

## Milk Decontamination and an Effective Milk Pasteurizing Method for Household Dairies

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This study investigated the bacteriological and the hygienic properties of some commonly sold canned milk and determined the effectiveness of pasteurization of milk contaminated with bacterial isolates spectrophotometrically. In this study, *Escherichia coli*, *Streptococcus spp.*, *Enterococcus aerogenes*, *Proteus spp.*, *Clostridium spp.*, *Lactobacillus spp.* and *Staphylococcus aureus* were isolated from the milk samples. The bacterial load of the milk samples ranged from  $5 \times 10^5$  to  $22 \times 10^5$  cfu/ml. The methylene reduction time indicating the hygienic quality of each milk sample varied from one milk to the other. While NNM and VSM samples not decolourized after 8 h were considered excellent and PNM and OFM decolourized methylene blue after 6 and 7 h respectively were considered good milk samples, CFM and TCM that decolourized methylene blue after 4 and 5 h respectively were considered fair milk samples. None of them was considered poor as they were not decolourized in less than 2 h. Pasteurizing milk contaminated with indigenous microflora (IMF), *E. coli* and *Strep. pyogenes* at 40°C, 50°C, 60°C and 70°C over a period of 30 min showed a varied degree of reduction in the absorbances and its effectiveness. A 10°C increase in temperature resulted in significant reduction in the absorbances obtained at different time intervals for all the milk samples. This investigation shows that pasteurizing milk with low heating would reduce the microbial loads of the milk samples. The method of pasteurizing milk in this study would be applicable in the household dairies where sophisticated equipment are not required to control microbial loads in milk.

**Keywords:** Contamination, methylene blue reduction, indigenous microflora, pasteurization.

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Milk is a highly nourishing and important food with potential therapeutic properties. It is an all-inclusive food consumed in a variety of different products by millions of people. Rolf<sup>1</sup>

described milk as colloids of butterfat globules within a water-based fluid containing dissolved carbohydrates. While containing proteins, fat, sugar, vitamins and minerals<sup>2</sup>, 97–98% lipids in milk are triacylglycerols, small amounts of di- and monoacylglycerols, free cholesterol and cholesterol esters, free fatty acids and phospholipids<sup>3</sup>. The fat portion of the milk contains fat-soluble vitamins,

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linoleic and linolenic acid<sup>4</sup>. Being a delicate agricultural product and an enrichment medium, its nutrient compositions easily support bacterial growth<sup>5-8</sup>. It may be contaminated from animal skin, environment, utensils and water used for adulteration<sup>9</sup> and the quality of the contaminated milk may deteriorate during transportation at ambient temperature<sup>10</sup>. Consequently, many bacterial strains including *Listeria monocytogenes*, *Salmonella sp.*, *Campylobacter sp.*, *Staphylococcus aureus*, *Bacillus cereus*, *Escherichia coli* and some species of *Streptococcus* and *Micrococcus* have been reported as pathogenic and spoilage bacteria in milk<sup>11-16</sup>. While many contaminating bacteria spoil the milk and others cause the souring of milk and its products naturally, many are known to cause gastroenteritis and diarrhea<sup>17</sup>.

Although De Man et al.,<sup>18</sup> and Kleter,<sup>19</sup> had earlier reported that freshly harvested milk usually have a low microbial load with total count usually less than  $10^3$  cfu/ml, Chye et al.<sup>20</sup> showed that the microbial load could increase when stored at ambient temperature for a long period of time. While the microbial content of milk determines its quality<sup>21,22</sup>, the presence of high microbial load in milk poses major economic loss for farmers and household dairies as milk price is determined by the value of the pathogenic bacteria count<sup>23,24</sup>. Since Vasavada<sup>25</sup> considered milk as a major vehicle for transmitting infectious agents, Coorevits et al.<sup>26</sup> and FAO<sup>9</sup> reported that bacterial contamination of milk could be from different sources. Consequentially, since careful hygiene is needed during processing to avoid microbial contamination and there is a significant threat to milk recipients who are vulnerable to food-borne infectious diseases due to low age and co-morbidities<sup>27</sup>, determining the bacteria load, characterizing the indigenous flora, pasteurizing and determining the absorbance rate of bacterial contaminated milk samples at different temperatures for varying durations and the potential of successfully practicing low scale pasteurization procedure becomes inevitable to ensure the quality and safety of milk consumed by the populace. This study was, therefore, aimed at investigating the bacterial contaminants and the effectiveness of pasteurizing milk contaminated with bacterial pathogens in a miniature experiment to show the possibility of pasteurizing milk by small scale farmers in a cost effective manner.

