ORIGINAL ARTICLE



COD reduction of textile effluent in three-phase fluidized bed bioreactor using *Pseudomonas aureofaciens* and *Escherichia coli*

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Abstract This paper investigated the effect of mixed cultures of Pseudomonas aureofaciens and Escherichia coli and air as gas on degradation of chemical oxygen demand (COD) in three-phase inverse fluidized beds (TPIFBs) for textile effluents with initial COD of 1200 ppm (effluent 1) and 640 ppm (effluent 2). Investigators focused on the importance of different design of gas spargers of the TPIFB for checking its effect on COD reduction performance. For this purpose, four different gas spargers were fabricated for gas flow entry into bioreactor with different % open area: Gas sparger 1 (18% open area), sparger 2 (10.42% open area), sparger 3 (8.077% open area) and sparger 4 (1.53% open area). These percentage open area decided the amount of gas flow in a unit time in batch studies, which in turn rely mainly on superficial gas velocity. The reduction in COD was measured at two different superficial gas velocity (0.00343 and 0.004068 m/s) and at two ratio of static bed height to diameter (H/D) of 0.5 and 0.25 in an aerobic mode. Textile effluent 1 resulted in 98.07% COD reduction with sparger 3 and discharged at 23.14 ppm in 28 h. Textile effluent 2 resulted in 96.5% COD reduction with sparger 2 and discharged at 22.4 ppm in 22 h. The

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² Department of Biotechnology, V. V. P. Engineering College, Rajkot, Gujarat 360005, India resulting COD values of Effluent 1 and effluent 2 were in range of discharge limit and resulted at a low gas velocity of 0.00343 m/s and low H/D ratio of 0.25. The gas holdup correlation was found to be $\varepsilon_{\rm g} = 0.0064 \ U_{\rm g}^{0.98}$.

Keywords Aerobic treatment \cdot Batch process \cdot Gas sparger \cdot Three-phase inverse fluidized bed bioreactor \cdot Textile effluent

Introduction

Textile industries produce considerable amounts of effluent with high chemical oxygen demand (COD), which is contributed by recalcitrant organics, toxicants, color and salts (Doble and Kumar 2008). The presence of dyes in the effluent poses the biggest problem since they are recalcitrant and toxic. Two percent of dyes produced are discharged directly in the effluent (Doble and Kumar 2008). Alkali or acids from the processing steps of textile manufacturing units contribute to extreme pH and high salt content. The effluents from textile industries are generally variable in compositions which may not be biodegradable. Textile effluent is diverse because of processes employed and varieties of chemicals employed in each of the processes. It is difficult to assess the effluent characteristics for evaluating the requirements for suitable wastewater treatment. Therefore, it is necessary to understand process operation and character of individual effluents. Characteristics of textile wastewater are reviewed and that indicated the initial COD for effluent from different processes ranges from 800 to 30,000 mg/mL (Correia et al. 1994). Due to the presence of grease, dirt as well as nutrients from dye bath additives, the wastewater discharged from a dying process in textile industry is high in COD whereas the



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acceptable limits is 250 ppm (Singh 2000). COD is always indicated as pollution loads resulting from each processing operation of various raw materials. Thus, COD removal is needed with more effective treatment.

COD removals from textile effluent are accomplished by many researchers using various types of reactor systems or via different process strategies. Researchers have worked on Pseudomonas spp. (Vimala et al. 2015; Sriram et al. 2013; Blumel et al. 2002), Escherichia coli (Sriram et al. 2013; Vimala et al. 2015) and Bacillus spp. (Shah et al. 2013; Sriram et al. 2013; Blumel et al. 2002) for degradation of textile wastewater in an aerobic mode. Few researchers have worked on anaerobic reactor system (Shin et al. 2011) and found that a high influent dissolved oxygen (DO) concentration have adverse impacts on organic removal efficiency. Farabegoli et al. (2010) have worked with the mix of aerobic-anaerobic sequencing batch reactor system and deduced the fact that the influent COD/dye ratio to be properly selected to provide the electron equivalents needed for the oxidation of dye molecule. Different reactor systems are used to treat textile effluent with different characteristics: like, submerged filtration using nano-membrane (Zheng et al. 2013), Upflow anaerobic sludge bed reactor (UASB) coupled with adsorption process (Conceição et al. 2013), fluidized bed reactor (Su et al. 2011), microbial degradation (Agarry and Ajani 2011), combination of fixed and fluidized bed (Baban et al. 2010), thermolysis (Kumar et al. 2008), ultrafiltration (Dhodapkar et al. 2007) and sequential anaerobic-aerobic treatments (Kapdan et al. 2003)-to name a few. Different reaction criteria for COD removal have also been worked upon, like: ratio of volume of settled bed to bioreactor (Sokol 2003), use of catalyst system (Kumar et al. 2008) with thermolysis and coagulation treatment as well as solid content of influent (Rajasimman and Karthikeyan 2007). Biotreatment of textile wastewater required few potential indigenous microorganisms for biodegradation of COD (Agarry and Ajani 2011). The appropriate combination of physical or chemical methods or specific chemical and biological conditions are demanded for treating textile effluents (Agarry and Ajani 2011; Baban et al. 2010; Dhodapkar et al. 2007; Kapdan et al. 2003).

As discussed above, there are many ways to treat textile effluents. The fluidized bed bioreactor is one of the most effective methods used in effluent treatment (Buffiére and Moletta 1999). Recently, industrial wastewaters have been treated by aerobic biological treatment in three-phase fluidized bed (Haribabu and Sivasubramanian 2014; Sabarunisha Begum and Radha 2014; Sokol 2012; Sokol and Woldeyes 2011; Rajasimman and Karthikeyan 2007; Ochieng et al. 2003). Air stream is bubbled directly into the liquid, creating three phase in the aerobic treatment of wastewater. In a three phase fluidized bed bioreactor, if the



low density solid media particles are employed then the process is referred as inverse fluidization where gas phase is introduced counter currently to the liquid phase at the bottom of the reactor and the bed expands downwards (Haribabu and Sivasubramanian 2014; Sokol and Korpal 2006; Nikolov et al. 2000; Comte et al. 1997). The role of solid media particles is to provide large surface area for microbes where they are retained from the growth. There is a formation of biofilm onto which microorganisms retain to digest organic waste (Rajasimman and Karthikeyan 2007; Sokol 2012). Moreover, three-phase inverse fluidized bed bioreactor (TPIFB) offers large specific support surface area, higher gas phase holdup and higher residence time in comparison to conventional fluidized bed. Therefore, the mass transfer coefficient reaches almost two times higher in the TPIFB than conventional fluidized bed (Nikolov and Karamanev 1987). TPIFB is efficient when used for biological aerobic wastewater treatment. It is 15 times more effective than airlift reactor (Nikolov et al. 2000) and preferred over suspended-cell systems (Sokol 2012).

