · 10^{-3} and an intercept of 0.002 ± 9.366 · 10^{-5} (approximately 2% of modeled A) ($F_{1,898}$ = 1.625 · 10^6, R^2 = 0.9994; P < 2.2 · 10^{-16}) (Fig. 4b). Similarly, the high respiration model for A had a slope of 0.977 ± 7.666 · 10^{-3} and an intercept of 0.004 ± 1.591 · 10^{-3} (approximately 2% of modeled A) ($F_{1,898}$ = 1.625 · 10^6, R^2 = 0.9994; P < 2.2 · 10^{-16}) (Fig. 4d). For daily A measured at a variable H$_d$, the slope for the low respiration model was 0.983 ± 9.909 · 10^{-3} and the intercept was 0.002 ± 1.128 · 10^{-3} (approximately 2% of modeled A) ($F_{1,556}$ = 9.834 · 10^5, R^2 = 0.9994; P < 2.2 · 10^{-16}) (Fig. 4a). For the high respiration model of daily A, the slope was 0.983 ± 9.909 · 10^{-3} and the intercept was 0.003 ± 9.909 · 10^{-4} (approximately 2% of modeled A) ($F_{1,556}$ = 9.834 · 10^5, R^2 = 0.9994; P < 2.2 · 10^{-16}) (Fig. 4c).

Figure 4 – The modified Arrhenius equation with the missing term (Equation 10) gives slight

differences in modeled total daily photosynthesis compared to Equation 3 for low R (a,b) and high R

(c,d) under scenarios where H_d is allowed to vary (a, c) and is fixed to 200 kJ mol^{-1} (b,d). "New"

indicates Equation 10, "Old" indicates Equation 3. A: photosynthesis, Low R: low respiration, High R:

high respiration. Black line indicates 1:1 line and grey dashed line indicates respective modeled

slopes and intercepts.

The ratio of the total daily photosynthesis : respiration (A/R) was also considered when comparing

models. Using a fixed H_d of 200 kJ mol^{-1}, the low respiration model of A/R had a slope of 0.993 ± 5.634

· 10^{-3} and the intercept was 0.025 ± 0.002 (approximately 1% of modeled A/R) ($F_{1,898}$ = 3.109 · 10^6, R^2 =

0.9997; P <2.2 · 10^{-16}) (Fig. 5b). The high respiration model of A/R was similar; the slope was 0.993 ±

5.628 and the intercept was 0.014 ± 0.001 (approximately 1% of modeled A/R) ($F_{1,898}$ = 3.115 · 10^6, R^2

= 0.9997; P <2.2 · 10^{-16}) (Fig. 5d). Using a variable H_d, the low respiration model of A/R had a slope of

0.994 ± 0.001 and the intercept was 0.030 ± 0.001 (approximately 1% of modeled A/R) ($F_{1,556}$ = 6.857 ·

10^5, R^2 = 0.9992; P <2.2 · 10^{-16}) (Fig. 5a). The high respiration model had a similar slope of 0.994 ±

0.001 and the intercept was 0.017 ± 0.003 (approximately 1% of modeled A/R) ($F_{1,556}$ = 6.812 · 10^5, R^2

= 0.9992; P <2.2 · 10^{-16}) (Fig. 5c).

Figure 5 – The modified Arrhenius equation with the missing term (Equation 10) gives slight

differences in modeled total daily A/R compared to Equation 3 for low R (a,b) and high R (c,d) under

scenarios where H_d is allowed to vary (a, c) and is fixed to 200 kJ mol^{-1} (b,d). "New" indicates

Equation 10, "Old" indicates Equation 3. A: photosynthesis, R: respiration, Low R: low respiration,

High R: high respiration. Black line indicates 1:1 line and grey dashed line indicates respective

modeled slopes and intercepts.

Lastly, total daily carbon (C) gain was also compared between models with different thermal response

parameters. Using a fixed H_d of 200 kJ mol^{-1}, the low respiration model of total C gain had a slope of

0.978 ± 7.445 · 10^{-4} and an intercept of 0.001 ± 7.609 · 10^{-5} (approximately 2% of modeled C gain)

($F_{1,898}$ = 1.726 · 10^6, R^2 = 0.9995; P <2.2 · 10^{-16}) (Fig. 6b). Likewise, the high respiration of total C gain

had a slope of 0.979 ± 7.261 · 10^{-4} and an intercept of 0.001 ± 1.116 · 10^{-4} (approximately 2% of

modeled C gain) ($F_{1,898}$ = 1.821 · 10^6, R^2 = 0.9995; P <2.2 · 10^{-16}) (Fig. 6d). This can be compared to a

variable H_d; starting with the low respiration model, total C gain had a slope of 0.985 ± 9.555 · 10^{-4}

and an intercept of 0.001 ± 1.223 · 10^{-3} (approximately 1.5% of modeled C gain) ($F_{1,556}$ = 1.06 · 10^6, R^2