## MATERIALS AND METHODS

### Source of samples

Five different evaporated canned milk and a soy milk sample were purchased randomly at different markets located in Ogun State, Nigeria. Each sample was labeled and surface sterilized with 70% ethanol before being opened with a heat sterilized tin cutter. Each milk sample was emptied in labeled sterilized autoclave-able plastic bottles. The milk samples were kept in ice packs before being kept in refrigerator at 3°C.

### Total viable aerobic bacterial count

Total viable aerobic bacterial and coliform counts of the milk samples was determined as described by Collins et al.<sup>28</sup>. Each of the samples was diluted tenfold in properly labeled 20 ml glass tubes containing 9 ml sterile normal saline as diluent. Each sample was aseptically diluted serially with separate sterile glass pipette (10 ml). One ml sample from tube dilution number  $10^{-8}$ ,  $10^{-7}$ ,  $10^{-6}$ ,  $10^{-5}$  and  $10^{-4}$  was dispensed on surfaces of nutrient agar and Eosin Methylene Blue (EMB) agar plates in triplicate before being spread evenly with sterile glass spreader. After incubating each plate at 37°C for 48 h, total colony forming units (cfu) of aerobic bacterial and coliform counts were recorded from the plate having 30-300 colonies.

### Isolation and identification of isolates from milk samples

Serially diluted milk samples were plated in duplicate in nutrient broth and McConkey broth using pour plate technique. Plates were incubated at 37°C for 24 - 48 h. Total viable counts were carried out on nutrient agar while the presence or absence of some organisms was done by plating on selective media. Pure cultures of discrete colonies were made by repeated sub-culturing on sterile nutrient agar with a sterile wire loop. Using overnight cultures, the pure strains were identified and confirmed using morphological, microscopy and biochemical tests following standard procedures described by Cowan et al.,<sup>29</sup> and Cheesborough<sup>30</sup>. Results were analyzed using Bergy's manual and other methods for bacterial identification<sup>31-33</sup>.

### Methylene blue reduction test

Methylene blue reduction test (MBR) was conducted to determine the hygienic status of each of the milk samples<sup>34</sup>. One ml of methylene blue solution (1:25000) was transferred to labeled and

sterile 20 ml screw capped test tube containing 10 ml of each of the milk samples. The tubes were capped and gently inverted three times to mix up the dye with each milk sample. Each of the tubes was incubated at 37°C and examined for discolourization at intervals up to 8 h. The time taken by the methylene blue in each milk sample to become colorless was recorded. The methylene blue reduction test depends upon the ability of bacteria in milk to grow and to consume the dissolved oxygen which reduces the oxidation-reduction potentials in the medium.

**Pasteurization and Spectrophotometry at different temperature and varied time intervals**

Pure strains of *Escherichia coli* and *Streptococcus pyogenes* were obtained from the

**Table 1.** Bacteria isolated from the different milk samples

Sample code	Gram reactions	Isolated bacteria
NNM	GNB	<i>Escherichia coli</i>
	GPC	<i>Streptococcus spp.</i>
	GNB	<i>Enterobacter aerogenes</i>
VSM	GNB	<i>Proteus spp</i>
	GPB	<i>Clostridium spp</i>
TCM	GPB	<i>Lactobacillus spp</i>
CFM	GPB	<i>Clostridium spp</i>
	GPC	<i>Staphylococcus aureus</i>
PNM	GNB	<i>Escherichia coli</i>
	GPB	<i>Lactobacillus spp</i>
OFM	GNB	<i>Proteus spp</i>

KEY: + Positive, - Negative, GPC - Gram Positive Cocci, GPB - Gram Positive Bacilli, GNB -Gram Negative Bacilli.

