

How enzymes are adsorbed on soil solid phase and factors limiting its activity: A Review**

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A b s t r a c t. A majority of biochemical reactions are often catalysed by different types of enzymes. Adsorption of the enzyme is an imperative phenomenon, which protects it from physical or chemical degradation resulting in enzyme reserve in soil. This article summarizes some of the key results from previous studies and provides information about how enzymes are adsorbed on the surface of the soil solid phase and how different factors affect enzymatic activity in soil. Many studies have been done separately on the soil enzymatic activity and adsorption of enzymes on solid surfaces. However, only a few studies discuss enzyme adsorption on soil perspective; hence, we attempted to facilitate the process of enzyme adsorption specifically on soil surfaces. This review is remarkably unmatched, as we have thoroughly reviewed the relevant publications related to protein adsorption and enzymatic activity. Also, the article focuses on two important aspects, adsorption of enzymes and factors limiting the activity of adsorbed enzyme, together in one paper. The first part of this review comprehensively lays emphasis on different interactions between enzymes and the soil solid phase and the kinetics of enzyme adsorption. In the second part, we encircle various factors affecting the enzymatic activity of the adsorbed enzyme in soil.

K e y w o r d s: adsorption of protein, adsorbed enzyme, enzymatic activity, soil solid phase, enzymes in soil

INTRODUCTION

The study on soil enzymatic activities has emerged to be a major research area in soil microbiology. This has been fostered by the need to understand the soil biological and chemical processes, as well as the catalysts and factors that foster or undermine these processes to enhance the soil productivity (Burns, 1982). Soil can be perceived as a living ecosystem through which numerous biochemical processes take place with the help of both intracellular and extracellular enzymes (Skujinš and Burns, 1976). In line with this, Nannipieri *et al.* (2003) argue that soil is a dynamic and complex biological system, whose complete microbial community composition is yet to be fully established, and hence understanding the link between the microbial activities, the community structure, and the soil composition is crucial. As such, the soil microorganisms play a crucial role in mediating the biochemical processes to ensure nutrient cycling, which entails physicochemical, chemical, and biochemical reactions. Different researchers hold it that enzymes present in soil are often released from the dead cells of living organisms or excreted from the microorganisms within the soil (Skujinš and Burns, 1976). Particularly, the extracellular enzymes often emanate from the death of animal and plant cells; these cells are associated with rapid reproduction. Further, the extracellular enzymes have been found to be remarkably stable with respect to soil biochemical processes, an aspect that is attributed to their humus and clay association. However, irrespective of the stability, the enzymatic activity is highly dependent on the ambience of the soil pH and temperature (Kedi *et al.*, 2013).

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Table 1. Classification of different amino acids

Amino acid with hydrophobic side chain	Amino acid with electrically charged side chain		Amino acid with polar uncharged side chain	Special cases
	Positive charge	Negative charge		
Alanine	Arginine	Aspartic Acid	Serine	Cysteine
Valine	Histidine	Glutamic Acid	Threonine	Seleno Cysteine
Isoleucine	Lysine		Asparagine	Glycine
Leucine			Glutamine	Proline
Methionine				
Phenylalanine				
Tyrosine				
Tryptophan				

At any given point in time, the soil enzymatic activity encompasses a wide array of activities, each of which is associated with different constituents, such as humic colloids, clay, cell debris, and different microorganisms (Burns, 1982). A crucial aspect associated with the soil biochemical activities is the process of the enzyme adsorption. Enzyme adsorption is a protein immobilization technique that takes advantage of the enzyme-carrier physical interactions including the hydrogen bonds, the ionic interactions, and the van der Waal forces (Jesionowski *et al.*, 2014). This comprehensive review is aimed at gaining a better understanding of the enzyme adsorption on the soil solid surface and factors that limit this process. The adsorption process, the driving forces, the adsorption isotherm and reversibility of adsorption, the enzyme immobilization and its catalytic properties, and the factors controlling enzyme adsorption have comprehensively been reviewed.

GENERAL OVERVIEW

Enzymes are essential constituents of biological systems. All chemical reactions that sustain life are catalysed by enzymes (McCord, 2000). Twenty-one different amino acids are attached together in one or more polypeptide chain to form an enzyme molecule (Atkins and Gesteland, 2000). These amino acids are glycine, alanine, valine, leucine, isoleucine, proline, 4-hydroxy proline, serine, threonine, aspartic acid, glutamic acid, asparagine, glutamine, lysine, arginine, histidine, phenylalanine, tyrosine, tryptophan, cysteine, and methionine (Atkins and Gesteland, 2000; Minervini *et al.*, 2014). The above twenty-one amino acids are basically divided into four groups on the basis of polarity and charge (Campbell *et al.*, 2006) (Table 1):

- amino acid with a hydrophobic side chain,
- amino acid with electrically charged side chain,
- amino acid with a polar uncharged side chain,
- special cases.

Enzymes are generally globular proteins (Koshland, 1963). The amino acid sequence in the polypeptide chain determines the enzyme structure. Like all proteins, enzymes are linear chains of amino acids that fold to produce

a three-dimensional structure. The three-dimensional structure is a net result of all different intra-molecular interactions between segments within the enzyme molecule and its surroundings (Fitzsimons *et al.*, 2007; Mulvihill and Donovan, 1987). The side chain of an amino acid present in the enzyme molecule varies its polarity. Some are hydrophobic while others are hydrophilic. As a result of different interactions, there are four levels of enzyme structures: primary, secondary, tertiary, and quaternary.

ADSORPTION PROCESS

The adsorption of enzymes on the soil solid phase is a quasi-reversible phenomenon with great significance. Adsorption can also stabilize enzymes leading to their resistance toward denaturation, compared to a free enzyme. However, the mechanism of the enzyme interaction with the soil surface is still unclear (Norde, 2008). The microbial life in the soil is influenced by adsorption of extracellular enzymes into the soil (Nannipieri *et al.*, 2003). An abundant amount of silica (SiO₂) is present in soil and is responsible for different physicochemical interactions between enzymes and soil. The outer surface of amorphous silica is exposed to silanol groups (Si-OH) and siloxane bridges (-Si-O-Si-). At pH above 3, these silanol groups lose the hydrogen ion and develop a negative charge. As a result, the outer surface of soil becomes negatively charged (Zhuravlev, 2000). Lysine and arginine are free *i.e.* not involved in intra-molecular interactions, and are thus available for electrostatic interactions with the silica surface (Karlin *et al.*, 1994, Musafia *et al.*, 1995).

