

Economic Importance of *Bacillus Thurnigiens* is bacterium for Bio Prospecting: A Review

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Abstract:

Background: *Bacillus thurnigiensis* is the most effective and playing important roles as dominant bacteria in biotechnological applications as well as industrial processes and products. Exploration of this bacterium holds great promise because of the role of microbes in nutrient cycling, environmental detoxification and novel metabolic abilities in pharmaceuticals and industrial processes and act as a major resource for agricultural, industrial, and medicinal applications

Methods: The researcher searched the PubMed database and other different Journals Online archives from September 2018 to February 2019 using the search term 'bioprospecting potential of the bacterium' to find articles published recently onwards.

Results: We included 36 published articles in our analysis. Most articles were on occurrence, diversity, insecticidal activity, metabolism, metabolic role, and industrial uses of bacterial and also economic uses of the bacterium. Members of *Bacillus* and related genera are used for the synthesis of a very wide range of important medical, agricultural, pharmaceutical and other industrial products.

Conclusion: With recent advances in biotechnology, the economic contributions that these organisms can make in biotechnological applications and industrial processes can be exploited further for large scale benefit of mankind.

Keywords: *Bacillus thurnigiensis*, Bioprospecting, Wolaita.

1. INTRODUCTION

Microbial diversity is a major resource for biotechnological products and processes. Exploration of microbial diversity holds great promise because of the role of microbes in nutrient cycling, environmental detoxification and novel metabolic abilities in pharmaceuticals and industrial processes and act as a major resource for agricultural, industrial, and medicinal applications (Handelsman *et al.*, 1998). Bacteria are the most dominant group of this diversity which exists in diverse ecological niches, including extreme environments present in both lithosphere and hydrosphere, where their metabolic abilities play a critical role in geochemical nutrient cycling and producing a wide range of products of industrial significance (Daniel, 2005).

The gram-positive bacteria form an important part of the microbiota in many soils. In particular, the low G+C content gram-positive bacteria, which are divided into three classes such as clostridia, mollicutes, and bacilli, play major roles in the mineralization of plant-derived materials, humus, pesticides and hydrocarbons in soil (Prescott *et al.*, 2002). Members of *Bacillus* and related genera are used for the synthesis of a very wide range of important medical, agricultural, pharmaceutical and other industrial products. These include a variety of antibiotics, enzymes, amino acids and sugars (Probanzaet *et al.*, 2002). They are major elements of the agronomic environment of plants, inhabiting the soil. Several reports have described the biodiversity, antibiotic production and plant growth-promoting effects of *Bacillus* and related genera.

The purpose of this review paper is to introduce the potential of *Bacillusthuringiensi*s isolate indigenous from Ethiopian ecology as a natural product in a soil of economic, environmental, industrial, agricultural etc. importance and to summarize the current status and developmental trends of biological control based on the published reports.