**Table 2.** Results for Bacterial Count of the different milk samples

Sample Codes	Number Of Colonies (cfu/ml)
TCM	6 x 10 <sup>5</sup>
NNM	9 x 10 <sup>5</sup>
CFM	19 x 10 <sup>5</sup>
OFM	16 x 10 <sup>5</sup>
PNM	22 x 10 <sup>5</sup>
VSM	5 x 10 <sup>5</sup>

Medical Laboratory Department of Babcock University Teaching Hospital, Ilishan Remo, Ogun State, Nigeria. The identity of these isolates were further confirmed by microbiological and biochemical tests. Overnight broth cultures of these bacteria were centrifuged at 1500 rpm for 5 min. The supernatants were discarded while the pellets were resuspended in normal saline. These were then centrifuged at 15000 rpm for 5 min and the supernatants were discarded again. The pellets were then resuspended in the normal saline. The absorbances of the bacterial suspensions were measured spectrophotometrically at a wavelength of 600 nm<sup>35</sup>. Dilutions of 10<sup>6</sup> cfu/ml of each each isolate from milk samples were prepared in triplicates. A predetermined volume of the diluted milk samples (9 ml) were pipette into the test tubes and 1 ml of adjusted *E. coli* and *S. pyogenes* suspensions was pipetted into two different sets of test tubes while the third set containing the indigenous flora (IMF) of the milk sample was not inoculated. The test tubes were then placed in the water bath at varying temperatures (40°C, 50°C, 60°C and 70°C). At specific time intervals of 10, 20 and 30 min, the absorbances of each contaminated milk sample were measured at a constant wavelength of 600 nm. The contaminated milk samples were later subcultured to check for the effectiveness of pasteurization procedure.

**RESULTS**

In this study, milk samples were tested for the presence of bacterial contaminants before being pasteurized. The pasteurized milk

**Table 3.** Results for methylene blue reduction test to determine the hygienic status of different milk samples

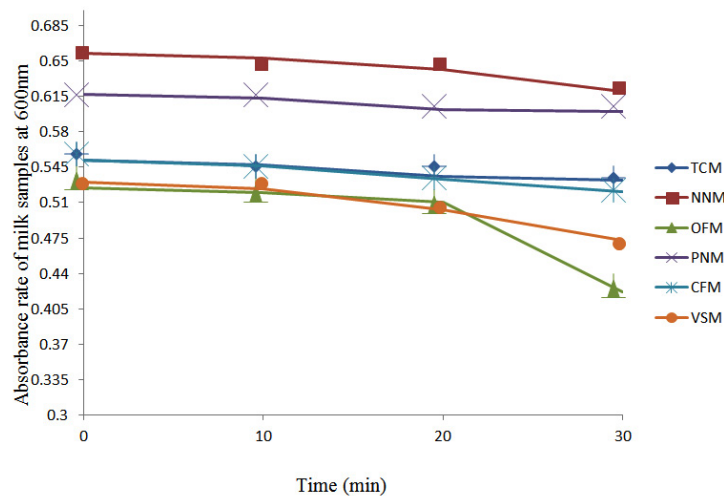
Sample Code	Methylene Blue Reduction Time (h)	Remark	Class
CFM	4	fair	3
TCM	5	fair	3
PNM	6	good	2
NNM	Not Decolourized	excellent	1
OFM	7	good	2
VSM	Not Decolourized	excellent	1

Key: Class 1 = > 8 h; Class 2 = < 8 h; Class 3 = < 6 h but not < 2 h.

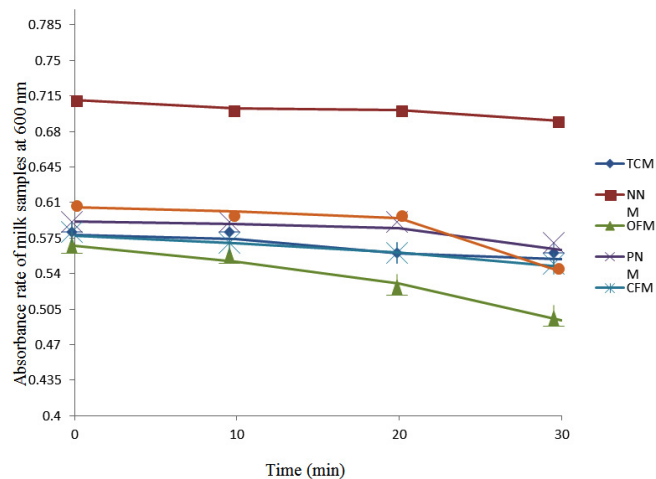
samples were then tested for methylene blue reduction to determine their hygienic status before being contaminated with the indigenous microbial flora (IMF), *Escherichia coli* and *Streptococcus pyogenes*. The bacteria isolated from the milk samples included *Escherichia coli*, *Streptococcus spp.*, *Enterococcus aerogenes*, *Proteus spp.*, *Clostridium spp.*, *Lactobacillus spp.* and *Staphylococcus aureus*. The result showed that the isolates from the milk samples included Gram negative bacilli (GNB) 45.45%, Gram positive cocci (GPC) 36.36% and Gram positive bacilli (GPC) 18.18% (Table 1). The bacterial loads of the different milk samples were negligible and ranged from  $5 \times 10^5$  to  $22 \times 10^5$  cfu/ml (Table 2). The