Forces involved in enzyme adsorption

Different forces are responsible for enzyme adsorption on the soil surface, but only electrostatic attraction is responsible for the first contact between the enzyme and the soil surface (Andrade and Hlady, 1986; Fenoglio *et al.*, 2011; Ikeda and Kuroda, 2011). Electrostatic attraction is due to the electrostatic force of attraction between

a charged amino acid and a negative charge present on silica. Arginine, histidine, and lysine have an electrically positive charged side chain above pH 4; hence, they are called basic amino acids. They are key amino acids responsible for electrostatic interactions with the electronegative silica surface. Haynes and Norde explained thermodynamically various interactions contributing to protein adsorption (Haynes and Norde, 1994). Different types of interactions take place between an enzyme and a soil solid surface such as Van der Waals forces, Lewis acid-base forces, and entropy-based effects such as hydrophobic interactions, conformational entropy, and restricted movements. In addition to the above forces, adsorption depends on forces within the helical structure of enzymes, which make them hard or soft.

Deformation of enzyme on soil surface

After being attached, the enzyme molecule undergoes structural deformations. Rigid and tightly coiled enzymes are not easily deformed on the soil surface and are less prone to adsorption. This structural deformation is due to the charged amino acid, which remains deep inside the helical structure of the enzymes (Secundo, 2013). Structural deformation is due to the conformation of an enzyme that belongs to free energy minimum in a solution state and is not necessary to have same free energy minimum conformation after being adsorbed on the soil surface. A greater contact surface area of the enzymes to the soil surface induces a gain in free energy. Hence, the enzymes tend to maximize their footprint through different conformational changes, as shown in different experimental works (Billsten *et al.*, 1995; Giacomelli and Norde, 2001; Santore and Wertz, 2005). The extent to which an enzyme is deformed strictly depends upon its structure, as explained by hard and soft protein (Galkin and Vekilov, 1999; Norde, 2003, 2008; Norde and Giacomelli, 2000). Instrumental techniques like ATR-IR (Karlsson *et al.*, 2005; Sharp *et al.*, 2002) and CD

spectroscopy provide direct information about the change in the secondary structure of the enzyme (Billsten *et al.*, 1995; Karlsson *et al.*, 2005; Vieira *et al.*, 2009).

Intermolecular forces within the helical structure of enzymes are mainly responsible for deformation of the enzyme on a solid surface, which happens just after the enzyme touches the soil surfaces. The reaction of the enzyme at interfaces is a net result of various types of interactions between different components of the system, *i.e.*, the enzyme molecules, the sorbent surface, and the solvent (water) molecules (Zoungrana *et al.*, 1997). An enzyme with weak structure can deform easily. As a result, more electrically charged side chains are exposed to oppositely charged silanol groups (Si-OH) present on the soil surface. Because of more interactions, the enzyme structure is deformed, and the enzyme is able to spread to maximum on the soil surfaces.

Percentage participation of each of the different types of the above-explained interactions depends on the nature of the system. For enzymes with strong intermolecular interactions, structural rearrangements do not contribute significantly to the adsorption process. Such enzymes are easily adsorbed on hydrophobic surfaces; however, they can adsorb on hydrophilic surfaces only if they are electrostatically attracted (Krisdhasima *et al.* 1992, Norde, 1986, 2003).

Enzymes with relatively weak intramolecular interactions are adsorbed on the hydrophilic surface even under extremely unfavourable conditions, *e.g.* an electrostatically repelling surface. In such a case, structural rearrangements are the main driving force for adsorption (Krisdhasima *et al.*, 1992; Malmsten, 1998; Norde, 2003).

Adsorption kinetics

Various steps of enzyme adsorption and desorption on the soil solid surface are shown in Fig. 1. They are: step 1. transport of the enzyme toward soil surface for the bulk,

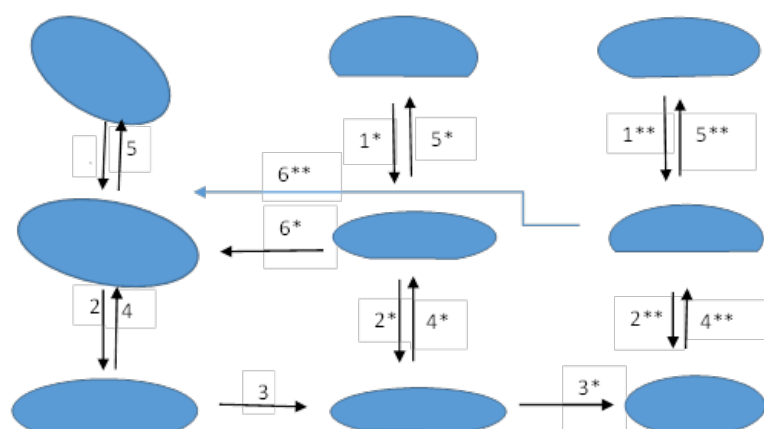


Fig. 1. Schematic presentation of the enzyme adsorption and desorption process. Explanation of step numbers in the text.

- step 2. deposition at the soil surface,
- step 3. unfolding of the enzyme,
- step 4. detachment of the enzyme from the soil surface,
- step 5. transport away from the soil surface,
- step 6. possible restructuring of the adsorbed enzyme.

An enzyme is transported near the soil surface by the convection process and its flow may be laminar or turbulent, but the last distance to the soil surface cannot be covered by the convection flow. The last step of attachment is always covered by diffusion. The enzyme and the adsorbing surface need to be dehydrated to overcome the hydrodynamic barrier just before the attachment of the enzyme to the soil surface (Kleijn and Norde, 1995). Before the attachment of the enzyme to the soil surface, the enzyme should be in proper spatial orientation towards the surface such that oppositely charged sites are in front of each other (Andrade and Hlady, 1986). Electrostatic attraction is the first as well as the rate-determining step. Without proper electrostatic attraction, there will be no initiation of adsorption irrespective of diffusion. The second step of the adsorption process depends on the degree of 'hardness-softness' of the enzyme or protein (Nannipieri *et al.*, 2003). Various steps of adsorption and desorption are shown in Fig. 1.

The process of enzyme adsorption on the soil surface is broadly divided into two individual steps:

1. transport of the enzyme from the bulk to the soil surface,
2. deposition of the enzyme on the soil surface.

