

Original Research Article

Mathematical Modelling of Hydrophilic Antibiotic Concentration, Binding Dynamics, and Drug Resistance under Competitive Inhibition in Disc Diffusion Assays

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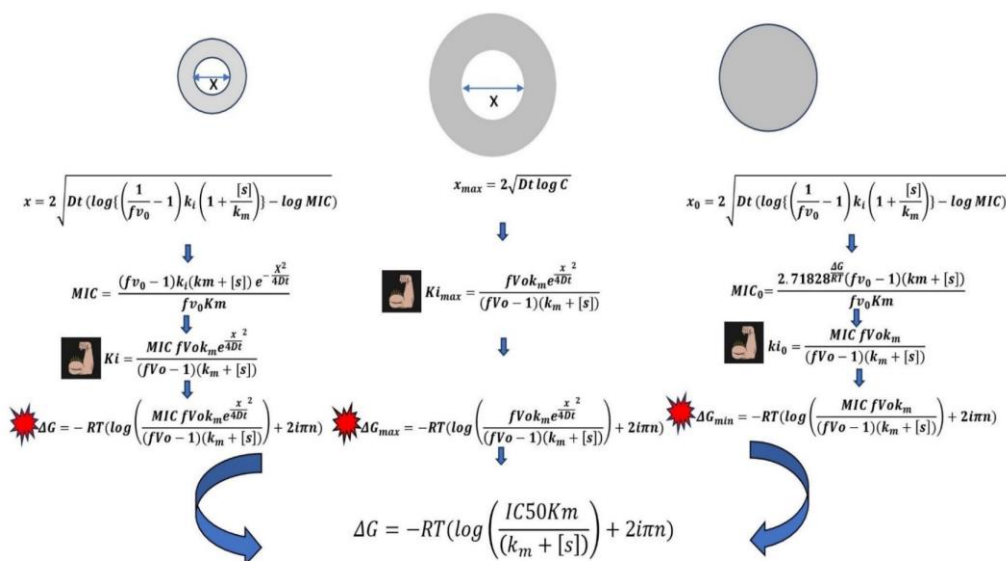
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Graphical Abstract



KEYWORDS	ABSTRACT:
Minimum inhibitory concentration, Zone of Inhibition, Gibbs Free energy, inhibition constant. Cheng–Prusoff, Binding affinity and Antibiotic Resistance	Understanding the intricate relationship between minimum inhibitory concentration binding Affinity and drug resistance is paramount for developing effective antimicrobial strategies, in this paper, we explore a mathematical relationship between these factors, under the condition of the disc diffusion method of antimicrobial susceptibility testing of antibiotics. In our study we considered the response as (ZOI) Zone of Inhibition denoted by "X" in mm, the response X is directly proportional to the concentration of antibiotics, Zone of Inhibition X is directly proportional to binding affinity and ΔG binding free energy of receptor. X is directly proportional to the -log of minimum inhibitory concentration (MIC), which contributes to resistance upon certain values by our mathematical derivation, we tried to give the direct relationship among binding affinity, minimum inhibitory concentration (MIC), zone of inhibition (ZOI) i.e. X and antibiotic concentration 'c' on the hypothesis of enzymatic kinetics the dimensions of Cheng–Prusoff, binding affinity, IC50, were calculated. For hydrophilic antibiotics following competitive inhibition kinetics.

