

## Original Research Article

# Isolation and identification of microbial deteriogens of fresh tomatoes stored at ambient temperature

### Abstract

- This study investigated the microorganisms associated with the deterioration of fresh tomato, *Lycopersicum esculentum*, stored at ambient temperature. A total of 60 tomatoes obtained from three different markets in Owerri metropolis, Nigeria were sampled and a total of 134 bacteria isolates and 140 fungal isolates were encountered. Bacteria isolated and identified were *Bacillus spp*, *Pseudomonas spp*, *Escherichia coli*, *Klebsiella spp* and *Salmonella spp*. The most prevalent bacteria isolate was *Bacillus spp* with 52.24% while *Salmonella spp* was the lowest prevailing isolate with 1.49%. The fungal isolates were *Fusarium spp*, *Aspergillus spp*, *Rhizopus spp*, *Saccharomyces spp*. Whereas *Aspergillus spp* was the most prevalent with 60.71% and was found in tomato sample from all the market, *Rhizopus spp* had the least prevalence of 0.71%. The presence of toxin producing fungi, *Aspergillus spp* and *Fusarium spp* which are capable of causing food poisoning as well as some bacterial isolates, raises concern over public health risks that may be associated with the consumption of unwholesome tomato.

### Introduction

Tomato (*Lycopersicum esculentum*) is a berry plant in the *solanaceae* family. It is a short lived perennial plant grown as an annual plant, typically growing about 3-5m approximately in height. Tomatoes are brightly red coloured berry which are eaten fresh in salads or processed and can be stewed, fried, baked and also used to produce soup, or used as juice. Tomato is one of the widely consumed fresh fruit worldwide since it contributes to a healthy well balanced diet which is rich in vitamins such as vitamin, B, C, and E. Carbohydrates such as fructose and glucose; and trace elements like iron, copper, zinc, and dietary fiber (Faostat, 2010).

Tomatoes grow best in light, free draining, and fertile loamy soil with PH of 5 – 7. Nonetheless, tomatoes can be grown in variety of soils (Naika et al., 2005). Concerning the fertilizer requirement, tomatoes requires an abundance of three vital elements namely, nitrogen, phosphorus, and potassium. The deep red colouration of rippened tomato is due to the presence of LYCOPENE, a form of B-carotenoid pigment and a powerful antioxidant that help to protect against prostate cancer, cardiovascular disease and diabetes, thus there is a great demand for tomatoes by consumer as a result of their healthy, tasty, convenient, and fresh appeal (Passam et al, 2007). Tomato, however, have serious

challenges to their existence. These include changes in climatic condition, pest, bacterial and fungal attack and over the years there has been an increase in the need to identify and isolate the microorganism associated with their deterioration.

Deterioration refers to any change in food, which the food becomes less palatable, or even toxic, these changes may be accompanied by alteration in taste, smell, appearance or texture (Berdegue *et al*, 2005). Numerous microbial defects (Signs and Symptoms) of tomatoes are characterized by the type of microorganism responsible for the deterioration, in the process of infection in the case of fungal invasion follows the development of fungal penetrating structure. The colonization process involves the ability of the microorganism to establish itself within the produce. Susceptibility of tomato to microbial colonization is due to its differential chemical composition such as high level of sugar, low pH (4.9-6.5) and its high water activity ( $p > 0.99$ ) which favors the growth of microorganism in tomato is recognized as a source of potential health hazard to man and animals, this is due to their production of toxins which are capable of causing disease like respiratory infection, meningitis, gastroenteritis, diarrhoea in man following ingestion (Beuchat *et al*, 2006).

Tomatoes was chosen for this study because they are referred to as ready-to-eat food since they are minimally processed and many people take tomatoes raw directly or via meals of salad usually served cold. Microbial spoilage and contaminating pathogens on this product poses a serious

problem in food safety. The Centre for Disease Control and Prevention (CDC) estimates that there are 76 million cases of food borne illness every year. Outbreaks with identified etiology are predominantly of microbial origins (Bihn *et al*, 2006), so that studying about the microbial ecosystem of fresh raw and spoilt tomato is necessary.