methylene reduction time indicating the hygienic quality of each milk sample varied from one milk to the other. While milk samples NNM and VSM were not decolourized after 8 h, others were decolourized between 4 and 7 h. The milk samples not decolourized were considered excellent. PNM and OFM decolourized after 6 and 7 h respectively were considered good milk samples while CFM and TCM decolourized after 4 and 5 h respectively were considered fair milk samples. None of the milk samples was considered poor as they were not decolourized in less than 2 h (Table 3).

After determining the initial microflora and the bacterial load of each milk sample, they were pasteurized before being contaminated with



**Fig. 1.** Effect of pasteurization on milk samples contaminated with Indigenous Milk Flora (IMF) at 40°C at 600 nm



**Fig. 2.** Effect of pasteurization on milk samples contaminated with *E. coli* at 40°C at 600 nm

10<sup>6</sup> cfu/ml of the indigenous microbial floral (IMF), *E. coli* and *Strep. pyogenes*. The results showed that a varied degrees of reduction in the absorbances were recorded for the contaminated milk samples pasteurized at 40°C, 50°C, 60°C and 70°C over a period of 30 min. With an increase in temperature of 10°C, there were significant reductions in the absorbances obtained at different time intervals for all the milk samples. The reductions in the absorbances of *E. coli* contaminated milk samples were higher than those obtained from IMF and *S. pyogenes* contaminated milk samples. Those of *E. coli* were followed by IMF contaminated milk samples while those of *S. pyogenes* contaminated milk samples were the least at the different

temperature used in pasteurization. In all these contaminated milk samples, the absorbances were reduced as time and temperature were increased. At 0 min when pasteurized at 40°C, TCM containing IMF, *E. coli* and *S. pyogenes* had an absorbances of 0.552 nm, 0.578 nm and 0.544 nm respectively. After 30 min of pasteurization, the absorbances were reduced to 0.532 nm, 0.554 nm and 0.472 nm respectively. At 0 min when pasteurized at 50°C, the absorbances were reduced to 0.491 nm, 0.490 nm and 0.426 nm from 0.534 nm, 0.559 nm and 0.472 nm in TCM contaminated with IMF, *E. coli* and *S. pyogenes* respectively. After 30 min of pasteurizing at 60°C, the absorbance was reduced from 0.491 nm, 0.490 nm, and 0.426 nm at 0 min to

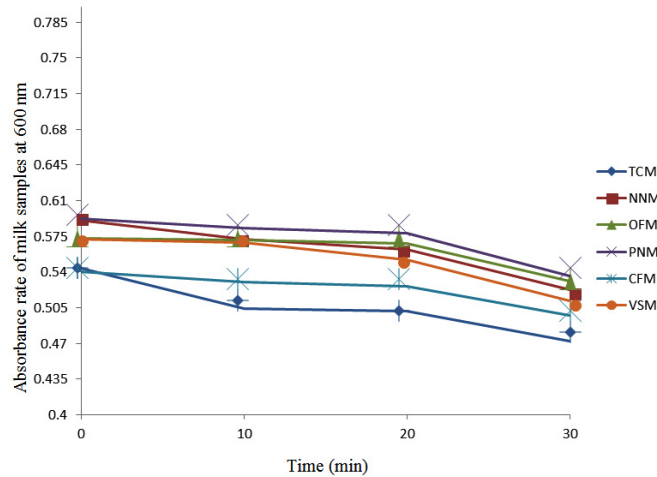


Fig. 3. Effect of pasteurization on milk samples contaminated with *S. pyogenes* at 40°C at 600 nm

