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Arrhenius Plot Analysis, Temperature Coefficient and Q_{10} Value Estimation for the Effect of Temperature on the Rate of Molybdenum Reduction by *Acinetobacter calcoaceticus* strain Dr.Y12

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ABSTRACT

Molybdenum is a micronutrient that is required as a co-factor for a variety of hydroxylation and redox transfer activities in both animal and plant physiological processes. The potential of over exposure to interfere with the sperm production and egg formation processes in a number of species, including fish, is the biggest danger of excessive exposure. Only recently has it been discovered that it can be utilised as a remediation method for molybdenum-reducing bacteria. The effect of temperature on molybdenum reduction is one of the variables to consider. It is possible to use many different models to estimate the growth rate of microbes on various media based on the temperature being utilised. The Arrhenius model is popular due to the fact that it contains a limited number of parameters. In general, temperature has an effect on the development and metabolic activity of microorganisms on their substrates. Because microorganisms are so tiny, they are very sensitive to changes in their environment's temperature. Growth on molybdenum by Acinetobacter calcoaceticus strain Dr.Y12 is described, with a discontinuous chevron-like graph of apparent activation energy with a breakpoint at 32.66 °C. Regression analysis results suggest that in the lower temperature range of 20-30 °C, growth on molybdenum had an activation energy of 66.48 kJ/mol, whereas at the higher temperature range of 37-45 °C, it had an activation energy of 99.5 kJ/mol. For the examined temperature range (20-30 °C) and (37-45 °C), Q_{10} values of 2.46 and 3.37 and theta values of 1.09 and 1.13 were obtained, respectively. This is study is very useful in predicting the breakdown of molybdenum and the movement of molybdenum during bioremediation.

INTRODUCTION

Our own actions are putting our ecosystem in danger right now. Heavily polluting the environment include heavy industry, urbanization, and agriculture, all of which have increased in tandem with the world's population growth [1–4]. Over exploitation of natural resources, as well as men's ignorance of natural laws, contribute to the escalation of the problem [5–8]. Over the years, the amount of pollution caused by hydrocarbons and metal ions has steadily increased across the globe. Toxic chemicals generated from metals and their compounds have been related to a range of acute and chronic toxicity cases in high-

exposure settings such as the workplace and the environment, according to research. Heavy metals may be present in the environment in their natural state. Heavy metal levels have increased significantly in recent years as a consequence of human activities dating back to pre-industrial times, according to the Environmental Protection Agency [9–13]. A large and indiscriminate release of toxins into the environment is happening in parallel with the increase in population and the intensity of industrial activity. The presence of high quantities of heavy metals over the critical load may have negative consequences for human health and the environment. Metals such as arsenic, cadmium, chromium, cobalt, copper, lead,

mercury, molybdenum, nickel, silver, and zinc are toxic in their elemental forms and different combinations, and they are also non-biodegradable in their elemental forms and various combinations. Metal accumulation in the food chain may represent a major threat to the ecosystem as a consequence of their carcinogenic and mutagenic properties, which are associated with metals. Heavy metal contamination has risen to the level of a global public health emergency in recent years, making it imperative to remove them from the environment as soon as possible [14–19].

Molybdenum is an important trace element that acts as a micronutrient and is required as a co-factor for over 50 enzymes. It promotes cellular activity in animal and plant physiology, for example, by catalysing a range of hydroxylation and redox transfer processes [20-25]. With molybdenum's widespread use in the industrial production of ceramics, glass and contact lens solution, metallurgical processes, lubricants, pigment, catalyst, electronic goods, and as colour additives in cosmetics, the dangers to people exposed to its toxicity have also increased [26– 32]. It has been reported that an increase in the amount of molybdenum in groundwater in mining sites of up to 0.5 mg/L has been found, which is higher than the World Health Organization (WHO) recommended limit of 0.07 mg/L in drinking water [28]. Animals that have had direct contact with molybdenum via drinking water or while foraging for plants are more likely to exhibit hypocuprosis signs or suffer from molybdenosis after a lengthy period of time [22].