## **Materials and methods**

Materials used for these analyses were standard microbiological materials and were gotten from the microbiology laboratory except the tomato samples that were purchased from different markets.

### **Samples collection**

Fresh tomatoes were obtained from the major markets in owerri metropolis, which include Ekeonunwa, Relief, and student market. A total of 60 tomatoes, 20 tomatoes from each market were sampled. The ripened tomato fruits selected were fresh, undamaged, firm and healthy. The sample were taken to the laboratory, washed and drained of water. The fruit samples were kept from dust and insects at room temperature for up to 14 days to undergo a natural process of deterioration before being used for study.

### **Processing of fresh spoilt tomatoes stored at ambient temperature**

The tomatoes fruit samples were ground using a sterile mortar and pestle. A homogenate of each sample was made by blending one gram in 9ml of sterile water and shaking them together. Serial dilution of

up to  $10^5$  of the homogenate was made in sterile test tubes.

### **Enumeration of microbial load**

A ten-fold serial dilution of each of the samples was carried out. Spread plate technique was employed by inoculating 0.1ml aliquot aseptically from the  $10^{-3}$  and  $10^{-4}$  dilutions onto nutrient and MacConkey agar plates for enumeration of bacteria and Sabouraud dextrose agar for fungi count. The agar plates were incubated at  $37^\circ\text{C}$  for 24-48 hours for bacterial count and at  $25^\circ\text{C}$  -  $28^\circ\text{C}$  for 5-8 days for fungal count. Each sample was inoculated in duplicate agar plates and the mean values of bacterial and fungal counts were recorded as colony forming unit per ml (cfu/ml).

### **Purification (subculture) of bacterial isolates**

Colonies from the primary plates were aseptically picked with a sterile wire loop and transferred onto freshly prepared sterile nutrient agar plate, with a streaking technique such that discrete colonies appear at the ends of streaked lines after incubation. The subculture plates were incubated at  $37^\circ\text{C}$  for 24 hours to 48 hours. Discrete colonies from the subculture plates were aseptically transferred and streaked on slant and incubated for another 24 hours at  $37^\circ\text{C}$  which were stored at  $4^\circ\text{C}$  and used subsequently for microscopic characterization and biochemical analyses.

### **Purification of fungal isolates**

Colonies from the primary plates were aseptically picked with a sterile inoculation

needle and transferred onto a freshly prepared sterile Sabouraud Dextrose Agar (SDA) plate with a streaking method and incubated for 5- 7 days at  $25^\circ\text{C}$ - $28^\circ\text{C}$ . Discrete colonies were aseptically transferred and stocked on slant and incubated for another 5 days at  $25^\circ\text{C}$ - $28^\circ\text{C}$ . Pure colonies were stored in the refrigerator at  $10^\circ\text{C}$ - $15^\circ\text{C}$  until needed for characterization and identification.

### **Characterization and identification of bacterial isolates**

All bacterial isolates were characterized and identified based on their cultural, morphological, microscopic examination and biochemical characteristics following the methods prescribed by (Cheesbrough, 2005). Biochemical test conducted include the following: Gram stain, Catalase test, Oxidase test, Motility test, Methyl red test, Citrate test and Urease test.

### **Identification of fungi isolate**

The complete identification of fungi isolate was done by comparing the result of their morphological characteristics with those of known taxa.