Both these steps have different limiting factors and different rate constants.

Step 1: transport of the enzyme from the bulk to the soil surface

As discussed above, the basic mechanisms of enzyme transport toward the soil solid surface are diffusion and convection (laminar or turbulent flow). Most transport processes take place through steady state convective diffusion driven by a fixed concentration gradient. In the steady state condition of laminar flow, the enzyme flux toward the soil solid surface is determined by transport coefficient K_m (depends on hydrodynamic condition) and the difference in enzyme concentrations from the surface to the bulk.

The enzyme flux can be expressed as:

$$J = K_m (C_b - C_s), \quad (1)$$

where: C_b and C_s are the enzyme concentrations in the bulk solution and at the surface respectively, and K_m is the transport coefficient.

Step 2: Deposition of the enzyme on the soil surface

Deposition of the enzyme on soil solid surface is directly proportional to the concentration gradient. It may be considered as a first order process.

The rate of change in the adsorbed amount is given by:

$$\frac{d\Gamma}{dt} = K_n (C_s - C_{eq}), \quad (2)$$

where: Γ is the adsorbed mass per unit soil solid surface area and K_n is the deposition rate constant. C_{eq} is the concentration in the bulk solution corresponding to the equilibrium value for Γ , as given by the adsorption isotherm. Enzyme deposition is now determined by the rate constant K_n . Rate constant K_n decreases with an increase in C_s until the equilibrium is reached. Thus, when the enzyme concentration at the soil solid surface is equal to the equilibrium concentration, the adsorption rate becomes zero.

Relaxation of adsorbed layer

After being attached to the soil solid surface, enzyme molecules start deforming from their native structure, which is termed as structural relaxation or unfolding of the enzyme. The unfolding of the enzyme is due to changes in the microenvironment of the enzyme. Structural relaxation is represented by step 3 in Fig. 1. Relaxation is difficult in the case of strong interaction between the enzyme and the soil surface. Structural relaxation leads to better spreading of enzyme molecules over the soil solid surface. Now the relaxed enzyme molecules are strongly attached and, therefore, more difficult to detach, which may result in structural heterogeneity in the adsorbed layer. During the initial stage of the adsorption process, the enzyme molecule gets sufficient area available for spreading. However, this is not the case for later stages of the adsorption process as the surface is already covered by enzyme molecules.

If the rate of spreading of the enzyme is faster than the rate of attachment, the shape of adsorbed enzyme molecules becomes flattened. However, when the enzyme flux increases, the adsorbed enzyme molecule acquires more globular conformation and, therefore, the adsorption mass per unit surface area increases.

DRIVING FORCE

The necessary driving force for the adsorption process is provided by negative Gibbs free energy G at constant temperature and pressure and is given by the expression:

$$\Delta G = \Delta H - T\Delta S < 0, \quad (3)$$

where: H and S are the enthalpies and the entropy of the system and where Δ indicates the change invoked by the adsorption process. The more negative value of Δ , the higher the adsorption affinity.

The last and final step of the adsorption process is carried out by structural rearrangement of proteins, their surface polarity, reshuffling of distribution of charges at the interface, and dehydration of the surface (Haynes and Norde, 1994). During the structural rearrangement of an enzyme of the adsorbing soil surface, different intramolecular interactions become weak to maintain its secondary structure. There is a continuous increase in disorderliness or entropy of the enzyme, which favours the adsorption process.

Enzyme adsorption can occur easily on the polar hydrated surface because some non-polar part of the enzyme is present on its surface, which easily attracts the nonpolar part

of the soil surface, but the strongest driving force is shown by dehydrated nonpolar absorbing surfaces (Micaêlo and Soares, 2007).

An electrical charge double layer forms at the soil surface and its counter charge remains present in the diffusion layer. This model is well described by Stern (Kirkwood, 1934). Electrostatic attraction between electrical double layers is formed on the surface of soil colloids and the enzyme molecule contributes to the negative value of Gibbs free energy.

ADSORPTION ISOTHERM AND REVERSIBILITY OF ADSORPTION

Adsorption data are typically presented as an adsorption isotherm. The measurement of the adsorption concentration versus the aqueous enzyme concentration gives an adsorption isotherm. Enzymes with globular structure form compact monolayer structure of molecules. In the case of an incomplete monolayer, a smaller plateau is obtained. This happens at a low concentration of the enzyme and depends on the orientation of the enzyme on the soil surface. The same enzyme has different patterns of orientation on soil surfaces at different concentrations. At first, the enzyme is attached to the adsorbed surface without any conformational change and this process is very fast. Later, the thickening of the adsorbed layer of enzymes occurs, and this is a slow process. The structure of the adsorbed layer depends on the kind of enzyme and type of surface. For example, if the shape of the enzyme is considered rectangular, two types of configuration may be present, end-on and side-on. At a low concentration of the enzyme, the side-on configuration is mainly found and, at a high concentration, the end-on configuration may be found (Andrade and Hlady, 1986; Kleijn and Norde, 1995; Nakanishi *et al.*, 2001).

ENZYME IMMOBILIZATION AND ITS CATALYTIC PROPERTIES: K_m AND V_{max}

Soil enzyme is immobilized by either adsorption to clay mineral or it can bind chemically to organic matter present in the soil (Theng, 1979; Ladd, 1985). An immobilized enzyme varies in catalytic properties due to its heterogeneous nature (Katchalski *et al.*, 2006). Heterogeneous catalysis involves the following processes:

- a) adsorption of reactants on the surface of the catalyst,
- b) chemical reactions of the activated forms on the surface,
- c) desorption of the reaction products.

Adsorption of the substrate on the surface of the catalyst and desorption of reaction products are a diffusion phenomenon governed by laws of diffusion.

Several studies show that enzyme immobilization decreases its activity, for example, protease (Butler and Ladd, 1969), oxidoreductase (Pflug, 1980; Sarkar and Burns, 1984), phosphatase, and invertase (Malcolm and Vaughan, 1979), and alters the kinetic parameters. Immobilized enzymes have lower V_{max} and higher K_m , compared to a free counterpart, indicating that binding results in structural

changes in enzyme molecules (Kiss *et al.*, 1998; McLaren and Packer, 2006). The immobilized enzyme shows an increase in stability towards physical, chemical, and biological degradation (Sarkar and Burns, 1984). Factors that may be responsible for decreased activity of immobilized enzyme are:

- a) conformational changes in enzyme structure upon immobilization on a solid support,
- b) steric limitations of the substrate to penetrate and reach the enzyme active site,
- c) diffusion limitation of the substrate from the bulk of the solution to the immobilized enzyme and product from the enzyme active site to the bulk of the solution.