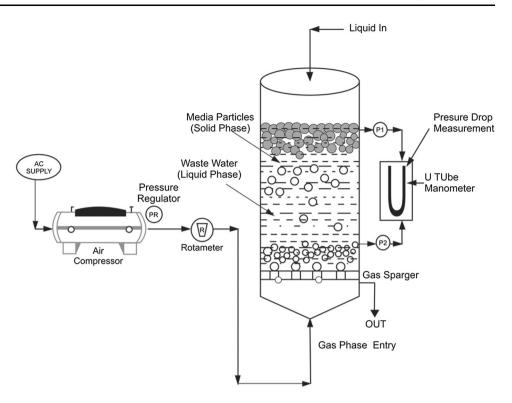
To investigate the effects of textile effluent characteristics on COD removal, two textile effluents with different COD values are taken and four gas spargers are designed. The purpose of the gas sparger of any fluidized bed is to evenly distribute the gas across the bed inlet (Sokol and Korpal 2006; Comte et al. 1997). Each of the gas spargers is characterized by open percentage area that it offers the air (gas phase) to pass through the wastewater (liquid phase). So each sparger controls the quantity of gas to flow at superficial velocity affecting the mass transfer.

Batch studies (only gas flow with no net liquid flow) for three-phase fluidized bed bioreactor in an inverse mode are countable and very few (Comte et al. 1997; Buffiére and Moletta 1999). The work on performance of three-phase inverse fluidized bed bioreactor (TPIFB) in a batch mode is still much far from complete. Therefore, the present work aims to investigate the performance of TPIFB through measurement of COD reduction in a batch mode with four spargers designed to treat two textile effluents. The effects of initial values of COD, effluent characteristics and effects of superficial gas velocities are discussed. Gas holdup correlation is derived.

Materials and methods

Air (density = 1.2 kg/m^3 , viscosity = 0.018 Pa s), effluent sample and solid media particles (polypropylene, 4 mm diameter, 723 kg/m³ density) comprised the three phases in the reactor column. The schematic diagram of three-phase inverse fluidized bed bioreactor for the batch process was given in Fig. 1. This apparatus was fabricated from acrylic column with a diameter of 0.097 m and height of 1 m. The

Fig. 1 Schematic diagram of fluidization in inverse mode



column consisted of liquid entry, sampling point and gas distribution. Synthetic wastewater was added from the top followed by addition of solid media particles. Air compressor was set to make air entry from the bottom. Air entered from the bottom of the reactor through air pressure regulator. Rotameter measured the flow (Online resources: ESM: Figure S1). Sufficient care was taken for the bioreactor operation and that, first the valve of compressor was opened and then valve at the bottom to maintain the required gas flow through the bioreactor. The solid media particles were added in quantities equivalent to the heights of the bioreactor column. This was calculated based on the ratio of static bed height to column diameter (H/D) of 0.5 and 0.25. Those values were equivalent to 4.85 and 2.43 cm heights of the particles (static bed height), respectively, in the bioreactor column.

Two raw untreated effluent samples were collected from industrial units in Rajkot district (Gujarat, India) with an initial COD of 1200 ppm (textile effluent 1) and 640 ppm (textile effluent 2). Compositions similar to these industrial effluents were fabricated in the laboratory (Online resources: ESM: Table S1). Major pollutants in textile effluents were characterized by the parameters like pH, COD, biological oxygen demand (BOD), hardness, NO₂⁻, and total suspended solids (TSS). The characteristics of textile effluents and the fabricated wastewaters were considered for experimental trials. The range of pH of the effluents were 7.15–8.5 and that of BOD was 107–182.4 (Online resources: ESM: Table S2). The reduction in COD was measured at static liquid conditions and low gas velocities. To facilitate the gas phase entry to the reactor, four different spargers were used. Each of them offered different area by their surface openings. Surface openings were referred as 'percentage openings' or 'open percentage area' in this paper. Percentage open area meant the proportion of open holes (m²) over the total surface of gas spargers. These were calculated to be 18% (sparger 1), 10.42% (sparger 2), 8.077% (sparger 3) and 1.53% (sparger 4). Each of the spargers was checked for its contribution in COD reduction since each of them allowed different volume in unit time to enter the bioreactor. Rotameter and pressure regulators were auxiliaries to this bioreactor column. The microbial cultures were added to the effluent for its degradation. Five different cultures of P. putida (NCIM 2872), P. aureofaciens (NCIM 2026), P. fluorescens (NCIM 2174), B. subtilis (NCIM 5433) and E. coli (NCIM 2805) were purchased from National Chemical Laboratory, Pune, India. The COD degradation capability of each of these cultures was determined. Finally, two microbial cultures of E. coli (NCIM 2805) and P. aureofaciens (NCIM 2026) were selected due to their higher degradation capabilities for further experimentation. These inoculated cultures were used for the entire bioreactor operation in a batch mode for three different sizes of inoculum-5, 10 and 20% of the working volume of the reactor. Batch process offered an advantage of sterility and easy handling. The bioreactor was ensured for sterility by two ways; firstly, by the use of air filter before the entry of air into the



bioreactor from compressor and secondly, by cleaning the media particles by acid solution (10% HCl) before adding into the bioreactor. Then the top of the reactor column was closed. The bioreactor was also washed with distilled water and 10% HCl solution after each batch.