### **Results and Discussion**

The results on the isolation and identification of microbial deteriogens of fresh tomato stored at ambient temperature are described as follows: The total heterotrophic count, total coliform count, total fungi counts are shown in table1. The result shows that tomato fruit samples from Relief market recorded the highest bacterial

and fungal count while the samples from Ekeonuwa market recorded the lowest bacterial and fungal count. Table 2 shows the characterization and identification of the isolates following biochemical procedures. The bacteria isolates identified from spoilt fresh tomatoes stored at ambient temperature were: *Bacillus spp*, *Escherichia coli*, *Pseudomonas spp*, *salmonella spp*, *Klebsiella spp*. The result on the identification of fungi are shown in table 3, the fungal isolates identified from spoilt fresh tomatoes stored at ambient temperature were *Fusarium spp*, *Aspergillus spp*, *Rhizopus spp*, *Saccharomyces spp*. The percentage occurrence of bacteria isolates from fruit samples obtained from different markets are presented in table 4 which indicates that From all the tomato fruit samples obtained from the 3 market, 134 bacteria was isolated of which *Bacillus spp* was the most prevalent with 52.24%, *Escherichia coli*(14.93%), *Pseudomonas spp*(23.88%), *salmonella spp* (1.49%), *Klebsiella spp* (6.72%).The isolation of soil bacteria *Bacillus spp*, from the fruit samples, suggests evidence of opportunistic contamination from human activity. The percentage occurrence of fungi isolates as presented in table 5 are as follows: in the characterization of fungi, a total of 140 organism were isolated of which 85(60.71%) were *Aspergillus spp*, followed

by *Fusarium spp* 44(31.43%), *Saccharomyces spp* 10(7.14%), *Rhizopus spp* was not significant ( $P>0.05$ ). Similar findings were reported by (Gosh,2009) who also asserted that *Aspergillus spp*, and *Fusarium spp*. were the major microorganisms that are responsible for the spoilage of tomato fruits. The implications of microbial contamination and growth on tomato produce causes spoilage, decreased sensory appeal and decreased shelf life leading to loss and wastage of product that have significant economic consequences. The microbiological safety of these products has also become a significant issue, as the incidence of food borne disease outbreaks associated with their consumption.

## Conclusion

As a consumer we need to recognize that food safety is important for fresh fruits and vegetable. Also individuals of the population especially those in developing countries who usually use spoilt and slightly decaying tomatoes as a result of their cheaper prices should be educated that these spoilage are often not due to mechanical damages but microbial colonization and physiological decays, they should be made to know that these organisms produces toxins and spores which are relatively heat resistant and can cause severe food poisoning resulting in fatal outcome.

**TABLE1: MICROBIAL LOAD OF SPOILT FRESH TOMATOES STORED AT AMBIENT TEMPERATURE (cfu/ml)**

	<b>Sample 1</b>	<b>Sample 2</b>	<b>Sample 3</b>
<b>Total bacterial count</b>	3.6x10 <sup>4</sup>	5.0x 10 <sup>4</sup>	4.8x10 <sup>4</sup>
<b>Total coliform count</b>	1.6x10 <sup>4</sup>	2.5x10 <sup>4</sup>	2.0x10 <sup>4</sup>
<b>Total fungi count</b>	4.0x10 <sup>4</sup>	5.3x10 <sup>4</sup>	4.6x10 <sup>4</sup>

**KEY**

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**Sample 1**= Result from tomatoes gotten from ekonunwa

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**Sample 2**= Result from tomatoes gotten from relief

**Sample 3** = Result from tomatoes gotten from Student market

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									Suspecting
	Size	Shape	Elevation	Margin	Surface	Opacity	Colour	texture	Organism
Sample 1(A)	Small	Round	Slighty raised	Entires	Smooth	Shiny	Grayish white	Mucoid	<i>Escherichia coli</i>
Sample 1(b)	Large	Round	Umbonate	Undulate	Smooth	Translucent	White	Mucoid	<i>Klebsiella spp</i>
Sample 1(c)	Medium	Oval	Unbonate	Wavy	Smooth	Transparent	Diffusible green	Mucoid	<i>Pseudomonas spp</i>
Sample 1(D)	Large	Irregular	Umbonate	Undulate	Rough	Opaque	White	Dry	<i>Bacillus spp</i>
Sample 2(A)	Medium	Oval	Unbonate	Wavy	Smooth	Transparent	Diffusible green	Mucoid	<i>Pseudomonas spp</i>
Sample 2(B)	Small	Round	Slighty raised	Entires	Smooth	Shiny	Grayish white	Mucoid	<i>Escherichia coli</i>