The above three factors are mainly responsible for the change in the value of kinetic parameters of immobilized enzymes. Immobilization of the enzyme results in a significant decrease in V_{max} values and an increase in K_m values of enzymes, compared to free enzyme. V_{max} values usually decrease after adsorption. Few studies have reported an increase in V_{max} after immobilization (Lei and Bi, 2007), but K_m values do not necessarily increase for all enzymes; they may increase or decrease depending upon the type of enzyme. Gianfreda proved experimentally that invertase and phosphatase decrease V_{max} and increase K_m values, compared to free enzymes after adsorption on pure and dirty clay. However, an increase in both V_{max} and K_m values was reported on all clay (Gianfreda, 2000; Gianfreda *et al.*, 1991; Rao and Rani *et al.*, 2000).

Enzyme immobilization is considered an important phenomenon because once immobilized, the enzyme is more resistant to inactivation and degradation to the extreme in situ soil conditions, and it has 90% catalytic activity of the native enzyme (Rani *et al.*, 2000). The immobilized enzyme, which is easily removed from the reaction solution, can later be reused.

Michaelis-Menten parameters can best explain enzyme inactivation upon adsorption to soil solid surface. The K_m value is the measure of the affinity of the enzyme for the substrate. Higher K_m indicates lower affinity and *vice versa*. The V_{max} value describes maximum velocity of conversion of substrate to the product when all the enzyme active sites are saturated with the substrate. The K_m and V_{max} can be useful parameters as they are independent of the enzyme concentration used (Schnell and Maini, 2003).

The Michaelis-Menten parameters can be simply applied to enzyme substrate reactions for the free enzyme *i.e.* homogenous reactions. Apparent kinetic values are obtained in case the enzyme is bound to the soil solid surface *i.e.* heterogeneous catalysis with diffusion limitation reactions. Two different types of enzyme inhibition occur upon adsorption of the enzyme to the soil solid phase (Fig. 2):

- a) competitive inhibition,
- b) noncompetitive inhibition (Fig. 2).

Competitive inhibition is reversible where the inhibitor competes with the substrate to bind to the same active site on the enzyme and vice versa. This may cause an increase in the K_m value without change in V_{max} .

This type of enzyme inhibition is similar to the temporary binding of organic matter to the enzyme active site.

In noncompetitive inhibition, the molecule binds to an enzyme surface other than the active site resulting in some conformational changes in the enzyme, such that the V_{max} value decreases but the K_m value remains the same (Fig. 2). This type of inhibition may be analogous to enzyme mineral or organic matter interaction.

Mixed inhibition occurs in those soil mineral and soil organic matter interactions that calculate Michaelis-Menten parameters. In most experiments, there is an increase in the K_m value, indicating decreased affinity of the enzyme toward the substrate. A decrease in the K_m value indicates an increase in enzyme-substrate affinity, which may occur due to the formation of an adsorption bond orientation, which causes enhanced active site accessibility to the substrate and increased enzymatic activity, mainly for montmorillonite.

FACTORS CONTROLLING ENZYME ADSORPTION

Enzyme adsorption is a very complex phenomenon and there are many internal and external parameters affecting enzyme adsorption on a solid surface. A few of them are: temperature, pH, soil moisture, and concentration of dissolved ions.

Temperature

Temperature is a crucial factor whose effect on the adsorption environment cannot be underestimated. While there are contradicting reports on the specific degree of the temperature effect on enzyme adsorption, almost all researchers agree that a change in temperature bears a significant effect on the enzyme adsorption, though there are variations in the recorded values. In an experiment using Sigmacell 50 and Sigmacell 20 and enzyme acquired from *T. viride*, Kim *et al.* (1998) unveiled that the Sigmacell adsorption capacity increases with an increase in temperature. Ooshima *et al.* (1983) also concluded that enzyme adsorption increases with an increase in temperature in an experiment entailing the use of Avicel at 5 and 50°C to adsorb cellulases. Bonomo *et al.* (2006) report that protein structures are affected by an increment in temperature, thereby exposing the hydrophobic sections of the protein to the adjacent environment. This, in turn, enhances the enzyme adsorption. Lee (2014) found only a small change in enzyme adsorption in an experiment using Solka floc SW-40 at 4 and 50°C to adsorb cellulases. Similarly, Barker *et al.* (1992) found out that the adsorption rate increased to maximum when the temperature of the enzyme mixture was increased from 55 to 60°C.

As observed, enzyme adsorption increases above room temperature. This increase in adsorption is due to structural rearrangements at a higher temperature. The kinetics

of enzyme adsorption is directly affected by even a small change in temperature and thereby altering the equilibrium state and all other kinetics parameters. With the change in temperature, the diffusivity of enzymes towards the soil surface is also affected.

The adsorption rate is likely to be increased due to the rapid diffusivity of the enzyme towards the soil surface. Enthalpy gain is the main driving force responsible for enzyme adsorption. It comes from structural changes taking place in the enzyme molecule and release of salt and water molecule from the surface (Andrade and Hlady, 1986; Malmsten, 1998; Norde, 1986; Ramsden, 1995). Hence, enzyme adsorption increases at higher temperatures (Koutsoukos *et al.*, 1983; Norde, 1986).

pH

The soil pH is a crucial factor responsible for controlling a wide array of microbial processes. A change in the soil pH may result in altered hydrophobicity of the soil solid surface, which in turn may significantly affect the enzymatic activities. It is crucial to note that different enzymes operate best at different pH levels. A study by Bezerra and Dias (2005) entailing the adsorption of cellulases QM 9414 on Sweco 270, Solka Floc SW 20, and Avicel PH 102 at 30°C with a pH of 3.8 to 5.5 unveiled that as the pH increased from 4.8 to 5.5, the enzyme adsorption decreased. This is seemingly contradicting to the findings by Tomme *et al.* (1990) who experimented on enzyme adsorption using *Trichoderma reesei* cellulases and concluded that 4.8 to 8.5 is the optimum pH for enzymatic hydrolysis. On the same note, an experiment by Kyriacou *et al.* (1988) using Solka floc BW-40 and fractionated *T. reesei* cellulases unveiled that even with an increase in the pH from 3 to 7, the maximum adsorption at 5 and 50°C was the same.