The experiments were carried out at all possible combinations of parameters including gas velocities (0.006, 0.004068 and 0.00343 m/s), bed heights (0.3, 0.6 and 0.9 m) and the H/D of 0.5 and 0.25. When the bioreactor operation began, gas bubbles entered the bioreactor through gas sparger. These bubbles were moving in different fashions developing flow patterns including inversion of bed where particles moved downward. This inversion was captured in camera through a series of steps in sequence and presented here (Online resources: ESM: Figure S2). These process sequences occurred for all four gas spargers. The details of these gas spargers are given in Table 1 and Fig. 2.

Pressure taps were mounted at a distance of 0.05 m from the top, 0.05 m from the bottom and in the middle of the reactor column for the measurement of pressure through manometer. It was useful in maintaining steady state. At a time, bed of solid media became steady and expansion of bed remained constant. This referred the condition of steady state after which COD removal was observed in batch mode. Very low gas velocities were characteristics of this experimental setup.

Results and discussions

Textile effluent 1 and textile effluent 2 were treated with liquid at zero velocity and particles remained "fluidized" with only gas flow in TPIFB. Air was sparged from the bottom of the bioreactor. The bioreactor operated at the minimum fluidization velocity of 0.002141 m/s. The gas flow was increased slowly in the reactor to ensure the inversion whereby the solid media particles moved downwards. Uniform size bubbles mainly contributed the bioreactor operation. After a certain time, bed of solid media became steady with expansion of bed and then

remained constant. Initially all the experimentations were accomplished at gas velocities of 0.004068 and 0.00343 m/s and at the ratio of static bed height to diameter (H/D) of 0.5 and 0.25. All these experimental trials were taken for 0.3 m bed height (based on the fact that bed height did not affect the minimum fluidization velocity in the experimental condition).

After raising gas velocities beyond minimum fluidization, the COD measurements were taken at superficial gas velocities of 0.00343 and 0.004068 m/s with gas spargers 1, 2, 3 and 4. Bed was shown inversion at 0.00343 and 0.004068 m/s. There was no significant movement of particles required for the criteria of inversion when bed operated below the gas velocity of 0.00343 m/s and above 0.004068 m/s gas velocity, there was formation of slugs. Therefore, the readings were taken at these two velocities. The steady state was confirmed by noting the steady pressure drop. Time taken to reach steady state was about 14 min. At steady state, the readings were taken in batch mode at regular interval time of 2 h each. The refinements to the time interval were for 10 min.

Initial trials in the bioreactor were conducted using the microbial cultures of *P. putida* (NCIM 2872), *P. aureofaciens* (NCIM 2026), *P. fluorescens* (NCIM 2174), *B. subtilis* (NCIM 5433), and *E. coli* (NCIM 2805) to investigate their COD reduction capacity (Online resources: ESM: Table S3). The higher COD reduction capacity was exhibited by *P. aureofaciens* (NCIM 2026) and *E. coli* (NCIM 2805) and, therefore, they were selected for COD reduction across all the experimentations further.

Flow patterns in bioreactor operation

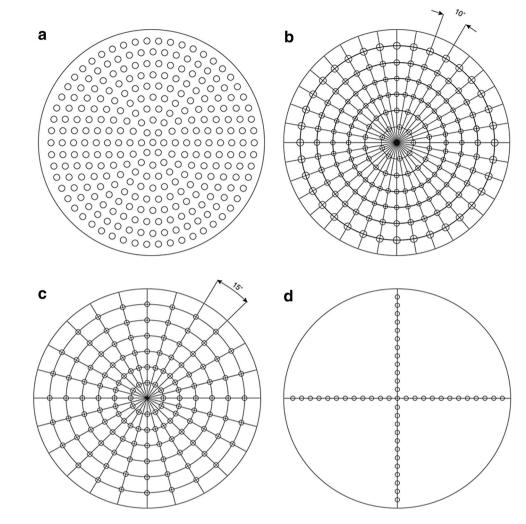
The bed was completely in packed condition initially. Gas was injected into the bioreactor apparatus with static liquid. Entire bed was fluidized and minimum fluidization velocity attained. Then superficial gas velocity was required to keep the particles in motion. The process resulted in homogeneous bed expansion. Bed had downward movements and it expanded up to the gas sparger. There was no movement past the gas sparger. At the onset of inverse fluidization

Table 1 Specifications for various gas spargers

Gas distributor	Description	Total Area (cm ²)	Sparger open area (cm ²)	Open area in (%)
Sparger 1	271 holes, 2.5 mm diameter each arranged in triangular pitch	73.86	13.279	18
Sparger 2	180 holes in total: 36 holes of 3 mm diameter, 36 holes of 2.5 mm diameter, 108 holes of 2 mm diameter, arranged at 10° angle	73.86	7.6986	10.42
Sparger 3	190 holes, 2 mm diameter each. Arranged at 15° angle	73.86	5.966	8.077
Sparger 4	39 holes, 2 mm diameter each at 90° angle	73.86	1.2246	1.53



Fig. 2 Designs of different gas spargers with specifications mentioned in Table 1



bubble started forming at superficial gas velocity and zero liquid velocity. The interaction between microbial cells and gas bubbles were governed by the flow patterns and it helped in understanding the extent of COD removal through all the four spargers. The flow patterns were influenced by sparger designs since each sparger allowed different volumes of gas into the bioreactor in a unit time. Initially, the flow represented dispersed regime with partially fluidized bed, for all the spargers. Homogeneous and consistent bubble flow was the main characteristics of sparger 2 and 3. However, there was a formation of slugs with the sparger 1 (higher open area) and sparger 4 (lower open area). These slugs were formed from homogeneous bubbly flow existing initially in the bioreactor. Later, the flow pattern indicated the bubbling fluidizing regime as the gas velocity was increased further. Bigger bubbles contributed in reduction of the surface area of interaction between microbes and bubbles. COD reduction dependence on open percentage area of the spargers, effluent characteristics, inoculum size and initial COD value were discussed further.

Effects of open percentage area

The characteristic design of each of the spargers offer different volumes of gas passing in TPIFB. Since the openings of gas spargers were different, they passed different volume of gas in the TPIFB. The effect of sparger openings were of interest from point of view of aeration. The aeration governed the growth of microorganisms. These microorganisms grew based on oxygen available at a time and degrade the organic pollutants as measured by COD. Their growth decided the reduction of COD for definite batch periods. Therefore, in turn, the gas sparger openings controlled the COD reduction rate. Four spargers were designed specifically with unique percentage open area (openings). The variations in designs of gas spargers were based on percentage open area (openings). So, the effects of these variations of designs of gas spargers were investigated in the performance of TPIFB for textile effluents of different nature.