									Suspecting
	Size	Shape	Elevation	Margin	Surface	Opacity	Colour	texture	Organism
Sample2(C)	Large	irregular	Umbonate	Undulate	Rough	Opaque	White	Dry	<i>Bacillus spp</i>
Sample 2(D)	Large	Round	Umbonate	Undulate	Smooth	Translucent	White	Mucoid	<i>Klebsiella spp</i>
Sample 3(A)	Small	Round	Slightly raised	Entire	Smooth	Shiny	Grayish white	Mucoid	<i>Escherichia coli</i>
Sample 3(B)	Large	Round	Umbonate	Undulate	Smooth	Translucent	White	Mucoid	<i>Klebsiella spp</i>
Sample 3(C)	Medium	Oval	Unbonate	Wavy	Smooth	Transparent	Diffusible green	Mucoid	<i>Pseudomonas spp</i>
Sample 3(D)	Large	irregular	Umbonate	Undulate	Rough	Opaque	White	Dry	<i>Bacillus spp</i>

## COLONIAL MORPHOLOGY OF BACTERIA ISOLATES

**TABLE 2 Identification of bacteria isolates**

A, B, C, D, E represents each colony on the culture plate from each sample

**TABLE 2.1      BIOCHEMICAL CHARACTERISTICS OF BACTERIAL ISOLATES**

	Gram staining	Catal ase	Coagulas e	Methyl red	Vp test (Voges proskauer)	motility	Citrate test	Indole test	Oxidase test	Probable organism
Sample 1(A)	-ve single rods	+ve	-ve	+ve	+ve	+ve	-ve	+ve	-ve	<i>E.coli</i>
Sample 1(b)	-ve single rods	+ve	-ve	-ve	+ve	-ve	+ve	-ve	-ve	<i>Klebsiella spp</i>
Sample1(c)	-ve single rods	+ve	-ve	-ve	-ve	+ve	+ve	-ve	+ve	<i>Pseudomonas spp</i>
Sample 1(D)	+ve single rods	+ve	-ve	-ve	+ve	+ve	-ve	-ve	+ve	<i>Bacillus spp</i>
Sample2(A)	-ve single rods	+ve	-ve	-ve	-ve	+ve	+ve	-ve	+ve	<i>Pseudomonas spp</i>
Sample 2(B)	-ve single rods	+ve	-ve	+ve	+ve	+ve	-ve	+ve	-ve	<i>E.coli</i>
Sample2(C)	+ve single	+ve	-ve	-ve	+ve	+ve	-ve	-ve	+ve	<i>Bacillus spp</i>

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	rods										
Sample 2(D)	-ve	single	+ve	-ve	-ve	+ve	-ve	+ve	-ve	-ve	<i>Klebsiella spp</i>
	rods										
Sample 3(A)	-ve	single	+ve	-ve	+ve	+ve	+ve	-ve	+ve	-ve	<i>E.coli</i>
	rods										
Sample 3(B)	-ve	single	+ve	-ve	-ve	+ve	-ve	+ve	-ve	-ve	<i>Klebsiella spp</i>
	rods										
Sample 3(C)	-ve	single	+ve	-ve	-ve	-ve	+ve	+ve	-ve	+ve	<i>Pseudomonas spp</i>
	rods										
Sample 3(D)	+ve	single	+ve	-ve	-ve	+ve	+ve	-ve	-ve	+ve	<i>Bacillus spp</i>
	rods										
Sample 3(E)	-ve	single	+ve	-ve	+ve	-ve	+ve	-ve	-ve	-ve	<i>Salmonella spp</i>
	rods										