Soil pH is a very important parameter for the progression of soil biochemical reactions. Soil pH has a direct effect on interfacial attraction. The electrostatic behaviour of the enzyme molecule is determined by the pH of the surroundings. When pH is equal to the isoelectric point of the enzyme, the numbers of negative and positive charges are balanced, resulting in the net neutral molecule. Hence, pH determines the electrostatic state of enzymes. Different enzymes have a different isoelectric point. At isoelectric pH, the net charge on enzymes remains neutral. At low pH, the net electric charge on the enzyme is positive whereas, at high pH, the net charge is negative. Minimum repulsive forces are expected at isoelectric point allowing a thicker layer of the adsorbed enzyme on the soil surface. However, maximum adsorption is observed at the isoelectric point. Hence, it can be asserted that the isoelectric point denotes the maximum adsorption (Bremer *et al.*, 2004; Demanèche *et al.*, 2009; Sethuraman and Belfort, 2005).

Soil moisture

Soil water stress occurs seasonally almost in all soils. Drying of soil can result in a decrease or complete cessation of enzyme activity. Many studies have been done on the impact of soil water stress on microbial communities. In most of the studies, enzyme activity was analysed (Zelles *et al.*, 1991). Enzyme adsorption at the soil solid surface can result in inactivation caused by conformational changes due to an increased concentration of the enzyme at soil-water interface (Durán and Esposito, 2000). Prokaryotes living in the thin water layer surrounding the soil particles are very sensitive to water depletion. Changes in soil as a result of drying are more physical than chemical. Enzyme activity decreases with a decrease in the size of soil organic matter and increases with a decrease in soil mineral matter (Killham, 1994). Chemical changes are usually very small (Wu and Brookes, 2005) and occur within 1-2 months of soil drying. Many cycles of drying and rewetting alter soil water potential creating osmotic stress (Halverson *et al.*, 2000), and it leads to microbial death unless they remain dormant until more favourable conditions occur (Schimel *et al.*, 2007).

Concentration of dissolved ions

The concentration of dissolved ions has a significant effect on enzyme adsorption. Increasing salt concentrations result in decreased enzyme adsorption at the interface until a complete suppression at high concentrations is reached. This is because of reduced attractive electrostatic interaction between the positively charged enzymes and negatively charged surfaces by charge screening (Jones and O'Melia, 2000). Further, high ionic concentration causes significant interference with the weak ionic bonds in the enzymes. It is crucial to note that enzymes comprise a non-protein called co-factor and a protein that is made of chains of various amino acids. As such, the enzymes are characterized by inter and intra-molecular bonds which keep the molecules in position (Liu and Zhu, 2010). The interference caused by the high ionic concentration leads to dysfunctional bonds between the amino-acids group. This, in turn, leads to denaturation, an aspect that entails the loss of the shape by an enzyme. Essentially, if an enzyme structural shape is compromised, it becomes challenging for it to react and accept substrate, and hence the adsorption rate is significantly affected (Liu and Zhu, 2010).

Earlier it has been known that an increased salt concentration supports the precipitation of colloidal substances. This is due to the absorption of water by salt ions. This phenomenon is known as a salting out effect. Hofmeister found that different salt ions have different ability to precipitate proteins from a solution, which leads to the concept of the Hofmeister series (Kunz *et al.*, 2004). Kosmotropes are ions that promote protein precipitation leading to a cha-

otropic effect. Chaotropic and kosmotropic effects are due to stabilization and destabilization of native conformations of proteins, which influence enzyme adsorption.

KEY FACTORS AFFECTING THE ENZYMATIC ACTIVITY OF ADSORBED ENZYMES IN SOIL

Substrate diffusion

Diffusion is an important process in the kinetics of rapid reactions in a solution. Diffusion may be the rate-limiting step in very rapid reactions and such reactions are termed as diffusion-limited reactions. Diffusion is also an important step for heterogeneous reactions occurring at the solid surface. Limitation of substrate diffusion towards the enzyme adsorbed on the soil surface could be one of the key factors for the decrease in enzyme activity (Arrio-Dupont and Béchet, 1989; Bille *et al.*, 1989; Douzou and Petsko, 1984; Konecny and Voser, 1977). Enzyme substrate reactions are zero order kinetic. Enzyme activity is maximum on the soil surface; hence, the rate of the reaction is maximum near the surface. The concentration of the substrate remains lowest adjacent to the surface relative to the bulk of the solution. The concentration gradient develops from the bulk of the solution to the soil surface (Bille *et al.*, 1989; Ku and Lentricchia, 1989). There can be three different situations if the diffusion is: slow, equal and fast relative to substrate consumption at the soil surface.

If diffusion is slow, it is the rate-limiting step and the reaction is diffusion limitation. If diffusion is equal to substrate consumption at the soil surface, then the steady state will eventually be established such that the rate of diffusion of the substrate in the unstirred layer (Nernst diffusion layer) equals the consumption rate of the substrate. In the case of faster diffusion, the reaction will be enzyme limited. In most of the soil conditions, the enzymatic activity is usually low, compared to the large surface area of soil (Datta *et al.*, 2014).

It should be noted that for a reaction to be diffusion-limited, it is necessary for the step following the initial combination of the enzyme and substrate to be faster than diffusion apart from the reactants; therefore, diffusion-limited reactions are indirect evidence for a conformational change following the combination of the enzyme and substrate.

Orientation effects

Enzymes are small asymmetric complex molecules of different shapes such as elliptical, rod-like, heart-like, or Y-shaped, *etc.* Enzymes can rotate freely in the solution whereas on the surface different enzymes adapt certain orientation. The orientation decides which part of the enzyme will face towards the soil surface and which part will expose towards the bulk solution. Orientation is an imperative factor for enzymatic activity. Any improper orientation can decrease or completely inhibit enzymatic activity. Spatial enzyme immobilization on a solid surface increases its activity and stability (Brady and Jordaan, 2009; Cao,

2005; Hanefeld *et al.*, 2009; Tischer and Wedekind, 1999). Orientation on the solid surface can be characterized by 'side-on' or 'end-on'. These orientations are best explained by Lu *et al.* (1998) and Su *et al.* (1998).