Once the bioreactor was in steady operating conditions, bubbles formed in consistent patterns. Out of four designs

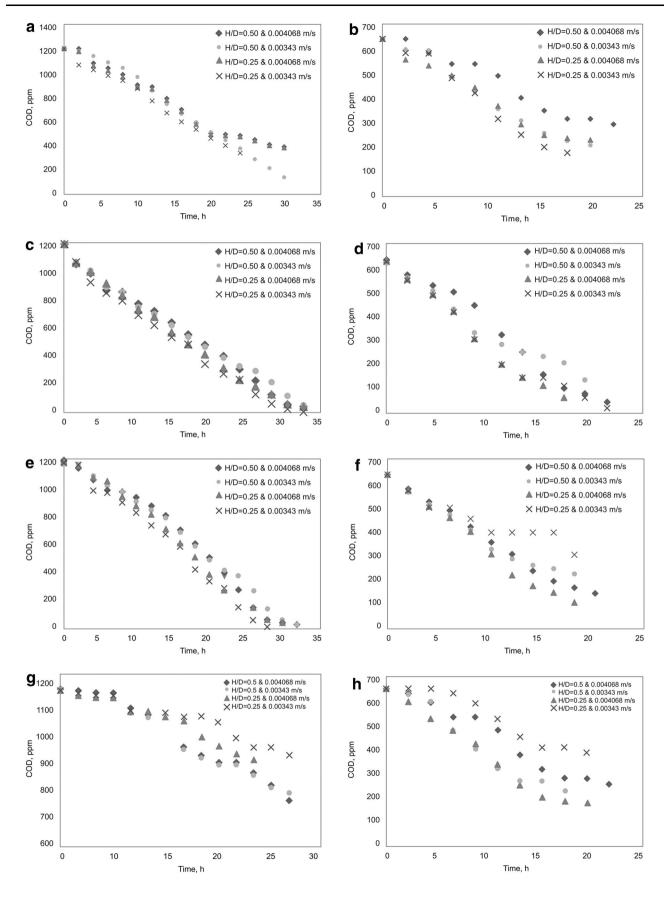


of spargers, Sparger 1 (Fig. 3a, b) was the one which had the highest opening area for gas to flow through the two phases- liquid effluent and solid media particles. Sparger 1 exhibited mainly bubble flow in case of textile effluent 1 and textile effluent 2. In the initial phase of the bioreactor operation, E. coli and P. aureofaciens mixed culture adopted to the chemical environment of the effluents in the period between 12 and 25 h for effluent 1 and between 8 and 17 h for effluent 2. These periods showed consistent growth of this mixed culture. The microorganisms retained on the surface of polypropylene particles to degrade textile effluent and were responsible for COD reduction. Turbidity was observed for these periods due to growth of E. coli and P. aureofaciens. Then after, there was no significant changes in COD reduction for effluent 1 and effluent 2, i.e., after 12 h for effluent 1 (Fig. 3a) and after 8 h for effluent 2 (Fig. 3b). Once this phase was over, gas holdup exhibited in the bioreactor column and created few slugs. Slug formation was not favorable for the growth of E. coli and P. aureofaciens. These slugs, as bigger irregular volumes of gas bubbles, were generated from smaller bubbles. Bigger bubbles contributed in reduction of the surface area of interaction between microbes and bubbles. This resulted in slower reduction of COD after 25 h for effluent 1 and after 17 h for effluent 2. Maximum removal efficiency achieved in sparger 1 was 88.3% at H/D of 0.5 with gas velocity of 0.00343 m/s for effluent 1. For effluent 2, maximum COD removal efficiency achieved was 81% at H/D of 0.25 and gas velocity of 0.00343 m/s with sparger 1. Both of these effluents required low gas velocities to achieve the maximum possible COD reduction.

Sparger 2 (Fig. 3c, d) had an open surface area of 10.42%. It exhibited consistent and homogeneous bubbly flow. However, the rate of degradation of COD in case of effluent 2 was little inconsistent between 8 and 16 h at high superficial gas velocity (0.004068 m/s) and H/D of 0.5. This occurred owing to the presence of steady gas bubbles in comparison to that for effluent 1 at 0.004068 m/s gas velocity and thus higher superficial gas velocity was not much desirable. Sparger 2 generated bigger gas bubbles at much faster rate compared to sparger 1. Hence the effective surface area for mass transfer was reduced for sparger 2. So, bigger bubbles were avoided by keeping superficial gas velocity as low as possible. The process of COD degradation was then dependent on both-gas velocity and sparger, apart from H/D ratio. Effluent 1 and effluent 2 resulted in 96.5% reduction in COD. There were consistent bubbles in both the spargers 1 and 2; however, the time taken to reach the value of maximum COD reduction was different since sparger 1 exhibited few slugs as mentioned earlier. This occurred at the gas velocity of 0.00343 m/s and 0.25 H/D for both the effluents. This strengthened the idea of COD degradation with low superficial gas velocity **Fig. 3** COD reduction in Sparger 1 (**a**, **b** for effluent 1, 2), Sparger 2 \triangleright (**c**, **d** for effluent 1, 2), Sparger 3 (**e**, **f** for effluent 1, 2), Sparger 4 (**g**, **h** for effluent 1, 2); at H/D ratio of 0.5 and 0.25 with gas velocities of 0.004068 and 0.00343 m/s

and low H/D ratio. Sparger 3 (Fig. 3e, f) was designed for open surface area of 8.077%. The experimental trails on sparger 3 resulted with same trend of flow pattern as that of sparger 2, but with different values of COD removal. Mass transfer occurred for these two spargers in consistent manner with discrete bubbles of air until full batch operation was over. The maximum possible COD reduction with the sparger 3 was 98.07% (at 0.00343 m/s and 0.25 H/D) for textile effluent 1. For effluent 2, it was 94.89%, but at higher gas velocity of 0.004068 m/s and higher H/D of 0.5.