1. Introduction

A traditional technique for figuring out the minimum inhibitory concentration (MIC) in semisolid media is agar plate diffusion [1]. The antibiotic is diffused into the bacterially loaded agarose medium using the disc diffusion method, which inhibits bacterial growth around the source and creates a clear zone devoid of bacterial lawns [2]. The diameter of these zones is directly correlated with the antibiotic concentration. A graph between inhibition zone radii x mm squared values x^2 are plotted against the logarithm of antibiotic concentration. The Intercept of linear regression gives the value of MIC at the squared size of these zoi[3].proposed by Boyan and James (2008)

2. Derivation

$$\ln \ln MIC = \ln \ln C - \frac{x^2}{4Dt} \dots \dots \dots eq(1)$$

Here “D” is the diffusion coefficient and it is not dependent on the “C” concentration of antibiotic and “t” the time of diffusion

Rearranging the equation (1) in terms of zoi = X

$$x = 2\sqrt{DtC - \log \log MIC} \dots \dots \dots eq(2)$$

In Case of competitive Inhibitor, [I]



Initial rate of reaction V_0 is given by [4]

$$v_0 = \frac{V[S]}{k_m \left(1 + \frac{[I]}{k_i}\right) + [S]} \dots \dots \dots eq(3)$$

Where K_m is Michaelis-Menten constant $[S]$ substrate concentration $[I]$ is the concentration of inhibitor, and V_{max} represents the maximum velocity achieved by the system, at maximum (saturating) substrate concentrations.

Rearranging the above equation for inhibitor concentration $[I]$

$$[I] = - \left(\frac{k_i(K_m v_0 + [S](v_0 - V_{max}))}{v_0} \right) \dots \dots \dots eq(4)$$

K_i = dissociation constant for the inhibitor [6] k_{-3}/k_3

To compute the concentration of competitive inhibitor [7] $[I]$ that yields the fraction $f v_0$ of velocity v_0 where $0 < f v_0 < 1$

$$[I] = \left(\frac{1}{f v_0} - 1 \right) k_i \left(1 + \frac{[S]}{k_m} \right) \dots \dots \dots eq(5)$$

K_i value can be interpreted as dissociation constant K_D of competitively binding inhibitor [8]

Substituting the value of equation (5) in equation (2)

$$x = 2 \sqrt{\left(\frac{1}{f v_0} - 1 \right) k_i \left(1 + \frac{[S]}{k_m} \right) \} - \log \log MIC} \dots \dots \dots eq(6a)$$

$$\therefore C = \left(\frac{1}{f v_0} - 1 \right) k_i \left(1 + \frac{[S]}{k_m} \right)$$

“C” concentration of antibiotic that can be termed as *an inhibitor* of the enzyme

Zone of inhibition “zoi” is the determining and critical factor in deciding the antimicrobial susceptibility [9] of antibiotic zoi x (directly proportional) $\propto \frac{1}{f v_0}$ where $f v_0$ is the ratio of $\frac{v_0}{v_i}$ where v_0 is the velocity without inhibitor and v_i is the velocity with inhibitor. The value of $f v_0$ decreases in two cases a) when v_0 rate of reaction between enzyme and substrate decreases [10] b) when v_i rate of reaction between inhibitor and enzyme gets increased, it will increase the value of zoi 'x' and decrease the value of MIC minimum inhibitory concentration, which means less inhibitor concentration is required to achieve the desired ZOI similarly, zoi may decrease if the value of $f v_0$ is increased in the denominator as a function of binding affinity of the enzyme with its natural substrate is more favouring for product formation and v_i velocity with inhibitor is relatively lower, it will increase MIC value, more inhibitor concentration will be required to inhibit the product formation. Inhibition constant K_i is inversely proportional to binding affinity [11], which means the higher the value of k_i , the lower the value of binding affinity that means the inhibitor is not tightly bound with the enzyme less the inhibitor will compete with the active site and that lowers the response of zoi and *vice-versa* [12].

As the k_m value increases to maintain maximum velocity V_{max} in case of competitive inhibition.