1.1. Taxonomy of *Bacillus Thurnigiensis*

Bacillus thuringiensis was first described by Berliner when he isolated a *Bacillus* species from the Mediterranean flour moth, *Anagastakuehniella*, and named it after the province Thuringia in Germany where the infected moth was found. Although this was the first description under the name *B. thuringiensis*, it was not the first isolation. In 1901, a Japanese biologist, IshiwataShigetane, discovered a previously un described bacterium as the causative agent of a disease afflicting silkworms. *Bt* was originally considered a risk for silkworm rearing but it has become the heart of microbial insect control. The earliest commercial production began in France in 1938, under the name Sporeine (Lambert, B. and M. Peferoen. 1992). A resurgence of interest in *Bt* has been attributed to Edward Steinhaus, who obtained a culture in 1942 and attracted attention to the potential of *Bt* through his subsequent studies. In 1956, T. Angus demonstrated that the crystalline protein inclusions formed in the course of sporulation were responsible for the insecticidal action of *Bt*. By the early 1980's, Gonzalez *et al.*, 1982 revealed that the genes coding for crystal proteins were localized on transmissible plasmids, using a plasmid curing technique, and Schnepf and Whiteley (Schnepf, H. E. and H. R. Whiteley. 1981) first cloned and characterized the genes coding for crystal proteins that had toxicity to larvae of the tobacco hornworm, from plasmid DNA of *Bt* subsp. *kurstaki* HD-1. This first cloning was followed quickly by the cloning of many other cry genes and eventually led to the development of *Bt* transgenic plants. In the 1980s, several scientists successively demonstrated that plants can be genetically engineered, and finally, *Bt* cotton reached the market in 1996 (Tabashnik *et al.*, 2005). It is classified under Eubacteria (kingdom); Bacteria (domain); Firmicutes (phylum); Bacilli (class); Bacillales (order); Bacillaceae (family); *Bacillus* (genus); *Bacillus cereus* group; *Bacillus thurnigiensis* (species). *Bacillus thurnigiensis* is a gram positive, rod-shaped, spore forming bacterium which often has insecticidal properties. *Bt* belongs to the *Bacillus cereus* complex which also includes *B. cereus*, *B. anthracis* and *B. mycoides*. The taxonomic relationships between members of the *B. cereus* group are not clear (Drobniewski 1994) and the cause of some concern as the differences between *B. cereus* and *B. thurnigiensis* are small and may be mainly plasmid based. DNA sequencing studies of conserved gene regions have suggested they may be strains of a single species. During sporulation in *B. thurnigiensis*, some strains produce one or more inclusions or parasporal bodies within a sporangium. The parasporal body is often toxic to specific insect groups and many different insecticidal crystal proteins (δ -endotoxins) can be found in different *B. thurnigiensis* subspecies and strains.

Many studies indicate and consider *B. thuringiensis* and *B. cereus* to be one species. However, their phenotypes greatly differ in that *Bt* produces crystal proteins despite the fact that crystal protein synthesis is controlled by plasmid genes which can be susceptible to loss and transmission to related bacteria. One response is that *Bt* strains produce enterotoxins (toxins released by micro-organisms in the lower intestine) that are involved in *B. cereus* pathogenesis and therefore signifies a fine-line between the two species (Cherif *et al.*, 2007).

1.2. Distribution of *Bacillus Thurnigiensis* in Ethiopian Soils

The distribution of *Bacillus thurnigiensis* all over the agro-ecological zones in all soils of Ethiopia (Zelege Wolde Tenssay and Mogessie Ashenafi, 2008). Ethiopia's diverse climatic and agro-ecological conditions may also favor rich diversity of microbial flora. But, the hitherto studies on larvicidal activity of indigenous *Bt* against disease vectors are limited (Aklilu Seyoum and Dawit Abate, 1997). The efficacy of standard *B. thuringiensis* subspecies *kurstaki* on Ethiopian crop pests was reviewed by Tsedeke Abate (1997). However, a thorough knowledge of the ecological distribution and diversity of *Bt* would have a substantial contribution to evaluate their exact role in the environment and their effects on non-target organisms. It also helps to choose appropriate soil types and insect habitats to isolate novel *Bt* strains. The frequency of *Bt* was highest in tepid to cool semi-arid lakes and rift valley soils followed by cold to very cold sub-humid afroalpine and hot to warm per humid low lands. Subhumid afroalpine is characterized by limited plant species due to its cold temperature whereas the other two zones are characterized by moderate to rich fertility, which may be associated with several factors such as soil nutrients and physical and biological components important in spore germination and recycling of *B. thurnigiensis*. *Bacillus thurnigiensis* was least frequent in soil samples collected from hot to warm sub-moist mid highlands and hot to warm moist lowlands. Several workers (Hossain *et al.*, 1997; Nicholson, 2002) have suggested that variation in abundance and distribution of *B. thurnigiensis* in the soils of different sources might be due to several