On the other side, Sparger 4 (Fig. 3g, h) was found with the least COD degradation efficiency amongst all four spargers, since it had the least open percentage area which allowed lower quantity of air into the bioreactor column. So gas bubbles required comparatively more time to reach the bottom of the static bed in the bioreactor. Due to this reason, there was consistent mass from sparging point to the bottom of the bed. This sparger generated clear slugs. Slugs formed due to accumulation of bigger bubbles, which in turn resulted from smaller sized gas bubbles. Due to slugging, the mixed culture comprising E. coli and P. aureofaciens made poor contacts with gas phase which resulted in poor mass transfer. This never happened for other three spargers (sparger 1, 2, 3) discussed earlier. After taking three trials on each of these spargers, it was found that, open percentage area of each of the spargers was an important parameter in deciding reduction of COD of the effluents. Table 2 exhibits the comparison of parameters during experimental trials on textile effluents and fabricated wastewater. It indicated that in most of the cases, low gas velocity together with low H/D ratio was sufficient in reducing COD. In case of sparger 2 and 3, bubbly flow was only responsible for the degradation process and that for sparger 1 was bubbly flow and few slugs. However, sparger 4 was not performing well in comparison with other spargers. All the trials revealed that the overall process of COD degradation was comprised of following phases when bioreactor is in operation: (1) attainment and maintenance of superficial gas velocity (2) adoption of mixed culture of E. coli and P. aureofaciens to the bioreactor environment (3) depletion of pollutants from textile effluents through biodegradation by E. coli and P. aureofaciens. Performance of TPIFB for textile effluent is compared with other effluents for various reactor systems in Table 3 exhibiting the time taken for COD reduction in each case with their respective modes-the batch and the continuous mode.



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Sparger	Max C	OD red	uction (%)	Time t	aken (l	h)		Gas veloc	ity (m/s)			H/D :	ratio		
	E1	FE1	E2	FE2	E1	FE1	E2	FE2	E1	FE1	E2	FE2	E1	FE1	E2	FE2
1	88.30	98.16	81.00	93.00	30.00	8.00	16.00	34.00	0.00343	0.004068	0.00343	0.004068	0.5	0.25	0.25	0.5
2	96.50	99.50	96.50	96.95	28.70	6.80	22.00	32.00	0.00343	0.004068	0.00343	0.00343	0.25	0.5	0.25	0.25
3	98.07	98.50	94.89	96.70	28.00	6.80	20.00	34.00	0.00343	0.004068	0.004068	0.00343	0.25	0.5	0.5	0.25
4	39.93	96.16	72.12	92.20	27.80	7.00	18.00	38.00	0.004068	0.004068	0.00343	0.004068	0.5	0.25	0.5	0.5

Table 2 Comparison of parameters for trials on textile effluents and fabricated wastewater

El effluent 1, E2 effluent 2, FE l fabricated effluent 1, FE2 fabricated effluent 2

Effect of effluent characteristics

Textile effluent 1 and textile effluent 2 were experimented with different gas velocities, H/D ratio and open area of spargers. The effluent 1 resulted in 98.07% reduction with final value of COD as 23.18 ppm. The effluent 2 resulted in 96.5% reduction reaching at 22.4 ppm with sparger 2. Both the textile effluents met the discharge standards. The time taken for maximum COD reduction by effluent 1 and effluent 2 were different and the difference in the reduction of COD was comparatively low. It meant that the mixed culture of E. coli and P. aureofaciens species degraded COD of textile effluents taking different time based on their initial values. Therefore, initial value of these pollutants load (COD) also played a role in achieving final value. Reduction in these pollutants was the first strategy in prevention of pollution. High amount of TSS was the indication of high turbidity. High turbidity was observed in case of sparger 1 with assisted slugs and resulted in lowering COD. So, high turbidity was not preferred for better reduction in COD through growth of E. coli and P. aureofaciens. Other important parameters generally found in most of the textile effluents are NO₂⁻ and BOD. They did not influence the degradation in COD that occurred by the cultures of E. coli and P. aureofaciens in TPIFB. Characterization and subsequent treatment of textile effluents 1 and 2 using E. coli and P. aureofaciens enhanced the performance of the treatment of textile effluent 1 and textile effluent 2 in TPIFB.

Effect of inoculums size

The growth rate of bacteria controlled the oxygen demand and hence, affected the COD reduction. For initial trials, three different inoculum size (volume) were taken as 5, 10 and 20% of the wastewater for COD degradation (Online resources: ESM: Table S4a and S4b). For all the volumes of the inoculum, mostly lower gas velocity imparted higher COD reductions, using *P. aureofaciens* and *E. coli*, for both effluent 1 and effluent 2. The visual observation revealed dominance of slugs and more turbidity in the bioreactor when 20% inoculum size was used. Since the slugs contributed insufficient phase contacts between gas



and liquid phases, the COD reduction was lowered. This effect was enhanced by turbidity due to overgrown cultures with 20% inoculum size of *P. aureofaciens* and *E. coli* and these two conditions were undesirable. Trials with 5% inoculum volume size resulted with more COD reduction in comparisons to that of 20% inoculum volume. Very less amount of turbidity was observed at 5% inoculum size. This problem of turbidity and slugs was eliminated when trials were taken with 10% inoculum size. For both the effluent 1 and the effluent 2, the highest COD reduction was achieved with 10% inoculum volume. Flow patterns (as discussed in earlier section) were normal ensuring good mass transfer properties in the trials with 10% inoculum volume and hence this inoculums size was carried on for all the experimentations.