The selective orientation of the enzyme on the electronegative soil surface is expected if a higher density of a positively charged amino acid is present at this face. In this regard, the following cases are expected:

- if the more positive face is not varying with a change in pH during the reaction process, and the active site is present on this face, then complete inhibition of the enzyme could be expected. No inhibition of activity is observed if the active site is on the opposite side;
- if a more positive charge faces changes with increased pH, an alkaline shift is observed. If the active site is present on this face, then an increase in optimal enzyme activity is observed due to more exposure of the active site towards the substrate.

Similarly, the acid shift is observed if the polarity of the enzyme changes due to a decrease in pH. Apart from explained cases, there are many situations where the active site is neither completely masked nor completely exposed. Enzymes of hard proteins are less prone to pH-dependent modification of conformation and enzymes with higher dipole moment are more likely to show preferential orientation.

Temperature

Temperature is also one of the factors that strongly affect and control soil enzyme activity. It can influence enzyme kinetics, stability, substrate affinity, and enzyme production. Thermal stability of an adsorbed enzyme is often increased in comparison to that of a free enzyme. Enzymes retain their functionality and selectivity after being adsorbed on the soil surface. Adsorption increases thermal stability two to three times when compared to the same enzyme in a solution (Nannipieri *et al.* 1982).

Most of the studies conducted for determining the effect of temperature on enzymatic activity have been done in lab conditions. Very few studies are done in field conditions by soil warming treatment. Studies done in field conditions show very little to no change in enzyme activity (Allison and Treseder, 2008). Temperature and climate change affect the activity of the adsorbed enzyme in different ways, which it is difficult to predict; most commonly, it is observed that the activity increases up to some optimum level (Wallenstein and Weintraub, 2008). Soil microbes may decrease their secretions after a certain level of warming (Allison, 2005). The rate of enzyme denaturation also increases with warming (Wallenstein *et al.*, 2012).

Enzyme-clay mineral interaction

Like proteins, enzymes show a strong affinity towards different mineral surfaces. Clay plays an important role in the adsorption of the enzyme surface catalysed process involving enzymatic protein and enzyme immobilization in soil (Boyd and Mortland, 1990).

Extracellular enzymes are physically and chemically adsorbed on clay mineral surfaces. Some studies put forward that electrostatic interaction dominates protein mineral interaction (Ding and Henrichs, 2002; Servagent-Noinville *et al.*, 2000). The mechanism of enzyme-clay mineral interaction is a complex phenomenon because enzymes are made up of amino acids with varied properties, *e.g.* they can be hydrophilic, hydrophobic, positively, negatively and neutrally charged. The shape of the amino acid may change in response to various environmental conditions.

Adsorption of an amino acid monomer and dimer shows significant relations between the net charge present on it and the charge density present on the mineral surface. No such relation is seen in the case of protein or enzyme adsorption on the clay mineral surface. Therefore, electrostatic force cannot be considered as a dominating factor in controlling enzyme or protein adsorption although the surface of many minerals is negatively charged and proteins contain some amount of negative or positive charge, due to their amphiphilic nature (Zimmerman *et al.*, 2004).

Hydrophobic interactions and a decrease in the free energy of the enzyme due to conformational changes are the most predominant mechanism of protein adsorption on the clay mineral surface. This mechanism can also explain the adsorption of proteins on hydrophobic surfaces under unfavourable electrostatic conditions.

The amount of an enzyme adsorbed on the clay and its property to form a complex with the clay mineral rigorously depend on many factors such as the nature and properties of the enzyme and clay (Naidja, 1995; Theng, 1979). Gianfreda proved experimentally that there is no significant relation between the surface area of clay mineral and the amount of enzyme adsorbed (Gianfreda *et al.*, 1991; Greco *et al.*, 1982).

Change in the kinetic properties of the enzyme is observed upon adsorption to the mineral surface. The activity of the enzyme is reduced on adsorption, and the extent of reduction depends on factors such as the type of enzyme, clay mineral, and pH (Gianfreda *et al.*, 1991, 1992; Violante *et al.*, 1999). The list of changes in enzymatic activity that has been observed upon adsorption to various adsorbents is shown in Table 2.

Different responses of adsorbed enzyme on clay mineral surface were attributed to different:

- arrangements held by the enzymes on clays, and
- kinds of electrostatic interactions occurring between enzyme molecules and binding sites on clay.

Table 2. Changes in enzyme activity following clay mineral interaction

Adsorbent	Enzyme	Activity change	Reference
Ca-montmorillonite	Peroxidase	0 to -69	Lozzi <i>et al.</i> , 2001
Na-montmorillonite	Peroxidase	-88 to -99	Lozzi <i>et al.</i> , 2001
Ca-montmorillonite	Catalase	-81 to -99	Calamai <i>et al.</i> , 2000
Kaolinite	Tyrosinase	0 to -75	Claus and Filip, 1988
Kaolinite	Acid phosphatase	-64	Gianfreda and Bollag, 1994
Montmorillonite	Acid phosphatase	-68	Gianfreda and Bollag, 1994
Montmorillonite complex	Peroxidase	0	Gianfreda and Bollag, 1994
Al(OH) _x	Invertase	-94 to -99	Gianfreda <i>et al.</i> , 1991
Al(OH) _x -montmorillonite	Invertase	-89 to -95	Gianfreda <i>et al.</i> , 1991
Na-montmorillonite	Invertase	-88 to -96	Gianfreda <i>et al.</i> , 1991
Al(OH) _x -montmorillonite	Urease	-49 to -67	Gianfreda <i>et al.</i> , 1992
Na-montmorillonite	Urease	-41	Gianfreda <i>et al.</i> , 1992
Goethite	Acid phosphatase	-32	Huang <i>et al.</i> , 2005
Kaolinite	Acid phosphatase	-43	Huang <i>et al.</i> , 2005
Ca-montmorillonite	Arylsulphatase	-52	Hughes and Simpson, 1978
Kaolinite	Arylsulphatase	-18	Hughes and Simpson, 1978
Goethite	Glucosidase	0 to 40	Quiquampoix, 1987
Kaolinite	Glucosidase	-13 to -100	Quiquampoix, 1987
Kaolinite	Glucosidase	0 to -88	Quiquampoix, 1987
Na-montmorillonite	Glucosidase	-35 to -100	Quiquampoix, 1987
Al hydroxide	Acid phosphatase	-55	Rao and Gianfreda, 2000
Al hydroxide	Phosphatase	-55	Rao <i>et al.</i> , 2000
Al(OH) _x -montmorillonite	Phosphatase	-42	Rao <i>et al.</i> , 2000
Na-montmorillonite	Phosphatase	-80	Rao <i>et al.</i> , 2000
Allophane	Glucose oxidase	-52	Ross and McNeilly, 1972
Ca-montmorillonite	Glucose oxidase	-77	Ross and McNeilly, 1972
Illite	Glucose oxidase	-21	Ross and McNeilly, 1972
Kaolinite	Glucose oxidase	-17	Ross and McNeilly, 1972
Elledge clay	Alkaline phosphatase	-75	Tietjen and Wetzel, 2003
Elledge clay	Glucosidase	55	Tietjen and Wetzel, 2003
Elledge clay	Protease	-82 to 82	Tietjen and Wetzel, 2003
Montmorillonite	Glucosidase	50	Tietjen and Wetzel, 2003
Montmorillonite	Protease	-100	Tietjen and Wetzel, 2003