Microorganisms are especially susceptible to molybdenum breakdown when exposed to high temperatures because of their small size. Physiology is influenced by temperature, which allows organisms to better adapt to their changing environments. When it comes to biodegrading chemicals, temperature is an essential element to take into consideration. For many years, the Arrhenius model has been widely employed in the study of bacterial growth and rates. It is often used to calculate the apparent activation energy, ΔH*, which is believed to exist for either growth or decay on various metabolic substrates [33–39].

Although being frequently employed in simulating the temperature impact in a limited temperature range, the Arrhenius model is less often used to larger ranges [40]. For most temperature ranges, the value of delta H (Δ H*) is approximately constant. However, for extreme ranges of temperature, this number may diverge three or fourfold depending on the range of temperatures being examined [41]. According to some studies, the model may not be accurate when used across the whole bacterial process temperature [42]. The Arrhenius plot may also display a previously discovered transition which is a rapid change in the activation energy [43].

Arrhenius's model has the fewest parameters, making it relatively universally accepted by researchers [40]. In other words, the Arrhenius models are utilised in understanding how temperature affects bacterial development because of this. The Arrhenius parameter estimate is calculated by drawing a linear regression on the Arrhenius plot. Several years ago, similar research looked at Q_{10} value estimates of Arrhenius plot analysis and impact of temperature on molybdenum growth done by *Pseudomonas* sp. strain DRYJ7 [44]. Other competing model, the Ratkowsky, is also built on the assumption of linear growth, but due to biological foundations, this model suffers from a lack of steady development and exhibits non-linear behaviour [45].

This research showed that there were many possible activation energies for the breakdown of molybdenum by a bacterium, which was previously unknown. It is interesting in terms of concepts, and it will also be extremely helpful in forecasting molybdenum removal and transport during bioremediation.

MATERIALS AND METHODS

The activation energy of growth on molybdenum by *Acinetobacter calcoaceticus* strain Dr.Y12

Acinetobacter calcoaceticus strain Dr.Y12 was grown and maintained in a low phosphate media (LPM) composed of magnesium sulphate pentahydrate MgSO4.7H2O (0.05%), disodium molybdate dihydrate Na2MoO4.2H2O (0.242 % or 10 mM), glucose (1%), ammonium sulphate (NH4)2.SO4 (0.3%), sodium chloride NaCl (0.5%), yeast extract (0.5%), and disodium phosphate anhydrous Na2HPO4 (0.071% or 5 mM) [46]. Molybdenum reduction rate data from Acinetobacter calcoaceticus strain Dr.Y12 was then processed as previous [39] by transferring the growth values at each temperature to the natural logarithm and calculating the value of the slope, which is equivalent to specific growth rate.

The Arrhenius equation [47] is as follows,

$$\mu = Ae^{-rac{E_a}{RT}}$$
 [eqn. 1]

Where T is the absolute temperature (Kelvin = ${}^{O}C + 273.15$), R is the universal gas constant (0.008314 kJ/molK-1), Ea is the activation energy (kJ/mol) and A physically signifies the rate constant at which all the participating molecules possess sufficient energy prior reaction (Ea = 0). A linearized form is given via the plot of log normal growth rate against 1/T and the equation is as follows;

$$\ln \mu = \ln A - \frac{E_a}{R} \cdot \frac{1}{T}$$
 [eqn. 2]

Coefficient of $Q_{I\theta}$ estimation

The Q_{10} value is estimated via the following equation;

$$Q_{10}=e^{\left(rac{Ea}{R}
ight)\left(rac{10}{T_2T_1}
ight)}$$
 [eqn. 3]

Following rearrangement,

$$\ln Q_{10} = \left(\frac{E_a}{R}\right) \left(\frac{1}{T_1 T_2}\right)$$
 [eqn. 4]

The coefficient of temperature or theta (θ) value (simplified Arrhenius temperature coefficient) is another important biological constant obtained from substitution of the obtained values into the reaction rates equation governed by the O_{10} rule;

$$KT = K20\Theta (T-20)$$
 [eqn. 5]

RESULTS AND DISCUSSION

The effect of temperature on the growth rate of the bacterium on molybdenum shows an increasing growth rate leading to a maximum rate at 30 °C and a decrease of growth rate at higher temperatures (**Fig. 1**). When plotting ln µm versus 1/T, we got a Chevron-like graph, which showed a discontinuous curve across the entire temperature range (**Fig. 2**). An interesting finding was the presence of a break point at 32.66 °C. Regression analysis results shown in **Table 1** and **Table 2** suggest that in the lower temperature range of 20-30 °C, growth on molybdenum had an activation energy of 66.48 kJ/mol, whereas at the higher temperature range of 37–45 °C, it had an activation energy of 99.54 kJ/mol. A previous study on the growth rate of *Pseudomonas sp.* strain DrYJ7 between 10 and 20 °C on molybdenum showed activation energy of 14.96 Kj/mol [44], which is much lower.