Effects of gas holdup

Gas hold up increased with increasing gas velocity at zero liquid velocity. The average value of gas holdup was measured based upon the increase in the height of the column by volumetric expansion method. It was calculated as a ratio of difference between final height and initial height to the final height of the bioreactor column. The average values of gas hold up were considered while developing the correlation. Gas holdup correlations from literature for different ranges of gas flow rates are compared in Table 4. The gas holdup values reported here were lower than reported by others. Chisti (1989) operated at much lower gas velocity and obtained n (0.80). At low gas flow rates, there existed almost a linear relation between gas holdup (ε_g) and gas velocity (U_g), as $\varepsilon_g = k U_g^n$, given in literature (Ochieng et al. 2003). The gas velocity (cm/s) was correlated with gas holdup to obtain n and k values. Fluidization was possible with lower gas flow rates in the inverse mode (lower density of solid media particles). The present work confirmed this and proposed the correlation for the gas holdup as below.

$$\varepsilon_{\rm g} = 0.0064 U_{\rm g}^{0.98}$$

0.00

Higher value of 'n' indicated that lower gas velocity was required for the bioreactor operation. The value of 'n' in

Effluent typeReactor systemMotimum (%) CODTime conductation (HTT or operation time) andProcess characteristicsTextile watewaterThree-phase inverse fluidized 8.07 $HTT = 17-34$ h, batch operation itme) andAreobic treatmentLow-strength syntheticThree-phase inverse fluidized bed reactor 9.07 $HTT = 0.27-1.33$ h, $3-46$ days of operation, semi-bed horsextorAreobic treatmentLow-strength syntheticAnerobic fluidized bed reactor 9.0 9.0 $HTT = 0.27-1.33$ h, $3-46$ days of operation, semi-bed horsextorAnerobic treatmentTextileSubmerged filtration $9.1.50$ $2.1.50$ $2.1.50$ days of operation, semi-bad horsextorAnerobic treatmentTextileSubmerged filtration $9.1.50$ $2.1.40$ h, batch nodePhysical-chemicalTextileHudszholed $9.1.50$ $2.1.40$ h, semi-continuous modePhysical-chemicalTextileHudszholed $9.0.50$ Reaction time = 10 m, hatch modePhysical-chemicalTextileHudszholed $7.0-93$ $2.1.40$ days of operation, unnotPhysical-chemicalTextileHudszholed $7.0-93$ $1.4.49$ s of operation, unnotPhysical-chemicalTextileHudszholed 9.91 $1.4.49$ s of operation, unnotPhysical-chemicalTextileUnnotine title (syc-but 3.90 $1.4.49$ s of operation, unnotPhysical-chemicalTextileUnnotine title (syc-but $1.4.49$ s of operation, unnotPhysical-chemicalTextileUnnotine title (syc-but $1.4.49$ s of operation, unnotPhysical							
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Low-strength syntheticAnaerobic Indidized bed reactor96HRT = 0.27-1.33 h, 3-46 days of operation, semit- continuous modeAnaerobic reamenti- anaerobic actionTextileSequencing batch reactor9724 byte/bit 2-97 days of operation, semit- continuous modeAnaerobic reactorTextileUASP reactor coupled with81.3024 byte/bit 2-97 days of operation, semit- continuous modePhysical-chemical Physical-chemicalTextileUASP reactor coupled with81.30Reaction time = 50 h, semi-continuous modePhysical-chemical Physical-chemicalTextileUASP reactor coupled with81.3087.449Single reaction time = 100 m, batch modePhysical-chemical 	1	Textile wastewater	Three-phase inverse fluidized bed bioreactor	98.07	HRT = $17-34$ h, batch operation	Aerobic treatment	Present work
TextileSequencing batch reactor9724 h cycle 12-97 days of operation, semi-Anaerobic-aerobicTextileSubmerged filtration1:00Reaction time = 50 h, semi-continuous modePhysical-chemicalTextileUASB reactor coupled with81.20HRT = 5.3 hPhysical-chemicalTextileUASB reactor coupled with81.20HRT = 5.3 hPhysical-chemicalTextileHudized bed reactor3.4.4950 days of operation, continuous modePhysical-chemicalTextileMicrobial degradation9714 days of operation, batch modeBiosegradationCotron rescribeCaalytic thermal treatment89.91Reaction time = 10. m, batch modeBiysical-chemicalTextileUtrafitation (UF) unit80-90Physical-chemicalTextileUtrafitation (UF) unit80-90Physical-chemicalTextileSequential treatment93.80 (optimal)HRT = 12-72 h, sludge age 10 daysArarobicRefinery wastewate	7	Low-strength synthetic wastewater	Anaerobic fluidized bed reactor (AFBR)	96	HRT = $0.27-1.33$ h, $3-46$ days of operation, continuous mode	Anaerobic treatment	Shin et al. (2011)
TextileSubmerged filtration91.50Reaction time = 50 h, semi-continuous modePhysical-chemicalTextileUASB reactor coupled with81.20HRT = 5.3 hPhysical-chemicalTextileUASB reactor coupled with81.20HRT = 5.3 hPhysical-chemicalTextileHudized bed reactor34.49Sof days of operation, batch modePhysical-chemicalTextileMicrobial degradation70-93HRT = 19.8-30.8 h, semi-continuous modePhysical-chemicalCotron textile (dye-bathFixed and fluidized bed70-93HRT = 19.8-30.8 h, semi-continuous modePhysical-chemicalCotron textile (dye-bathFixed and fluidized bed70-93HRT = 19.8-30.8 h, semi-continuous modePhysical-chemicalCotron =extileCatalytic thermal treatment89.91Reaction time = 4 h, 60 m regulation time forPhysical-chemicalTextileUltrafiltration (UF) unit80-90Physical-chemicalTextileSequential treatment90HRT = 12.72 h, sludge age 10 daysAnaerobic-AterobicRefinery wastewaterThree-phase fluidized Bed90HRT = 12.72 h, sludge age 10 daysAcrobic treatmentStarch wastewaterThree-phase fluidized bed bioreactor93.80 (optimal)HRT = 2.72 h, sludge age 10 daysAcrobic treatmentDomestic wastewaterInverse fluidized bed bioreactor93.80 (optimal)HRT = 2.75-80 h, continuous modeAcrobic treatmentDomestic wastewaterInverse fluidized bed biological90HRT = 2.75-80 h, continuous modeAcrobic treatment<	ŝ	Textile	Sequencing batch reactor	97	24 h cycle 12-97 days of operation, semi- continuous mode	Anaerobic-aerobic	Farabegoli et al. (2010)