Substituting the value of $f v_0$ with $\frac{v_0}{v_i}$ equation (6a) becomes (6b)

$$x = 2 \sqrt{\left(\frac{v_i - v_0}{v_0}\right) k_i \left(1 + \frac{[s]}{k_m}\right) \} - \log \log MIC) \dots \dots \dots eq(6b)$$

Rearranging the equation (6a) for minimum inhibitory concentration MIC

$$MIC = \frac{(f v_0 - 1) k_i (k_m + [s]) e^{-\frac{x^2}{4Dt}}}{f v_0 K_m} \dots \dots \dots eq(7)$$

As postulated from the equation MIC is directly proportional to the exponent $e^{-\frac{x^2}{4}}$. This is the form of Gaussian function [13]. $e^{-\frac{x^2}{4}}$ that is the negative square of the zone of inhibition (zoi) by 4 as the zoi increases the minimum inhibitory concentration decreases exponentially ($MIC = -x^2/4$). The value of inhibition constant K_i influences the MIC directly [14], smaller k_i represents greater binding affinity which means smaller amount of ligand (inhibitor) is needed to inhibit the activity of the enzyme, which lowers the value of minimum inhibitory concentration. A high value of K_i leads to less binding affinity hence more the value of MIC. In competitive inhibition inhibitors and substrate compete for the same active site.

In competitive inhibition, V_{max} remains unchanged and K_m increases and shifts towards the right, as the K_m gets increased, affinity of the enzyme with its substrate decreases and the affinity of the enzyme with the inhibitor increases so MIC decreases and vice-versa

$$MIC = \frac{\frac{\Delta G}{RT} e^{-\frac{x^2}{4Dt}} (f v_0 - 1) (k_m + [s])}{f v_0 K_m} \text{ Since } k_i = \text{Exp} \left(\frac{\Delta G}{RT} \right) \dots \dots \dots eq(8a)$$

$$MIC = e^{\frac{\Delta G}{RT} - \frac{x^2}{4Dt}} \left(\frac{1}{f v_0} - 1 \right) \left(1 + \frac{[s]}{k_m} \right) \dots \dots \dots eq(8b)$$

Since value of $e = 2.71828$

$$MIC = \frac{2.71828^{\frac{\Delta G}{RT} - \frac{0.25x^2}{Dt}} (f v_0 - 1) (k_m + [s])}{f v_0 K_m} \dots \dots \dots eq(8c)$$

In the above equation Michaelis-Menten K_m unit is mole, R gas constant in kcal/molK and T is Temperature in Kelvin/

Rearranging the equation (6a) concerning the K_i inhibition constant

$$K_i = \frac{MIC f v_0 k_m e^{\frac{x^2}{4Dt}}}{(f v_0 - 1) (k_m + [s])} \dots \dots \dots eq(9)$$

$$k_i = \text{Exp} \left(\frac{\Delta G}{RT} \right) \dots \dots \dots \text{eq}(10)$$

Where k_i = inhibition constant [15]

ΔG = Binding Energy kcal/mol

R = Gas Constant 1.985×10^{-3} kcal/mol K

T = Temperature 298.15K

Here the inhibition constant k_i was obtained from the binding energy by using the formula as mentioned in equation (10)

Substituting the value of K_i of equation (10) to equation (6) the value of $zoi = x$ is given by

$$x = 2 \sqrt{\left(\frac{1}{f v_0} - 1 \right) \text{Exp} \left(\frac{\Delta G}{RT} \right) \left(1 + \frac{[s]}{k_m} \right) \} - \log \log \text{MIC}} \dots \dots \dots \text{eq}(11)$$

Substituting the value of K_i of equation (10) in equation (9) we get

$$\Delta G = -RT \left(\log \left(\frac{\text{MIC } f V o k_m e^{\frac{x^2}{4Dt}}}{(f V o - 1)(k_m + [s])} \right) + 2i\pi n \right) \dots \dots \dots \text{eq}(12)$$