ecological factors that affect the viability of *B. thurnigiensis* spores and recycling of *B. thurnigiensis* either by growth at the expense of nutrient present in the environment, or, to some extent, by association with diversity of insect types which may be distributed in those agro-ecological zones. Soil samples from the natural vegetation yielded *B. thurnigiensis* in a comparable rate to soils of cultivated fields. Soil samples collected from vegetable fields had lowest *B. thurnigiensis* yield. It is generally known that the number and types of microorganisms in soil vary depending on the type of plant cover. Naturally, insect pest distribution also differs with vegetation cover (Janzen, 1973; Bianchi et al., 2006), and if *B. thurnigiensis* distribution has any correlation to crop pest distributions, it is apparent that its distribution and abundance will differ with vegetation types of the soil.

1.3. Occurrence and Role in the Environment

An understanding of the ecology of *Bacillus thurnigiensis* in the environment is essential in assessment of its environmental risk. While originally recovered mainly from insects, improved isolation and identification techniques have indicated that Bt may be ubiquitous in soil (Martin and Travers 1989). The lowest percentage recovery of Bt from soil reported was in the USA (60% of soils sampled) (Meadows 1993). In New Zealand, (Chilcott and Wigley 1993) found that between 60-100% of soils sampled contained Bt, depending on source (urban, horticulture etc.). Bt is also indigenous in many other environments, being found in stored products, dust, on deciduous and coniferous plants and in aquatic environments. Bt has also been isolated from insect habitats such as rotting wood, wasp nests and stored products.

There are several theories on the ecological niche filled by Bt. Unlike most insect pathogenic microbes, Bts generally recycle poorly and rarely cause natural epizootics in insects, leading to speculation that Bt is essentially a soil micro-organism that possesses incidental insecticidal activity (Martin and Travers 1989). Evidence to support this view is that Bts are commonly reported in the environment independent of insects and there is a lack of association between occurrence and insect activity (van Frankenhuyzen 1993). Meadows (1993) suggested four possible explanations for the presence of Bt in soil: 1) rarely grows in soil but is deposited there by insects; 2) may be infective to soil-dwelling insects (as yet undiscovered); 3) may grow in soil when nutrients are available; and 4) an affinity with *B. cereus*. Alternatively, Smith and Couche (1991) proposed that Bt is a natural component of the phylloplanemicro flora and has evolved in a symbiotic or mutualistic association with plants to provide protection.

2. ECONOMIC IMPORTANCE OF *BACILLUS THURNIGIENSIS*

2.1. Plant Growth Promoting Agents

Microbial production of secondary metabolites that can promote or constrain plant growth has often been found to be finely tuned, controlled by environmental conditions, medium components, and influenced by root activities. Plant-stimulatory effects exerted by plant growth-promoting bacteria (PGPB) might also be due to an enhanced availability of limited plant nutrients such as nitrogen, phosphorus, vitamins and amino acids in the rhizosphere, caused by phosphate-solubilizing and diazotrophic bacteria (Rozycki, 1999). *Bacillus* is the most abundant genus in the rhizosphere, and the PGP activity of some of these strains has been known for many years, resulting in a broad knowledge of the mechanisms involved (Probanza et al., 2002).

2.2. Persistence and Activity in the Environment

Bacillus thuringiensis is the most common environmentally-friendly insecticide used and is the basis of over 90% of the pesticides available in the market today (Cherifet et al., 2007). Having sequenced the entire genome, scientists can witness and determine how mutations can create different variant strains that may eventually grow stronger against agricultural pests. *Bacillus thurnigiensis* has many properties which make it a superior mosquito control agent; for example it is environmentally benign, easily produced and host specific. One characteristic which contributes to a lack of environmental risk is lack of persistence. Generally, Bt persists for days rather than months as reported for some of the more toxic chemicals for mosquito control. Early reports showed that a primary powder formulation of Bt (serotype H-14) had virtually no residual effect against mosquito larvae beyond application, although the delta-endotoxin remained chemically stable in neutral and acid waters (Sinegreet et al. 1980).