TextileUASB reactor coupled with adsorption 81.20 HRT = 5.3 hPhysical-chemical adsorptionTextileHuidized bed reactor 34.49 Single reaction time = 100 m, batch modePhysical-chemical BiodegradationTextileMicrobial degradation 97 14 days of operation, batch modeBiodegradationTextileMicrobial degradation 97 14 days of operation, batch modeBiodegradationCotton =textileCatalytic thermal treatment 89.91 Reaction time = 4 h, 60 m regulation time forPhysical-chemical modeCotton =textileUltrafiltration (UF) unit 80.90 $ -$ Physical-chemical modeTextileSequential treatment 90 HRT = $12-72$ h, studge age 10 daysAmerobic-Aetrobic biologicalTextileSequential treatment 90 HRT = $12-72$ h, studge age 10 daysArerobic treatmentRefinery wastewaterThree-phase fluidized Bed 93.80 (optimal)HRT = $12-72$ h, studge age 10 daysArerobic treatmentStarch wastewaterInverse fluidized bed biofing 90 HRT = $12-72$ h, studge age 10 daysArerobic treatmentDomestic wastewaterInverse fluidized bed bioractor 93.80 (optimal)HRT = $12-72$ h, studge age 10 daysArerobic treatmentDomestic wastewaterInverse fluidized bed bioractor 93.80 (optimal)HRT = $12-72$ h, studge age 10 daysArerobic treatmentDomestic wastewaterInverse fluidized bed biofing 90 HRT = $12-72$ h, studge age 10 daysArerobic treatment	4	Textile	Submerged filtration	91.50	Reaction time $= 50$ h, semi-continuous mode	Physical-chemical	Zheng et al. (2013)
TextileHurdized bed reactor34-49Single reaction time = 100 m, batch modeChemicalTextileMicrobial degradation9714 days of operation, batch modeBiodegradationCotton textile (dye-bath)Fixed and fluidized bed70-93HRT = 19.8-30.8 h, semi-continuous modeBiodegradationCotton textileCatalytic thermal treatment89-91Reaction time = 4 h, 60 m regulation time forPhysical-chemicalCotton =textileCutalytic thermal treatment80-90Physical-chemicalTextileUltrafiltration (UF) unit80-90Physical-chemicalTextileSequential treatment90HRT = 12-72 h, sludge age 10 daysPhysical-chemicalTextileSequential treatment90HRT = 58 h, batch culturing 2 weeksAnterbio:-ArtobicRefinety wastewaterThree-phase fluidized Bed90HRT = 58 h, batch culturing 2 weeksAerobic treatmentBioreactor95.60 (maximum)HRT = 58 h, batch culturing 2 weeksAerobic treatmentBioreactor95.60 (maximum)HRT = 67 h, batch nulturing 2 weeksAerobic treatmentIndustrial wastewaterInverse fluidized bed biofin95.60 (maximum)HRT = 6.40 h, batch nulturing 2 weeksAerobic treatmentIndustrial wastewaterInverse fluidized bed biofin95.60 (maximum)HRT = 6.40 h, batch nuodeAerobic treatmentIndustrial wastewaterInverse fluidized bed biofin96HRT = 6.7, continuous modeAerobic treatmentIndustrial wastewaterInverse fluidized bed	S	Textile	UASB reactor coupled with adsorption	81.20	HRT = 5.3 h	Physical-chemical	Conceição et al. (2013)
TextileFundized bed reactor34-49Single reaction tune = 100 m, batch modeChemicalTextileMicrobial degradation9714 days of operation, batch modeBiodegradationCotton textileAnalytic thermal treatment89-91Reaction time = 4 h, 60 m regulation time forPhysical-chemicalCotton =textileCatalytic thermal treatment89-91Reaction time = 4 h, 60 m regulation time forPhysical-chemicalTextileUltrafiltration (UF) unit80-90-Physical-chemicalbiologicalTextileSequential treatment90HRT = 12-72 h, sludge age 10 daysAnaerobic-AerobicRefinery wastewaterThree-phase fluidized Bed93.80 (optimal)HRT = 12-72 h, sludge age 10 daysAnaerobic-AerobicStarch wastewaterThree-phase fluidized bed93.80 (optimal)HRT = 54 h (optimal)Aerobic treatmentIndustrial wastewaterInverse fluidized bed bioreactor95.60 (maximun)HRT = 6-40 h, batch modeAerobic treatmentIndustrial wastewaterInverse fluidized bed bioreactor97.50HRT = 65 h, continuous modeAerobic treatmentIndustrial wastewaterInverse fluidized bed biological96HRT = 65 h, continuous modeAerobic treatmentIndustrial wastewaterInverse fluidized bed biological96HRT = 65 h, continuous modeAerobic treatmentIndustrial wastewaterInverse fluidized bed biological96HRT = 65 h, continuous modeAerobic treatmentIndustrial wastewaterInverse fluidized bed biological96HRT = 65 h	,	:			200 days of operation, continuous mous		
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Cotton textile (dy-bath)Fixed and fluidized bed70-93HRT = 19.8-30.8 h, semi-continuous modePhysical-chemicaleffluent)Cotton =textileCatalytic thermal treatment89.91Reaction time = 4 h, 60 m regulation time forPhysical-chemicalTextileCultrafiltration (UF) unit80-90Physico-chemical andTextileSequential treatment90HRT = 12-72 h, sludge age 10 daysPhysico-chemical andTextileSequential treatment90HRT = 24 h (optimal)Aerobic treatmentStarch wastewaterThree-phase fluidized bed93.80 (optimal)HRT = 24 h (optimal)Aerobic treatmentDomestic wastewaterInverse fluidized bed biofilm95.60 (maximum)HRT = 40 h (max), semi-continuous modeAerobic treatmentDomestic wastewaterInverse fluidized bed biofilm95.60 (maximum)HRT = 65 h, continuous modeAerobic treatmentIndustrial wastewaterInverse fluidized bed biofilm96HRT = 65 h, continuous modeAerobic treatmentIndustrial wastewaterInverse fluidized bed biofilm96HRT = 27.3-80 h, continuous modeAerobic treatmentIndustrial wastewaterInverse fluidized bed biofilm96HRT = 27.3-80 h, continuous modeAerobic treatmentRefinery wastewaterInverse fluidized bed biofilm96HRT = 27.3-80 h, continuous modeAerobic treatmentRefinery wastewaterInverse fluidized bed biological96HRT = 27.3-80 h, continuous modeAerobic treatmentRefinery wastewaterInverse fluidized bed biologic	٢	Textile	Microbial degradation	97	14 days of operation, batch mode	Biodegradation	Agarry and Ajani (2011)