Since $\Delta G = RT(\log \log (k_i) + 2i\pi n)$

The value ΔG is the binding energy, R is gas constant and $\left(\frac{\text{MIC } f V o k_m e^{\frac{x^2}{4Dt}}}{(f V o - 1)(k_m + [s])} \right)$ is inhibition constant

k_i , T is the temperature in kelvin. The function $2i\pi n$ is the periodic function [16], accounts for the multiple branches of the logarithm function in the complex plane, which states that the ΔG may have finite values depending on the integer “n” by setting $n=0$ the periodic influence of the term $2i\pi n$ can be eliminated, hence a constant value of ΔG can be achieved at a particular T temperature. The equation ΔG (directly proportional) $\propto -\log \text{MIC}$ indicates a mathematical relationship where change in the variable ΔG (binding energy) is inversely related to the $\log \text{MIC}$, this implies that as the value of MIC increases due to resistance, the term $\log \text{MIC}$ grows positively leading to the decrease in ΔG .

The function shows the logarithmic decline in the value of binding energy in response to increase in minimum inhibitory concentration MIC.

In condition the value of ΔG binding energy increases the value of MIC. the left-hand side of the equation becomes more positive. Since ΔG is defined as $-\log \text{MIC}$, an increase in ΔG implies that $-\log \text{MIC}$ becomes increasingly positive, which in turn means that $\log \text{MIC}$ must become increasingly negative. This indicates that MIC must decrease because as the logarithm of a positive number becomes more negative.

Rearranging the equation (12) in terms of ΔG is the binding energy

$$\Delta G = -RT \left(\log \log \left(\frac{\text{MIC } f V o k_m e^{\frac{2Dt\sqrt{x}}{4Dt}}}{(f V o - 1)(k_m + [s])} \right) + 2i\pi n \right) \dots \dots \dots \text{eq}(13)$$

on simplifying the equation (11)

$$x = 2 \sqrt{Dt \log \left[\frac{\left(\frac{1}{f v_0} - 1 \right) (k_m + [S])}{k_m} \right] - \log \log MIC} \dots \dots \dots eq(14)$$

When the IC50 value is known, the Cheng–Prusoff [17] equation is frequently applied to determine the equilibrium dissociation constant (Kb) of a competitive antagonist.

$$Ki = \frac{IC50}{1 + \frac{[S]}{Km}} \dots \dots \dots eq(15)$$

Substituting the value of Ki of equation (17) in equation (6)

$$x = 2 \sqrt{\left(\frac{1}{f v_0} - 1 \right) \frac{IC50}{1 + \frac{[S]}{Km}} \left(1 + \frac{[S]}{k_m} \right) \} - \log \log MIC} \dots \dots \dots eq(16)$$

The term $1 + \frac{[S]}{Km}$ cancelled out from the equation

$$x = 2 \sqrt{\left(\frac{1}{f v_0} - 1 \right) IC50 \} - \log \log MIC} \dots \dots \dots eq(17)$$

Rearranging the above equation (19) for IC50

$$IC50 = \frac{f v_0 MIC e^{\frac{x^2}{4Dt}}}{(f v_0 - 1)} \dots \dots \dots eq(18)$$

Driving the above equation for MIC by equation (20)

$$MIC = \frac{IC50(f v_0 - 1)e^{-\frac{x^2}{4Dt}}}{f v_0} \dots \dots \dots eq(19)$$

MIC is the minimum inhibitory [18] concentration used to determine the lowest concentration of the substance that inhibits visible growth of microorganism

The term $e^{-x^2/4}$ this term involves the exponential function with negative argument, as ZOI ‘x’ increases the value of MIC decreases. The factor 4 in the denominator inside the exponent makes this change more sensitive

The diffusion coefficient ‘D’ represents the rate at which molecules diffuses through the medium. The increased rate of diffusion can potentially lead to more effective inhibition of microbial growth thereby radius of MIC

IC₅₀ half maximal concentration is directly related to the MIC as they both measure the potency of antibiotic in inhibiting the growth of microorganism

A lower IC₅₀ generally corresponds to a lower MIC, indicating the greater potency and effectiveness in inhibiting microbial growth.