2.3. As Biotechnological Agents in Industrial Processes and Production

Bacillus thuringiensis produces the Cry crystal toxin that attacks the gut of pests and kills them internally. These Cry toxins are good agricultural tools for growing plants. Instead of using chemicals that may have adverse effects on humans, genetic engineers integrate the *Bt* toxin into the plant's genome. These *Bt* Cry toxins are safe for humans and kill off species of pests that are susceptible to the *Bacillus thuringiensis* endospore. Each single strain of organism produces a large number of enzymes which include the function such as hydrolyzing, oxidizing or reducing, and metabolic in nature. But the absolute and relative amounts of the various individual enzymes produced vary markedly between species and even between strains of the same species. Amylase, lipase, protease and cellulase constitute a very important part of microbial enzymes that are used in food, pharmaceutical, textile, paper, leather, and other industries. It is estimated that *Bacillus* spp. enzymes make up about 50% of the total enzyme market (Schallmeyer *et al.*, 2004).

Food fermentations mediated by *Bacillus*, can provide insight into some of the potential industrial properties of the genera or species involved. Recent developments in genetic engineering and fermentation technology have contributed to improvements in D-ribose productivity by *Bacillus* fermentations. Most members of the genus *Bacillus* are able to produce antibiotics. Interestingly, the majority of these antibiotics are low-molecular-weight peptides, which possess different biological activities, including antimicrobial, antiviral, and antitumor activities (Giacomodonato *et al.*, 2001).

2.4. *Bacillus Thurnigiensis* Insecticidal Agents

Global use of insecticides for mosquito vector control in recent decades has caused environmental pollution of aqueous ecosystem and resulted in insecticide resistance in many mosquito species. The last decade has witnessed and increased interest in biological control agents. Biological means to control vectors, based on entomopathogenic bacteria has been studied for more than 20 years. More number of biological control agents was screened for their efficacy, mammalian safety and environmental impact. Many organisms have been investigated as potential agents for vector mosquito control, including viruses, fungi, bacteria, protozoa, nematodes, invertebrate predators and fish (Zulfaidah P. and Nobukazu N. 2014). Only a few spore forming bacteria, copepods and fish have reached operational use and are undergoing extensive field trials. The discovery of bacteria like *B. thuringiensis serovar israelensis* de Barjac, which are highly toxic to dipteran larvae have opened up the possibility of its use as potential bio-larvicides in mosquito eradication programs in the over the world (Poopathi and Tyagi, 2002; Poopathi *et al.*, 2002). The larvicidal substances of these preparations are based on endotoxin proteins accumulated as parasporal crystals produced by the bacterial cells during the sporulation growth phase. These biological preparations have some important advantages over conventional insecticides in mosquito control operations, besides being safe to nontarget organism including human beings. Also, it is harmless to the environment (Prabakaran and Balaraman, 2006). The *B. thuringiensis serovar israelensis* has been used operationally for the control of mosquitoes for over two decades and its formulations are highly effective against *Anopheles*, *Aedes* and *Culex* mosquitoes (Poopathi and Abidha, 2010).

3. HOST RANGE

Bacillus thurnigiensis is highly pathogenic against Culicidae (mosquitoes) and Simuliidae (blackflies), and has some virulence against certain other Diptera, especially Chironomidae (midges). There are few records of susceptibility outside these dipteran groups, and most other hosts within Diptera require high doses to kill. *Bt* is generally regarded as specific to larvae of the Nematocera, which include filter-feeding mosquitoes and Simulium (blackflies, sandflies) (Clarke, 1994). However, the vast majority of susceptible hosts are recorded in nematoceros Diptera. Among mosquitoes, different preparations of *Bt* have shown differing levels of toxicity to host species. Generally, *Culex* and *Aedes* are highly susceptible while *Anopheles* are less susceptible, but can still be killed with *Bt* (Balaraman *et al.*, 1983). However, even within one genus, some species are more susceptible than others (Chui *et al.*, 1993).