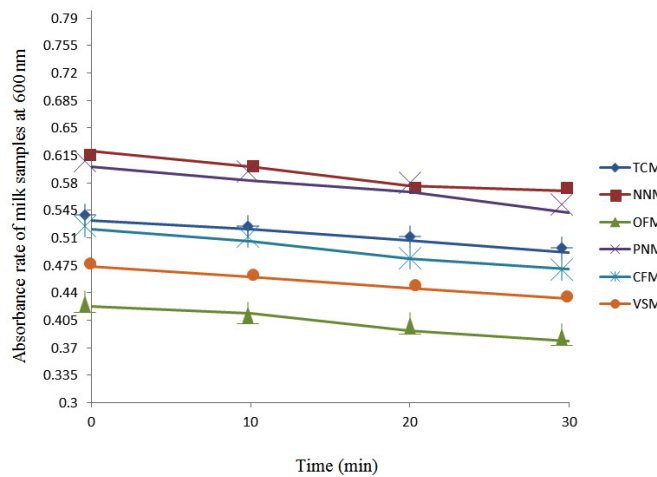


Fig. 4. Effect of pasteurization on milk samples contaminated with Indigenous Milk Flora (IMF) at 50°C at 600 nm

0.403 nm, 0.408 nm and 0.322 nm. After 30 min of pasteurizing at 70°C, the absorbances were reduced from 0.403 nm, 0.408 nm and 0.322 nm at 0 min to 0.262 nm, 0.283 nm and 0.221 nm. The same trend of reductions in the absorbances were obtained in the different milk samples contaminated with IMF, *E. coli* and *S. pyogenes* (Fig. 1 – 12). To ensure sterility, most of the contaminated milk subcultured on sterile nutrient showed no bacterial growth.

### DISCUSSION

Pasteurization is a moderate heat treatment of milk. It destroys disease producing organisms and limits fermentation in milk by heating milk to 63°C for 115 min while slow growing organisms and spore formers are hardly affected<sup>36</sup>. While pasteurization destroys many microorganisms in milk, improper handling after pasteurization could

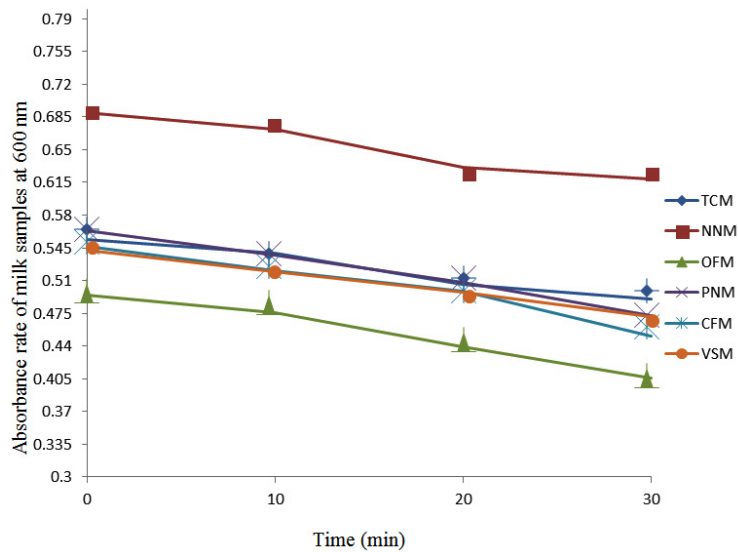


Fig. 5. Effect of pasteurization on milk samples contaminated with *E. coli* at 50°C at 600 nm

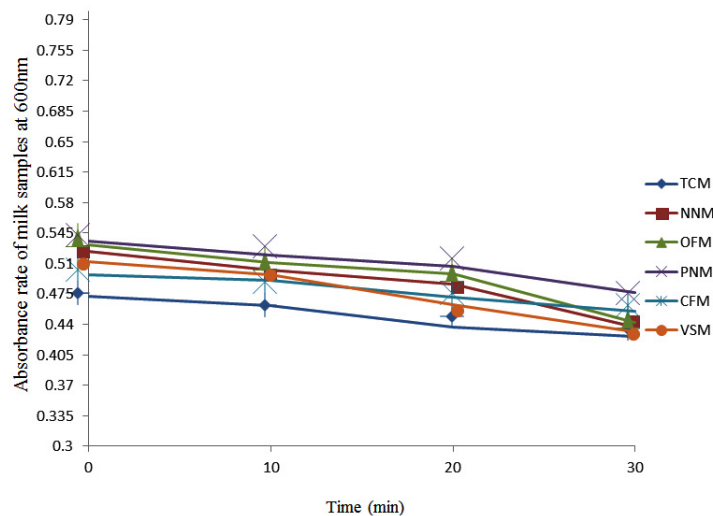


Fig. 6. Effect of pasteurization on milk samples contaminated with *S. pyogenes* at 50°C at 600 nm