In the condition of resistance where the zoi x_0 , $x=0$ or $x \geq 0$ then the exponent $e^{-\frac{x^2}{4Dt}}$ becomes one, the value of inhibition constant Ki_0 , minimum inhibition concentration MIC_0 , and change in binding energy ΔG_{min} is given by.

$$x_0 = 2 \sqrt{\left(\frac{1}{fv_0} - 1\right) k_i \left(1 + \frac{[s]}{k_m}\right) \{-\log \log MIC\}} \dots (20) \text{ (where } x=0 \text{ or } x \geq 0)$$

$$MIC_0 = C \dots \dots (21)$$

The minimum inhibitory concentration (MIC) is almost equal Antibiotic concentration “c”

$$MIC_0 = \left(\frac{1}{fv_0} - 1\right) k_i \left(1 + \frac{[s]}{k_m}\right) \dots \dots \dots eq(22a)$$

$$MIC_0 = \frac{k_i(fv_0-1)(k_m+[s])}{fv_0K_m} \dots \dots \dots eq(22b) \text{ where } k_i = e^{\frac{\Delta G}{RT}}$$

$$MIC_0 = \frac{2.71828^{\frac{\Delta G}{RT}}(fv_0-1)(k_m+[s])}{fv_0K_m} \dots \dots \dots eq(23) \text{ (Where } e=2.7182)$$

$$ki_0 = \frac{MIC fv_0k_m}{(fv_0 - 1)(k_m + [s])} \dots \dots \dots eq(24)$$

Here the ki_0 Value is given when $x=0$ or $x \geq 0$ this reveals that the enzyme had loses the binding affinity with the inhibitor it is the critical value with no zoi or negligible zoi showing resistance to Antibiotics

$$\Delta G_{min} = -RT(\log \left(\frac{MIC fv_0k_m}{(fv_0 - 1)(k_m + [s])}\right) + 2i\pi n) \dots \dots \dots eq(25)$$

The binding energy is at its critical value denoted by ΔG_{min} at it lowest below which no zoi will be detected as no microorganism killed on nutrient agar plate, the exponent $e^{-\frac{x^2}{4Dt}}$ must have certain value (x) so that ΔG in negative should have certain value more than ΔG_{min} to show some antimicrobial activity and zoi to be appeared. In equation (25)

$Ki = \left(\frac{MIC fv_0k_m}{(fv_0-1)(k_m+[s])}\right)$ is the Inhibition constant, and R is the gas constant T

is temperature and $2i\pi n$ is the periodic function.

Substituting the value zoi of $x=0$, or $x \geq 0$ in equation (19) the condition of resistance to antibiotic is given by.

$$x = 2 \sqrt{\left(\frac{1}{fv_0} - 1\right) IC_{50} \} - \log \log MIC_0} \dots \dots \dots eq(26)$$

$$\left(\frac{1}{fv_0} - 1\right) IC_{50} = MIC_0 \dots \dots \dots eq(27)$$

$$MIC_0 = \frac{IC_{50}(fv_0 - 1)}{fv_0} \dots \dots \dots eq(28)$$

$$IC_{50} = \frac{fv_0 MIC}{(fv_0 - 1)} \dots \dots \dots eq(29)$$

Relative comparison of MIC with resistant and non-resistant species

When $x \geq 0$, then Value of MIC_0 is given by eq (28) the value of minimum inhibitory concentration

MIC (without resistance) is always lower than MIC_0 (with resistance) by the factor of $e^{-\frac{x^2}{4Dt}}$

therefore $MIC = (MIC_0) e^{-\frac{x^2}{4Dt}}$ can be written as

$$MIC = MIC_0 \exp\left(-\frac{x^2}{4Dt}\right) \dots \dots \dots eq(30)$$

The magnitude of resistance between resistance strain of microorganism and non-resistance strain of microorganism can be explained mathematically by x' it is the difference of zoi in between resistance and non-resistance strain of microorganism subjected to be treated by same antimicrobial agent and same species with different strains.