4. FUTURE PROSPECTS OF *BACILLUS THURNIGIENSIS*

The use of Bt spray as an insecticide has several disadvantages; 1) Bt spray cannot be applied uniformly to all parts of the plant, 2) it cannot be delivered to pests that are inside plant tissues, and 3) Bt is susceptible to rapid degradation by UV light and removal by water runoff. Therefore, multiple applications are required to provide extended pest protection (Schnepfet *et al.*, 1998). Moreover, since McGaughey reported resistance to Bt in Indian meal moth populations (Gonzalez *et al.*, 1982) resistance to Bt sprays has also evolved in greenhouse populations of the cabbage looper and in field populations of the diamondback moth (Tabashnik, B. E. 1994). Transgenic crops with Bt cry genes might overcome these kinds of disadvantages.

This feature eliminates difficulties in targeting pests that burrow into plant tissues, as well as the labor and expenses associated with applying sprays. At present, field-evolved resistance to Bt crops has not been documented. (Tabashnik *et al.*, 2005) reported long-term resistance levels in transgenic Bt crop fields. Generally, it has been a concern that an increase in the wide planting of Bt crops might lead to rapid evolution of resistance to Bt toxins by pests (Tabashnik, B. E. 1994).

However, bioassay results show no net increase from 1997 to 2004 in the mean frequency of pink bollworm resistance to Bt toxin (Tabashnik *et al.*, 2005). This delay in resistance can be explained by the presence of refuges of cotton without Bt toxin, recessive inheritance of resistance, incomplete resistance, and fitness costs associated with resistance.

Basically, the advantages of Bt and Bt crops apparently includes 1) no harmful effects on vertebrates and humans, or the ecological environment, 2) low impacts on non target organisms, and 3) a narrow spectrum of primarily leaf-feeding lepidopteran targets. These merits, including the short field life of the spray, are still this biopesticide's greatest challenge in the pesticide market (Whalon, M. E. and B. A. Wingerd. 2003). For the sustainable use of Bt, it is imperative that there be 1) collections of Bt isolates, crystal proteins, and strains of related species, 2) investigations into the persistence of crystal proteins and possible long-term effects on non target organisms and the environment, 3) development of improved resistance management strategies, and 4) genetic engineering of Bt genes into the plastid genomes of transgenic crops (Nester *et al.*, 2002). Environmentally safe-insect control strategies based on the Bts and their insecticidal crystal proteins are going to increase in the future, especially with the wide adoption of transgenic crops. In conclusion, the discovery of new toxins and new ways of presenting the toxin to the target insects, which includes the development of recombinant microorganisms and proteomic technology, could be adapted to the study of Bt crystal proteins; additionally, interaction studies between Bt and target insects involving modes of action of Bt Cry proteins and resistance mechanisms should be carried out, all of which are fundamental studies that will allow for improvement of existing Bt application strategies and the ability to design alternative options.

5. CONCLUSION

According to the studies cited in different literatures *Bacillus thurnigiensis* reported as the most effective and playing important roles as dominant bacteria in biotechnological applications as well as industrial processes and products. And, with recent advances in biotechnology, the economic contributions that these organisms can make in biotechnological applications and industrial processes can be exploited further for large scale benefit of mankind. Therefore, Ethiopian Biodiversity Institute Microbial Directorate collected, isolated and identified several *Bacillus* spp. And *Bacillus thurnigiensis* is one of them collected across the country. There are number of potential *Bacillus* species in the gene bank of the institute. One of the major *Bacillus* species are *Bacillus thuringiensis*. Therefore, we encourage any bio prospecting company to access this genetic resource for its potentials in biotechnological applications as well as industrial processes and products for access and benefit sharing.

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