lead to its recontamination. Thus, the isolation of different strains of bacteria from the milk samples used in this study could have been through recontamination from pasteurization equipment prior packaging. Although pasteurization is not sterilization and meant to reduce the pathogens, the negligible level of microbial loads ranging from  $5 \times 10^5$  to  $22 \times 10^5$  cfu/ml indicated the effectiveness of the pasteurization methods employed by the various manufacturing companies. However, since the microbial content of milk is a major feature in determining its quality<sup>37,38</sup>, the methylene reduction time indicating the hygienic quality of each milk sample showed that NNM and VSM samples were

excellent milk samples, PNM and OFM were considered good milk samples while CFM and TCM were considered fair milk samples.

However, since El-Mohandes *et al.*<sup>39</sup> reported that 30% of contaminated milk samples in their study had more than 30,000 cfu/ml in which the highest infection was related to *S. epidermidis*, the presence of microbial contaminant in the samples showed that they are potential source of microbial infections depending on the virulence, pathogenicity and inoculum sizes. Though the microbial loads were negligible, that milk is nutritious and could support microbial growth indicated that the bacterial load of the

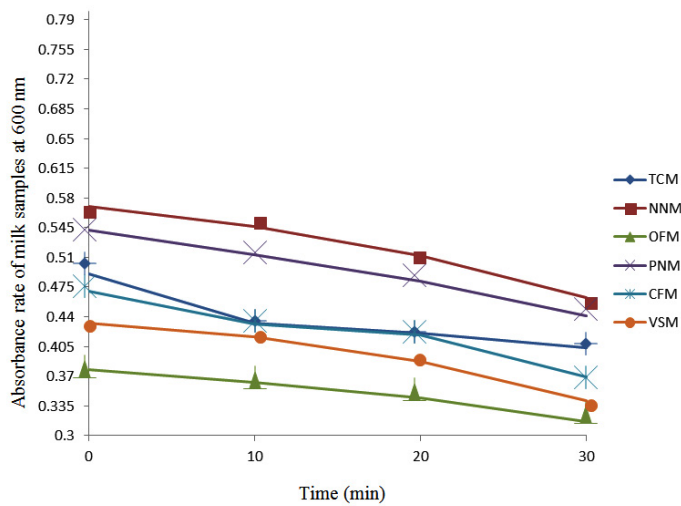


Fig. 7. Effect of pasteurization on milk samples contaminated with Indigenous Milk Flora (IMF) at 60°C at 600 nm

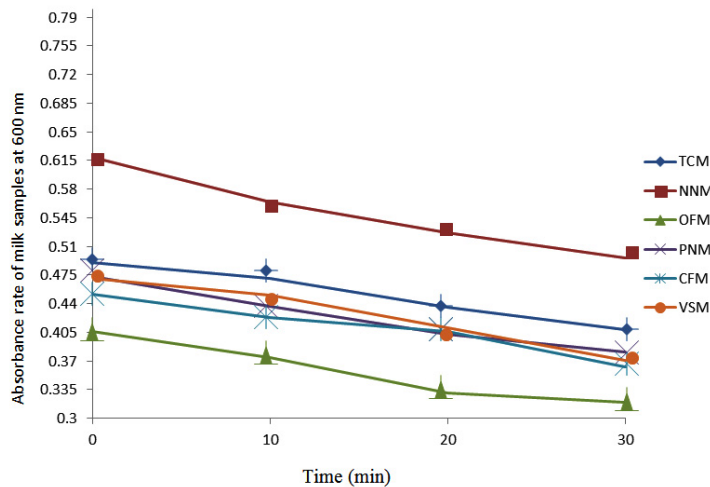


Fig. 8. Effect of pasteurization on milk samples contaminated with *E. coli* at 60°C at 600 nm

indigenous microbial flora could increase under favourable conditions. Since poor milk hygiene have implications on the structure of milk, processing value, shelf life and edible food loss<sup>40</sup> and indirectly on human health and bacterial contamination<sup>41-45</sup>, a high biological count in raw milk potent possible problems with product safety<sup>46</sup>.