$$x' = \sqrt{\log\left(\frac{MIC}{MIC_0}\right) + 2i\pi n 2\sqrt{Dt}} \dots \dots \dots eq(31)$$

putting $n=0$ the value of $2i\pi n = 0$

$$x' = \sqrt{\log\left(\frac{MIC}{MIC_0}\right) 2\sqrt{Dt}} \dots \dots \dots eq(32)$$

$$\text{rearranging the eq(32) } x' = \sqrt{\log(MIC) - \log(MIC_0) 2\sqrt{Dt}} \dots \dots \dots eq(33)$$

$$x' = 2\sqrt{Dt(\log(MIC) - \log(MIC_0))} \dots \dots \dots eq(34)$$

This value of x zoi will give the difference between resistance and non-resistance strains and the where $MIC_0 > MIC$

Ideal Condition

While considering a hypothetical condition where MIC value is negligible then eq (2) becomes

$$x_{max} = 2\sqrt{DtC - \log \log MIC} \because \log MIC \text{ is negligible}$$

$$x_{max} = 2\sqrt{Dt \log \log C} \dots \dots \dots eq (35)$$

The eq (35) it is the hypothetical case where the MIC is negligible and all the concentration of drug converts to the response x_{max} the maximum response of zoi at negligible value of log MIC. For maximum response $K_{i \max}$ is given by

$$K_{i \max} = \frac{fVok_m e^{\frac{x^2}{4Dt}}}{(fVo - 1)(k_m + [s])} \dots \dots \dots eq(36)$$

Similarly, max energy without MIC is given by

$$\Delta G_{max} = -RT \left(\log \left(\frac{fVok_m e^{\frac{x^2}{4Dt}}}{(fVo - 1)(k_m + [s])} \right) + 2i\pi n \right) \dots \dots \dots (37)$$

Gibbs-Cheng equation for competitive inhibition.

The intensity of an inhibitor in preventing an enzyme's activity is measured by the inhibition constant (Ki). It is a particular type of equilibrium dissociation constant that shows how well an inhibitor binds to its specific enzyme [19]

The relationship between the Gibbs free energy change and the inhibition constant can be expressed mathematically by the equation

$$\Delta G = -RT \ln Ki \dots \dots \dots eq(38)$$

This relationship implies that as the inhibition constant decreases (indicating stronger inhibition), the Gibbs free energy change becomes more negative, reflecting increased stability of the enzyme-inhibitor complex[20]

$$Ki = \frac{IC50}{1 + \frac{[s]}{K_m}} \dots \dots \dots eq(39)$$

$$k_i = \text{Exp} \left(\frac{\Delta G}{RT} \right) \dots \dots \dots eq(40)$$

$$\text{Exp} \left(\frac{\Delta G}{RT} \right) = \frac{IC50}{1 + \frac{[s]}{K_m}} \dots \dots \dots eq(41)$$

The IC50 is the concentration of an inhibitor at which the enzyme's activity is reduced by 50%. A lower IC50 indicates a more potent inhibitor. Km represents the substrate concentration at which the reaction rate is half of its maximum value. It reflects how tightly the enzyme binds to the substrate. [S] is the concentration of substrate available for the enzyme.

$$\Delta G = -RT \left(\log \left(\frac{IC50K_m}{(k_m + [s])} \right) + 2i\pi n \right) \dots \dots \dots eq(42)$$

ΔG is the change in free energy, which tells us if the reaction or process is favourable (negative ΔG = spontaneous) or unfavourable (positive ΔG = non-spontaneous).