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**KEY:**

**+ = Positive**

**- = Negative**

**TABLE 3: IDENTIFICATION OF FUNGAL ISOLATES**

	<b>Macroscopy</b>	<b>Microscopy</b>	<b>Fungal isolates</b>
<b>Sample 1(A)</b>	Greenish, filamentous with profuse proliferation of black velvety spores	Septate hyphae, branched conidiophore with secondary branches. The conidiophore is enlarged at the tip forming rounding vesicle-like chains	<i>Aspergillus spp</i>
<b>Sample2(B)</b>	Initially white and cottony but later develop pink centre with a lighter periphery.	Septate hyphae with canoe-shaped macroconidia, conidiophores bear conidia singly or in cluster.	<i>Fusarium sp.</i>
<b>Sample2(A)</b>	Initially white and cottony but later develop pink centre with a lighter periphery.	Septate hyphae with canoe-shaped macroconidia, conidiophores bear conidia singly or in cluster.	<i>Fusarium sp.</i>
<b>Sample2(B)</b>	Greenish, filamentous with profuse proliferation of black velvety spores	Septate hyphae, branched conidiophore with secondary branches. The conidiophore is enlarged at the tip forming rounding vesicle-like chains	<i>Aspergillus spp</i>

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	Macroscopy	Microscopy	Fungal isolates
<b>Sample3(A)</b>	Initially white and cottony but later develop pink centre with a lighter periphery.	Septate hyphae with canoe-shaped macroconidia, conidiophores bear conidia singly or in cluster.	<i>Fusarium sp.</i>
<b>Sample3(B)</b>	Greenish, filamentous with profuse proliferation of black velvety spores	Septate hyphae, branched conidiophore with secondary branches. The conidiophore is enlarged at the tip forming rounding vesicle-like chains	<i>Aspergillus spp</i>
<b>Sample3(C)</b>	Cottony white, filamentous, coenocytic, stolons, rhizoids.	Ovoid sporangiospores, tall sporangiospores in groups	<i>Rhizopus spp</i>
<b>Sample3(D)</b>	Colonies of <i>Saccharomyces sp.</i> grow rapidly. They are flat, smooth, moist glistening or dull, and cream to tannish cream in color	Multilateral budding is typical Pseudohyphae, if present are rudimentary. Hyphae are absent. <i>Saccharomyces sp.</i> produces ascospores.	<i>Sacharomyces spp</i>

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**TABLE 4: THE PERCENTAGE OCCURRENCE OF BACTERIA ISOLATES IN SPOILT FRESH TOMATOES STORED AT AMBIENT TEMPERATURE**

Bacteria isolates	Sample 1(no of colonies)	Sample 2(no of colonies)	Sample 3(no of colonies)	Frequency	Percentage(%)
<i>Bacillus spp</i>	20	25	25	70	52.24
<i>Pseudomonas spp</i>	8	10	14	32	23.88
<i>Salmonella spp</i>	Nil	Nil	2	2	1.49
<i>Escherichia coli</i>	5	10	5	20	14.93
<i>Klebsiella spp</i>	3	5	1	9	6.72
Total	36	50	48	134	100

#### **KEY**

**Sample 1**= Result from tomatoes gotten from ekonunwa market

**Sample 2**= Result from tomatoes gotten from relief market

**Sample 3** = Result from tomatoes gotten from Student market

**TABLE 5: THE PERCENTAGE OCCURRENCE OF FUNGI ISOLATES IN SPOILT FRESH TOMATOES STORED AT AMBIENT TEMPERATURE**

Fungal isolates	Sample1	Sample 2	Sample 3	Percentage (%)
<i>Fusarium spp</i>	20	16	8	31.43
<i>Aspergillus spp</i>	20	37	28	60.71
<i>Rhizopus spp</i>	Nil	1	Nil	0.71
<i>Saccharomyces spp</i>	Nil	Nil	10	7.14
Total	40	54	46	100

**Sample 1**= Result from tomatoes gotten from ekonunwa market

**Sample 2**= Result from tomatoes gotten from relief market

**Sample 3** = Result from tomatoes gotten from Student market

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