In agreement with previous studies El-Mohandes *et al*<sup>37</sup>, Lindemann *et al*<sup>47</sup> and Rozolen *et al*<sup>48</sup>, the isolation of *Escherichia coli*, *Streptococcus spp.*, *Enterococcus aerogenes*,

*Proteus spp.*, *Clostridium spp.*, *Lactobacillus spp.* and *Staphylococcus aureus* from milk samples used have been reported. While El-Mohandes *et al*<sup>37</sup>, Lindemann *et al.*<sup>47</sup>, Rozolen *et al.*<sup>48</sup> and Carroll *et al.*<sup>49</sup> showed that frequently isolated microorganisms from milk samples included *S. epidermidis*, *S. aureus*, *Enterobacter*, Group B *Streptococcus*, *Escherichia coli* and *Klebsiella*, Karimi *et al.*<sup>50</sup> reported *Klebsiella*, *S. epidermidis*, *Enterobacter*, *Escherichia coli*, *Pseudomonas aeruginosa*, *Acinetobacter*, *S. saprophyticus*, *Citrobacter* and *S. aureus*. To forestall outbreaks

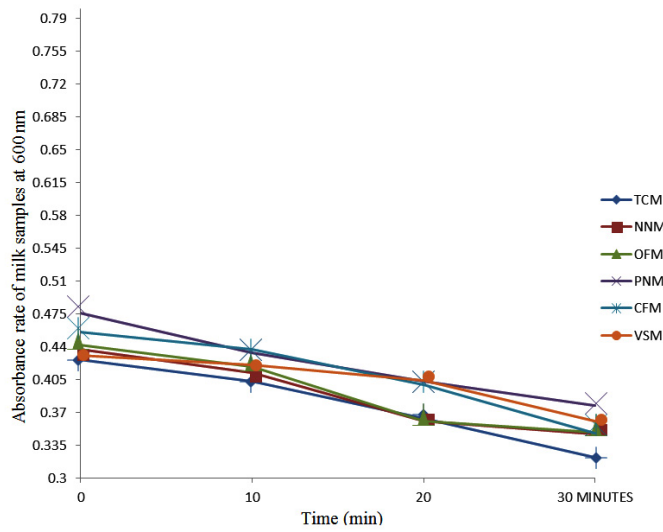


Fig. 9. Effect of pasteurization on milk samples contaminated with *S. pyogenes* at 60°C at 600 nm

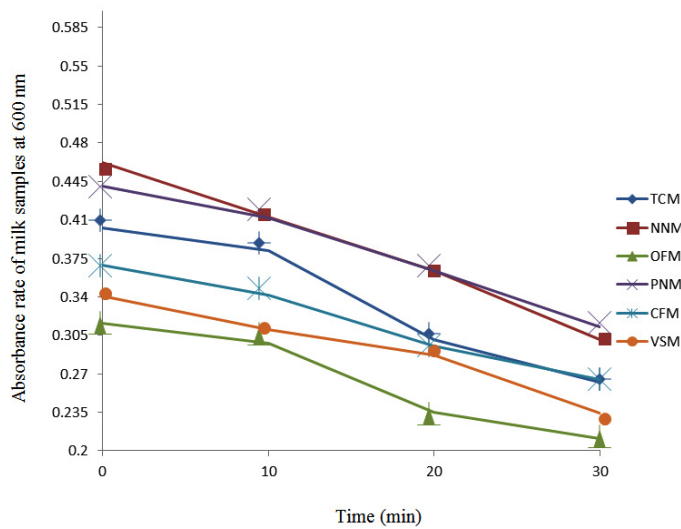


Fig. 10. Effect of pasteurization on milk samples contaminated with Indigenous Milk Flora (IMF) at 70°C at 600 nm



of infectious diseases from milk contaminated with these pathogens, several milk samples have been pasteurized at varied temperature to ascertain its effectiveness in decontaminating milk samples. Wallner *et al.*<sup>51</sup> showed that pasteurization is effective in destroying the bacteria in milk that cause tuberculosis, salmonellosis, diphtheria, typhoid fever and other illnesses without adversely affecting the milk's nutritional content, flavor or quality. Grant *et al.*<sup>52</sup> reported that milk contaminated with *Mycobacterium paratuberculosis* was

decontaminated with a high-temperature and short-time pasteurization. While Orloff *et al.*<sup>53</sup> reported that human immunodeficiency virus Type I was not recovered from human milk after pasteurization, Egberé *et al.*<sup>54</sup> pasteurized kunun-zaki at 70°C for 30 min and discovered that the bacterial load was drastically reduced from  $9.4 \times 10^5$  cfu/ml at 0 minutes to  $3.1 \times 10^1$  cfu/ml after 30 min. At 15 min, most of the *Streptococcus* spp present in the sample had been killed.