The numerator $IC_{50} \cdot K_m$ combines the effect of inhibitor potency (IC_{50}) and the enzyme-substrate binding affinity (K_m). A lower IC_{50} (stronger inhibition) increases this value. A lower K_m (tighter substrate binding) also increases this value. The denominator $k_m + [S]$ captures the contribution of the substrate concentration ($[S]$) and another kinetic factor, k_m . As $[S]$ increases, the denominator grows, reducing the overall value of the fraction, which could mean a lower ΔG . The k_m term represents the intrinsic catalytic activity or background rate, and can act as a baseline in the absence of a high substrate concentration. ΔG becomes more negative (favorable) as IC_{50} decreases (stronger inhibition), K_m decreases (higher substrate affinity), $[S]$ increases (more substrate present).

3. Discussion

A mathematical model of antibiotics concentration, binding energy, minimum inhibition concentration, Inhibition constant and IC_{50} is developed. It is a novel approach to incorporate the kinetics in the equation (2) Boyan and James (2008) for competitive Inhibition.

On introducing Inhibitory concentration as a function of $f v_0$ fractional velocity, inhibitory concentration K_i and substrate /Michales Menten. ratio.

Further Inhibition constant K_i is substituted with the equation $\Delta G = -RT \log K_i$ and K_i was substituted by Cheng-Prusoff, The equation reflecting the two dimensions in term of ΔG binding free energy and IC_{50} Inhibitory concentration 50%, by eq (6b) Zoi is related to the square root logarithmic term changes in Michales Menten will directly affect the value of “x”, in two cases scenario a) Michales Menten will change according to the kind and nature of antibiotics use. b) on fixing the antibiotic, value of Michales Menten is constant for that particular moiety. If the K_m value increases the ratio of $[S]/k_m$ decreases, this in term reduces the value of $1 + \frac{[S]}{k_m}$ so the logarithmic value leading to reduce zoi . The value of zoi is significantly decreases by $-\log MIC$ value.

The MIC is (directly proportional) $\propto (f v_0 - 1)$ where $f v_0$ is the ratio of $\frac{v_0}{v_i}$ where v_0 is the velocity without inhibitor and v_i is the velocity with inhibitor, as enhancement of v_i leads to decrease the MIC and *vice-versa*, similarly v_0 progression will promote the value of MIC on higher side. The relationship ($MIC \propto \frac{v_0}{v_i} - 1$) On the condition's if $v_0 > v_i$ MIC will have a positive value, if $v_0 < v_i$, MIC becomes negative because the ratio $\frac{v_0}{v_i}$ is less than 1, so subtracting 1 yield negative result that's a hypothetical condition. If $v_0 = v_i$, then $MIC = 0$ it is similar to ideal situation as describe in equation (35). Contrary to that $f v_0$ is in denominator eq (8c), 21 and 28. It give an undefined value of MIC. The expression $e^{-\frac{x^2}{4Dt}}$. This term relates to the exponential function with negative input, as the value of MIC drops as ZOI 'x' increases. This adjustment is extremely sensitive because of the factor 4 in the denominator inside the exponent.

4. Conclusion

The mathematical model can be used to simulate the antimicrobial susceptibility test disk diffusion method in silico and to predict and correlate resistance, intermediate and susceptibility mathematically, we can calculate the binding energy, zoi and minimum inhibitory concentration

of antibiotic. We can also relate the various factors involved in invitro conditions, for drugs Hydrophilic in nature. This model also gives an insight of IC₅₀ and MIC as they both represents concentration of the antibiotic.

IC₅₀ is the concentration needed to inhibit process (e.g, enzyme activity or microbial growth) by 50% while MIC refers to the lowest concentration that inhibits the visible growth. For antimicrobial activity the value of binding energy ΔG should be greater than ΔG_{min} for the particular inhibitor to show antimicrobial activity. The Gibbs-Cheng equation a novel attempt that effectively models how the balance of inhibitor potency, enzyme-substrate affinity, and substrate concentration contribute to the overall free energy of the system. It's a hybrid equation connecting thermodynamic principles (ΔG) with enzyme kinetics (IC₅₀, K_m, [S]).

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