= 0.9995; P <2.2 · 10^{-16}) (Fig. 6a). The high respiration of total C gain also had a slope of 0.985 ± 9.110 ·

10^{-4} and an intercept of 0.001 ± 1.290 · 10^{-4} (approximately 1.5% of modeled C gain) ($F_{1,556}$ = 1.17 ·

10^6, R^2 = 0.9995; P <2.2 · 10^{-16}) (Fig. 6c).

Figure 6 – The modified Arrhenius equation with the missing term (Equation 10) gives slight

differences in modeled total carbon gain compared to Equation 3 for low R (a,b) and high R (c,d)

under scenarios where H_d is allowed to vary (a, c) and is fixed to 200 kJ mol^{-1} (b,d). "New" indicates

Equation 10, "Old" indicates Equation 3. C: carbon, Low R: low respiration, High R: high respiration.

Black line indicates 1:1 line and grey dashed line indicates respective modeled slopes and

intercepts.

<u>Discussion</u>

We sought to determine whether the missing term in Equation 3 had a meaningful impact on fitted

temperature response parameters due to its prevalence in photosynthetic temperature response data

and vegetation modeling (Kattge & Knorr, 2007; Duursma & Medlyn, 2012; Rogers et al., 2017; Smith

& Dukes, 2017; Stinziano et al., 2018; Stinziano et al., 2019; Kumarathunge et al., 2019). Our present

analysis suggests that there is no large impact on the fitted temperature response parameters for

V_{cmax} and J_{max}, with differences ranging between 0 and 3.8% depending on the parameter. In general,

fitting Equation 10 instead of Equation 3 results in slightly reduced values for E_a and ΔS, with

essentially no impact for k_{25} and H_d when H_d is fit. When H_d is fixed, there are no meaningful

differences in the fits except for ΔS for J_{max}, where Equation 10 values are ~4% lower than Equation 3

values, and intercepts in the E_a regressions of ~-2 kJ mol^{-1}, which represents <5% differences between

the two equations in most instances. These findings are promising in that one of the parameters to

which modelled carbon gain is particularly sensitive, H_d (Stinziano et al., 2018), is minimally affected

by the missing term. However, since temperature responses are non-linear, small changes in the

shape of an accelerating curve can have a strong impact on the integral of the response (Jensen,

1906). This caused the differences in temperature response parameters to cause ~1.5 – 2.2%

difference in daily C balance. The daily C balance was most affected by the missing term in Equation 3

when comparing models where H_d was fixed, leading to a 2.2% change. Overall, comparisons of low-

respiration to high-respiration scenarios resulted in similar percent changes in daily C balance;

however, the variation was consistently larger under high-respiration across all variables measured.

Similar trends were found when comparing differences in daily photosynthesis and A/R.

Based on the above analysis, the impact of the missing term in the modified Arrhenius equation may

appear negligible based on the parameter outputs. However given that carbon balance is the time

integral of net CO_2 assimilation, this may lead to substantial impacts over a long time period. It is thus

possible that the apparently small differences we observed could accumulate to relatively large

carbon flux errors across large spatial and temporal scales with fluctuating temperatures. A growing

body of literature on using Optimality Theory for estimating V_{cmax25} and J_{max25} (Ali et al., 2016; Walker

et al., 2017; Smith et al., 2019) still relies on Equation 3 for temperature scaling. Given the impacts we

observed, Optimality Theory may be even more successful if the temperature scaling were done

either with Equation 10, or, as we recommend, macromolecular rate theory which is better grounded

in thermodynamics (e.g. Hobbs et al., 2013; Liang et al., 2018).

Since Equation 3 is incorrect, there are three arguments to support moving beyond the modified

Arrhenius paradigm: 1) there are new approaches to modeling biological temperature responses that

are better-grounded in thermodynamics (e.g. Macromolecular Rate Theory; Hobbs et al., 2013; Liang

et al., 2018); 2) uncertainties in model outputs may accumulate due to the missing term across large

spatial and temporal scales, and 3) the modified Arrhenius model as implemented is categorically

incorrect due to a missing term. Therefore, we conclude that future modeling efforts should move

beyond this incorrect paradigm.

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