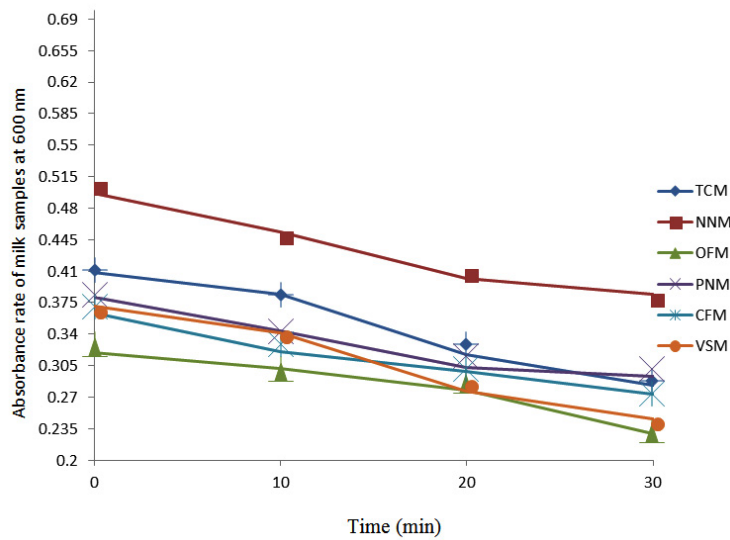


Fig. 11. Effect of pasteurization on milk samples contaminated with *E. coli* at 70°C at 600 nm

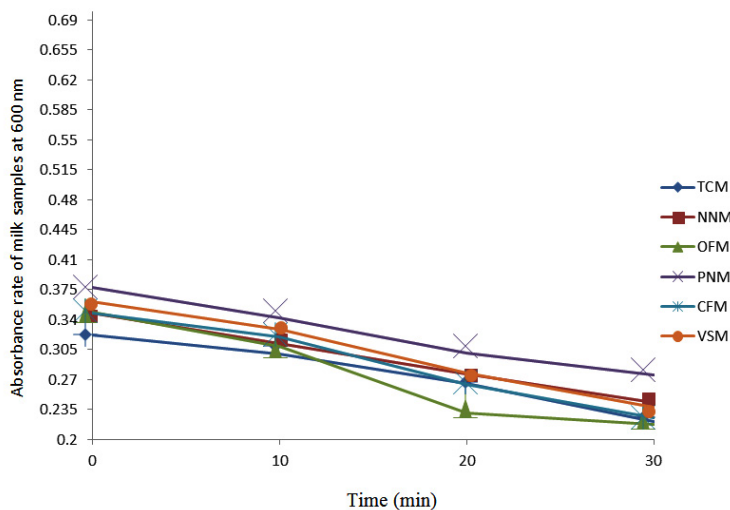


Fig. 12. Effect of pasteurization on milk samples contaminated with *S. pyogenes* at 70°C at 600 nm

From this study, the decreased absorbances of each milk sample contaminated with the indigenous milk flora (IMF), *E. coli* and *S. pyogenes* showed that the pasteurized milk contained very few organisms and agreed with the reports indicating that heating at 63°C for 30 min or 72°C for 15 min followed by quick cooling at 13°C destroys most milk-borne pathogens or reduces them to levels acceptable by international standards<sup>55-57</sup>. The spectrophotometric analysis of the pasteurized contaminated milk samples showed that all the samples obeyed Huygens' principle stating that light passing through a suspension of microorganisms is scattered and the amount of scattered light is an indication of the biomass present in the samples<sup>57</sup>. The absorbance rates of all tested pasteurized milk samples was reduced because the concentration and number of organisms present in all the milk liquids were reduced by the pasteurization.

In conclusion, milk is one of the most consumed foods in the world and contained virtually all nutrients needed by man. If properly pasteurized, it should not contain contaminants, pathogenic organisms or coliforms indicating fecal contamination. This study indicates that, although not all milk samples were made sterile by the pasteurization, pasteurization of milk would reduce the number and concentration of organisms present in milk to levels acceptable by international community as most organisms present are killed by heating at 70°C for 30 min. The method of pasteurizing milk in water bath at varied temperature as used in this study would be applicable in the homestead where sophisticated equipment are not required to control microbial loads in milk.

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