Activation energy estimated using the Arrhenius model was within the published literature's range of activation energy for various biodegradation of xenobiotics (Table 2). The connections seem to require more energy to break apart. Increasing the temperature uses less energy. Of the many reports on the activation energy calculated from rates of metabolic process at various temperatures, very few works report on the presence of two activation energies opting instead to report for only one activation energy spanning a large range of temperature. Of the reports, two contrasting difference is seen wherein one study, a higher activation energy is reported at higher temperatures compared to a lower range of temperature while in another study, an opposite phenomenon is observed (Table 3). A case in point is the growth of Bacillus sp. JF8 on polychlorinated biphenyl (PCB) where the activation energy was 12.1 Kj/mol from 20 to 46 °C and 31.4 Kj/mol from 50 to 70 °C [48]. In another contrasting study, the growth of phenol by Pseudomonas sp. AQ5-04 shows activation energy of 38.92 Kj/mol from 15 to 30 °C and 11.34 Kj/mol from 35-45 °C [38].

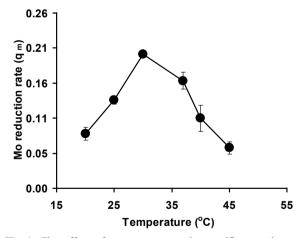


Fig 1. The effect of temperature on the specific growth rate of *Acinetobacter calcoaceticus* strain Dr.Y12 on molybdenum. Error bars represent mean ± standard deviation (n=3).

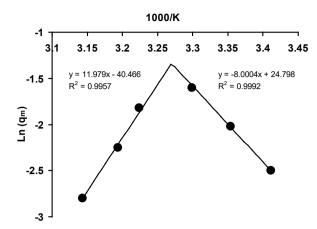


Fig 2. Arrhenius plot of the molybdenum reduction rate by *Acinetobacter calcoaceticus* strain Dr.Y12.

Table 1. The two-part linear regression analysis for the Arrhenius plot of molybdenum reduction rate by *Acinetobacter calcoaceticus* strain Dr.Y12.

Temperature range °C 20,25,30 Regression equation $y = -8.0004x + 24.798$ Coefficient of determination 0.99 tan $a \pm S$ tandard error -8.00 ± 0.23 $E_a \pm S$ tandard error, kJ mol ⁻¹ 66.48 \pm 1.91
Regression equation $y = -8.0004x + 24.798$ Coefficient of determination 0.99 tan $a \pm Standard error$ -8.00 ± 0.23 $E_a \pm Standard error$, kJ mol $^{-1}$ 66.48 ± 1.91
Coefficient of determination 0.99 $\tan a \pm \text{Standard error}$ -8.00 ± 0.23 $E_a \pm \text{Standard error}$, $kJ \text{ mol}^{-1}$ 66.48 ± 1.91
tan a \pm Standard error -8.00 ± 0.23 $E_a\pm$ Standard error, kJ mol ⁻¹ 66.48 ± 1.91
$E_a \pm \text{Standard error, kJ mol}^{-1}$ 66.48±1.91
t-Statistic -34.73
Degrees of freedom 2
Left part
Temperature range °C 37,40,45
Regression equation $y = 11.979x - 40.466$
Coefficient of determination 0.99
tan a \pm Standard error 11.98 \pm 0.79
$E_a \pm \text{Standard error, kJ mol}^{-1}$ 99.54±6.58
t-Statistic 15.13
Degrees of freedom 2
Break points data
Intersection coordinates, (x, y) 3.27,-1.34
Break point temperature °C 32.66
Q_{10} (20-30 °C) 2.46
Theta (20-30 °C) 1.09
$Q_{10}(37-45 ^{\circ}\text{C})$ 3.37
Theta (37-45 °C) 1.13

Table 2. Summary of nonlinear regression of the effect of temperature on the rate of molybdenum reduction *Acinetobacter calcoaceticus* strain Dr.Y12.