Adsorption of the enzyme on the solid soil surface may be modified by the presence of metal oxides and hydroxides such iron aluminium and manganese (Rao *et al.*, 2000). The presence of aluminium hydroxide on montmorillonite surfaces resulted in lower activity for invertase, tyrosinase, and urease, whereas the same metal oxide increases the activity of phosphatase (Gianfreda *et al.*, 1991, 1992; Naidja and Huang, 1996; Rao *et al.*, 1996).

Enzyme organic matter interactions

Apart from clay minerals, enzymes can also be adsorbed on soil humus (Ladd and Butler, 1975). The enzyme soil organic matter interaction plays an important and vital role

in controlling enzymatic activity, especially in soil comprising high organic matter content (Pflug, 1982). In soils with higher sand and organic matter, enzyme organic matter interactions are of greater significance in regulating soil enzyme activity and stability than the enzyme mineral interaction (Table 3).

Many mechanisms have been proposed for the formation of the enzyme-humus complex such as:

- covalent attachment of the enzyme to soil humic substances,
- physical entrapment in humus,
- hydrogen bonding, and
- ionic bonding (Burns, 1982).

Table 3. Changes in enzyme activity following organic matter interaction

Adsorbent	Enzyme	Percentage change in activity	Reference
2,4,6-Trimethylphenol	Laccase	0 to -30	Cañas <i>et al.</i> , 2007
Acetosyringone	Laccase	-10 to -50	Canas <i>et al.</i> , 2007
Acetovanillone	Laccase	-70 or +50	Canas <i>et al.</i> , 2007
Benzoquinone	Pronase	-32 to -96	Rowell <i>et al.</i> , 1973
Ferulic acid	Laccase	-30 or +80	Canas <i>et al.</i> , 2007
	Acid phosphatase	-20 to -65	Allison, 2006
	Catalase	100	Serban and Nissenbaum, 1986
	Glucosaminidase	-75 to -90	Allison, 2006
	Glucosidase	-70 to -100	Allison, 2006
	Malate dehydrogenase	-16 to -57	Pflug and Ziechmann, 1981
	Papain	+88 to +160	Butler and Ladd, 1969
	Papain	+49 to +226	Freeman <i>et al.</i> , 2004
Humic acid	Peroxidase	+18 to +125	Serban and Nissenbaum, 1986
	Phosphatase	+10 to +100	Boavida and Wetzel, 1998
	Polyphenol oxidase	-35 to -70	Allison, 2006
	Pronase	-40 to -73	Butler and Ladd, 1969
	Pronase	-25 to -35	Ladd and Butler, 1969
	Tyrosinase	-5 to +19	Ruggiero and Radogna, 1988
	Urease	-10 to -35	Allison, 2006
p-coumaric acid	Laccase	-85 or +250	Cañas <i>et al.</i> , 2007
Phenolic compound	Chitinase	-22	Freeman <i>et al.</i> , 2004
Syringaldehyde	Laccase	0 to -50	Canas <i>et al.</i> , 2007
Tannic acid	Acid phosphatase	-49 to -93	Rao <i>et al.</i> , 2000
Vanillin	Laccase	-50 to -60	Canas <i>et al.</i> , 2007

Sometimes non-polar interaction is also found between the enzyme and natural organic matter.

Soil organic matter cannot be thoroughly extracted out except humic substances because enzymes remain covalently bound to soil and, therefore, they become inseparable from the soil. Due to this, it is impossible to separate the enzyme humic acid complex by any purification methods (Rao *et al.*, 1996, 2000).

Enzymes react chemically with lignin, phenols, tannins, humic acid, and quinones to form complex organic matter components of soil. In this regard, strong pH-dependent bonds such as covalent, electrovalent, and hydrogen are known to occur. There are several studies on extraction and purification of the enzyme humic acid complex, but none of them has shown any single type of association. In fact, a different combination of mechanisms has been proposed by different groups. Such combinations are covalent bonding, hydrogen bonding ion exchange, and physical entrapment methods (Ruggiero and Radogna, 1988).

Many groups have shown the impact of the electrostatic force on the enzyme-organic matter complex. Inorganic cation can be used to inactivate the complexed enzyme. Use of sodium pyrophosphate in extracting soil organic matter bound extracellular enzyme was put forward by Ceccanti *et al.* (1978) and Nannipieri *et al.* (1982). They also suggested that large fractions of enzymes are covalently bound to the humic surface than electrostatic interaction (Mayaudon and Sarkar, 1975; Pflug, 1982). The correlation between enzyme activity in soil and extracted organic carbon or nitrogen suggests binding of enzymes to the humic material (Nannipieri *et al.*, 1980).

The porous structure of humic acid facilitates inward diffusion of the small substrate molecule and outward diffusion of the product molecule, but at the same time, the movement of larger molecules is restricted. Hydrophobic interactions are mainly responsible for enzyme humic acid interaction (Tan *et al.*, 2009; Zang *et al.*, 2000, 2001). It is observed that enzyme activity of the organic matter bound enzyme is highly specific for the particular enzyme. After interaction with organic matter, some enzyme activities increase while others decrease. Deactivation occurs due to the orientation of the enzyme active site by intermolecular forces. Soil porosity is also an important factor affecting enzymatic activity. Soil sieved through fine size mesh shows greater enzymatic activity than sieved through bigger mesh size sieve. This is due to exposure of a greater enzyme-adsorbed soil surface area to the substrate (Datta *et al.*, 2014).