Segmental linear regression	
Best-fit values	
intercept1	-39.08
slope1	11.54
X_0	= 3.270
slope2	-8.236
Std. Error	
intercept1	1.314
slope1	0.4104
slope2	0.3676
95% CI (asymptotic)	
intercept1	-43.26 to -34.90
slope1	10.24 to 12.85
slope2	-9.406 to -7.066
Goodness of Fit	
Degrees of Freedom	3
R squared	0.9963
Sum of Squares	0.003631
Sy.x	0.03479
Constraints	
X_0	$X_0 = 3.27$

Table 3. Arrhenius temperature characteristics for metal reduction.

Microorganisms	Tempe- Substrate rature range		ΔH^* apparent activation Ref energy (kJ.mol ⁻¹)	
	(°c)			
Ochrobactrum	25-35	Chromate	120.69	[49]
intermedium				
BCR400				
Arthrobacter sp. SUK	25-60	Chromate	36.21	[50]
1201				
Aspergillus niger	30-60	Chromate	8.56	[51]
Bacillus sp.	25-40	Chromate	22.0	[52]
Thermus scotoductus	65	Chromate	35 (membrane bound	[53]
SA-01			enzyme)	
			40.3 (soluble)	
Thermus scotoductus	60-65	Iron	30	[53]
SA-01				
Shewanella profundo	ı 4-37	Iron	50.3	[54]
LT13a				
β-Proteobacteria	15-40	Vanadate	36	[55]
Shewanella	25-40	Selenate	Control system 62.90	[56]
oneidensis · MR-1			TPPS-supplemented	
			system 47.33	
			•	
Acinetobacter	20-45	Molybdate	66.48 (20-30 °C)	This
calcoaceticus strair	ı	-	99.54 (37-45 °C)	study
Dr.Y12			, ,	,

Note: N(TPPS) Meso-tetrakis (4-sulfonatophenyl) porphyrin mediator

The higher the activation energy, the more energy the bacterium needs to use to metabolize xenobiotics. Based on **Table 3**, the values obtained in this study for both temperature ranges are within the activation energy for numerous metal reduction by microbial species. However, the activation energy for the typical mesophilic bacteria is between 33.5 and 50.3 kJ/mol [57], indicating that the activation energy for one of the temperature range studied in this study was relatively higher. The higher activation energy for the higher range of temperature was within the range reported by the chromate-reducing *Ochrobactrum intermedium* BCR400 (49) (**Table 3**).

In the current study, we found that the activation energy is not constant, rather it depends on the temperature chosen [58]. While we can't accurately estimate all of the interacting complex biological processes that are taking on at the same time, the model functions as an observational model. Activation energy thus should not be thought of as the activation energy utilised in chemical processes, but rather the total temperature response of the microorganism [59].

Even with these problems, the model is in use worldwide. The activation energy, which depends on the temperature change, plays an important role in the metabolic activity of microorganisms, and it has been shown in a variety of conditions not limited to metal-reducing activity and include processes such as the decolourization of various dyes [43,60–63], chromate reduction [49,64] and phenolics biodegradation [38,40,65–67] and molybdenum [44].

The details of the process that causes the change are still unknown, but two hypotheses provide two plausible explanations. The first is that water characteristics change as it transitions and a hypothesis of "bottleneck" hypothesises that a limited number of events occur simultaneously in rapid succession [68]. Based on various measured Arrhenius breakpoint temperatures, the first theory does not seem to be correct [43]. Following the "bottle-neck" idea, since each of the chained enzymes has its unique thermal characteristics, it is impossible to verify the "bottle-neck" hypothesis.

When taking into consideration the ambient temperature, the cell membrane will also vary [69]. The "bottleneck" theory continues to hold strong among academics [43,70]. Alternatively, the Arrhenius plots may be used to estimate the Q_{10} values, or they can be calculated by measuring the rates of growth for different incubation temperatures with ten degrees of variation [71]. When the bioreduction and growth rates have been logarithmically plotted against 1000/temperature (Kelvin), the Arrhenius curve is the slope of the resulting plot (**Fig. 1**).