RECENT TOOLS AND TECHNIQUE FOR STRUCTURAL DETERMINATION OF IMMOBILIZED ENZYME AND ITS SUPPORT

The three-dimensional orientation of an immobilized enzyme is crucial for its activity and stability. Immobilization is the net result of various conformational changes that occur within and outside the enzyme surface. Immobilization could cause altered enzyme activity (Schröder *et al.*, 2006). The formation of individual biomolecules and their orientation on surfaces can easily be monitored by different microscopic and spectroscopic techniques.

Fourier transformed infrared spectroscopy (FTIR) is a widely used technique to analyse immobilized enzyme structure. FTIR can easily determine changes occurring in

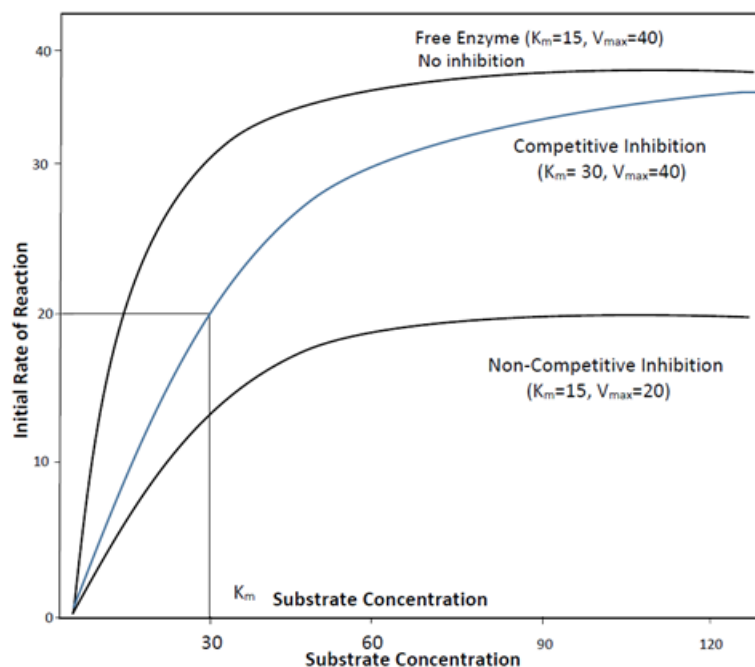


Fig. 2. Michaelis-Menton plot showing predicted kinetics change with 3 different types of enzyme inhibition.

the secondary and tertiary structure of the enzyme upon immobilization, the formation of the hydrogen bond, and other enzyme surface interactions (Barth, 2007; Carlsson *et al.*, 2014; Yang *et al.*, 2015). The Fourier transformed technique has been used to observe the structure of many enzymes such as lipase, protease, and glucose oxidase (Gole *et al.*, 2001; Portaccio *et al.*, 2014).

Circular dichroism (CD) spectroscopy serves as a powerful method in structural determination and enzyme immobilization studies. CD elucidates the structure information of enzyme solid surface interaction (Carlsson *et al.*, 2014; Thyparambil *et al.*, 2015). CD spectroscopy is an alternative to FTIR spectroscopy in the characterization of an immobilized enzyme. Many immobilized enzymes obtained with different methods have been analysed using CD. Modification in the enzyme secondary structure after immobilization has also been observed by CD spectroscopy (Barth, 2007; Thyparambil *et al.*, 2015; Verma *et al.*, 2013).

Atomic-Force Microscopy (AFM) is a unique microscopic technique that can visualize enzyme molecules at a single molecular level with high accuracy (da Silva *et al.*, 2015). It can visualize the enzyme surface, being attached to support (Gole *et al.*, 2001; Hanefeld *et al.*, 2009). Surface-induced changes due to enzyme surface interaction and changes due to enzyme interaction are easily visualized by AFM (da Silva *et al.*, 2015). Hence, the technique is widely used in many surface-based protein interactions (da Silva *et al.*, 2015; Gole *et al.*, 2001).

Electron microscopy visualizes an immobilized enzyme at nanometer level resolution with limited information about the internal structure. Scanning electron microscope (SEM) and field emission scanning electron microscope (FESEM) are highly applied techniques for morphological characterization of the enzyme and surface (Kröger *et al.*, 1999; Reátegui and Aksan, 2009). FESEM helps to visualize the very small change in enzyme morphology upon immobilization (Sang and Coppens, 2011). Transmission electron microscopy (TEM) also provides good structural morphology, but it provides only a two-dimensional surface structure (Lei *et al.*, 2006).

Raman spectroscopy is also potentially a strong technique to analyse enzyme structure in dried or frozen solution samples (Thyparambil *et al.*, 2015; Xiao *et al.*, 2012). There is increasing use of Raman spectroscopy to characterize immobilized enzyme. (Ash and Vincent, 2012; Xiao *et al.*, 2012).

Latest technique such as Foster resonance energy transfer (FRET) measurements (Mohamad *et al.*, 2015), solid-state nuclear magnetic resonance spectroscopy (NMR) (Noureddini and Gao, 2007), Trp/Tyr fluorescence measurements (da Silva *et al.*, 2015), and TOF-secondary ion mass spectrometry (Thyparambil *et al.*, 2015) are gathering strength and more advancement are expected in near future.

RECAPITULATION

Enzyme adsorption on the soil solid phase is a complex phenomenon; different forces acting together at the same time make it more complex. Enzyme organo-mineral relationship directly affects enzymatic activity in any soil. Although continuous research has been done on enzyme interactions on soil solid phase over the past decades, which helps us in advancing our understanding on enzyme adsorption, it is still unclear how an enzyme behave when it comes in contact with the soil surface. Further advancement in technology is necessary to understand in depth the enzyme organo-mineral interaction. More advanced spectroscopic techniques can help in determining changes occur in the tertiary structure of enzyme after adsorption.

Many studies on adsorption of proteins on solid surfaces focus on enzymatic activity in soil, but none of them emphasize on adsorption with soil perspective. In this regard, this review chiefly endeavours to simplify the process of enzyme adsorption on soil surfaces, clarifying further the effect of enzyme adsorption on enzymatic activity. Needless to say, enzyme adsorption of on soil surfaces must not be omitted while discussing the enzymatic activity. Furthermore, this article is obtrusive as it has exhaustively reviewed all important publications on protein adsorption on the soil surface and enzymatic activity done separately before compiling these two relevant aspects together in one paper.

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