For the examined temperature range (20-30 °C) and (37-45 °C), $Q_{I\theta}$ values of 2.46 and 3.37 were obtained, respectively. However, since biological processes are dynamic, there may be more than one $Q_{I\theta}$ value for a distinct temperature range being investigated. In the reduction of molybdate to molybdenum blue, a 2.038 value was obtained [72] while in another molybdenum reducer; Morganella sp, a $Q_{I\theta}$ value of 2.31 was obtained. When attributing the growth process to a distinctive biological activity, this value is essential. $Q_{I\theta}$ was calculated to be 2.7 for oil biodegradation in a beach gravel column in previous studies [73]. However, different research on soil polluted with decane and toluene shows a $Q_{I\theta}$ value of 2.2 [74].

Both bacteria's ability to break down petrochemicals and the effects of temperature on it were determined to have a Q_{10} of 2.2 [75], while, immobilized bacterial systems at temperatures ranging from 25 and 45 degrees Celsius produce molybdenum and its Q_{10} value is 2.8 [76]. Increasing the value of Q_{10} as the temperature decreases is often true [77,78]. In another research, *Pseudomonas* sp. strain AQ5-04 produced a Q_{10} value of 1.834 (38) while a Q_{10} value of 2.17 was calculated for the growth rate of this organism on molybdenum. A lower Q_{10} value of 2.17 is reported in another study on molybdenum reduction [44].

For the examined temperature range (20-30 °C) and (37-45 °C), theta values of 1.09 and 1.13 were obtained, respectively (**Fig. 3** and **Fig. 4**), which was similar to a theta value of 1.08 calculated for the molybdenum reduction by the bacterium *Serratia* sp. strain HMY1 [72]. In the growth rate on molybdenum by the Antarctic bacterium *Pseudomonas* sp. strain DRYJ7, a theta value of 1.03 was obtained [44]. The theta value is also within the range for many biological processes that are from 1.1 to 1.7 although higher values of up to 16.2 have been reported for the degradation of other xenobiotics [79].

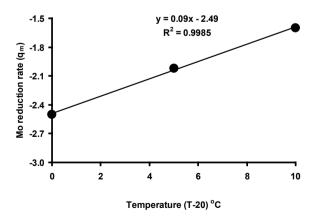


Fig. 3. Estimation of theta value for rate of Mo-reduction within the temperature range of 20 to 30 $^{\circ}$ C.

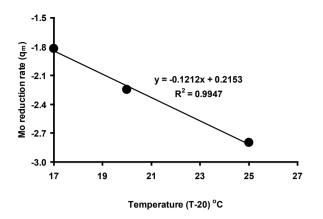


Fig. 4. Estimation of theta value for rate of Mo-reduction within the temperature range of 37 to 45 $^{\circ}$ C.

CONCLUSION

This is the first study demonstrated that the activation energy needed for the biodegradation of molybdenum by a bacterium which displays a broken profile with two activation energies observed in the Arrhenius plot. Temperature generally affects microbial growth and metabolic activity on their substrates. The small nature of microbes makes them susceptible to change in the surrounding temperature. Growth on molybdenum by Acinetobacter calcoaceticus strain Dr.Y12 is described, with a discontinuous chevron-like graph of apparent activation energy with a breakpoint at 32.66 °C. Regression analysis results suggest that in the lower temperature range of 20-30 °C, growth on molybdenum had an activation energy of 66.48 kJ/mol, whereas at the higher temperature range of 37-45 °C, it had an activation energy of 99.5 kJ/mol. For the examined temperature range (20-30 °C) and (37-45 °C), Q₁₀ values of 2.46 and 3.37 and theta values of 1.09 and 1.13 were obtained, respectively. The quantum, especially in between 15 and 20 °C, is relatively a bit higher than the typical energies observed in mesophilic microorganisms. The amide bond is postulated to hold much higher activation energy to be broken. Additional work is under investigation, particularly on parameters themselves, to determine the effects of temperature on growth kinetics. The values obtained in this work are within the normal range for many biological processes. The values obtained in this work are within the normal range for many biological processes.

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