Cotton =textileCatalytic thermal treatment8.9.1Reaction time = 4 h, 60 m regulation time forPhysical-chemical reactionTextileUltrafiltration (UF) unit80-90-Physico-themical reactionPhysico-themical biologicalTextileSequential treatment90HRT = 12-72 h, sludge age 10 daysAnaerobic-Aerobic 	8	Cotton textile (dye-bath effluent)		70–93	HRT = 19.8-30.8 h, semi-continuous mode	Physical-chemical	Baban et al. (2010)
TextileUltrafiltration (UF) unit80-90-Physico-chemical and biologicalTextileSequential treatment90HRT = 12-72 h, sludge age 10 daysPhysico-chemical and biologicalRefinery wastewaterThree-phase fludized Bed90HRT = 58 h, batch culturing 2 weeksAnaerobic-AerobicRefinery wastewaterThree-phase fludized bed93.80 (optimal)HRT = 58 h, batch culturing 2 weeksAerobic treatmentStarch wastewaterThree-phase fludized bed93.80 (optimal)HRT = 54 h (optimal)Aerobic treatmentDomestic wastewaterInverse fluidized bed bioreactor95.60 (maximum)HRT = 40 h (max.), semi-continuous modeAerobic treatmentDomestic wastewaterInverse fluidized bed bioreactor97.50HRT = 6-40 h, batch modeAerobic treatmentIndustrial wastewaterInverse fluidized bed biological96HRT = 65 h, continuous modeAerobic treatmentRefinery wastewaterInverse fluidized bed biological96HRT = 27.5-80 h, continuous modeAerobic treatmentRefinery wastewaterInverse fluidized bed biological96HRT = 27.5-80 h, continuous modeAerobic treatmentRefinery wastewaterInverse fluidized bed biological96HRT = 27.5-80 h, continuous modeAerobic treatment	6	Cotton =textile	Catalytic thermal treatment	89.91	Reaction time = $4 h$, $60 m$ regulation time for reaction	Physical–chemical	Kumar et al. (2008)
TextileSequential treatment90HRT = 12–72 h, sludge age 10 daysAnaerobic-AerobicKiRefinery wastewaterThree-phase fluidized Bed90HRT = 58 h, batch culturing 2 weeksAerobic treatmentSoStarch wastewaterThree-phase fluidized bed93.80 (optimal)HRT = 24 h (optimal)Aerobic treatmentReStarch wastewaterThree-phase fluidized bed93.80 (optimal)HRT = 24 h (optimal)Aerobic treatmentReDomestic wastewaterThree-phase fluidized bed bioreactor95.60 (maximum)HRT = 40 h, batch modeAerobic treatmentReDomestic wastewaterInverse fluidized bed bioreactor97.50HRT = 6-40 h, batch modeAerobic treatmentReIndustrial wastewaterInverse fluidized bed biological96HRT = 65 h, continuous modeAerobic treatmentSoRefinery wastewaterInverse fluidized bed biological96HRT = 27.5-80 h, continuous modeAerobic treatmentSoRefinery wastewaterInverse fluidized bed biological96HRT = 27.5-80 h, continuous modeAerobic treatmentSo	10	Textile	Ultrafiltration (UF) unit	80–90	1	Physico-chemical and biological	Dhodapkar et al. (2007)
Refinery wastewaterThree-phase fluidized Bed90HRT = 58 h, batch culturing 2 weeksAerobic treatmentSoStarch wastewaterThree-phase fluidized bed93.80 (optimal)HRT = 24 h (optimal)Aerobic treatmentRacobic treatmentStarch wastewaterThree-phase fluidized bed93.80 (optimal)HRT = 24 h (optimal)Aerobic treatmentRacobic treatmentDomestic wastewaterInverse fluidized bed bioreactor95.60 (maximum)HRT = 40 h (max.), semi-continuous modeAerobic treatmentRacobic treatmentIndustrial wastewaterInverse fluidized bed bioreactor97.50HRT = 6-40 h, batch modeAerobic treatmentMacobic treatmentIndustrial wastewaterInverse fluidized bed biological96HRT = 67.5-80 h, continuous modeAerobic treatmentSoRefinery wastewaterInverse fluidized bed biological96HRT = 27.5-80 h, continuous modeAerobic treatmentSoreactorreactorreactor16HRT = 27.5-80 h, continuous modeAerobic treatmentSo	11	Textile	Sequential treatment	06	HRT = 12-72 h, sludge age 10 days	Anaerobic-Aerobic	Kapdan et al. (2003)
Starch wastewater Three-phase fluidized bed 93.80 (optimal) HRT = 24 h (optimal) Aerobic treatment Racobic treatment So Industrial wastewater Inverse fluidized bed biological 96 HRT = 67.5-80 h, continuous mode Aerobic treatment So Refinery wastewater Inverse fluidized bed biological 96 HRT = 27.5-80 h, continuous mode Aerobic treatment So reactor reactor 16 HRT = 27.5-80 h, continuous mode Aerobic treatment So	12	Refinery wastewater	Three-phase fluidized Bed bioreactor	06	HRT = 58 h, batch culturing 2 weeks	Aerobic treatment	Sokol (2003)
Domestic wastewaterInverse fluidized bed bioreactor 97.50HRT = 6-40 h, batch modeAerobic treatmentHaIndustrial wastewaterInverse fluidized bed biofilm96HRT = 65 h, continuous modeAerobic treatmentSoRefinery wastewaterInverse fluidized bed biological96HRT = 27.5-80 h, continuous modeAerobic treatmentSoRefinery wastewaterInverse fluidized bed biological96HRT = 27.5-80 h, continuous modeAerobic treatmentSo	13	Starch wastewater	Three-phase fluidized bed bioreactor	93.80 (optimal) 95.60 (maximum)	HRT = 24 h (optimal) HRT = 40 h (max.), semi-continuous mode	Aerobic treatment	Rajasimman and Karthikeyan (2007)
Industrial wastewaterInverse fluidized bed biofilm96HRT = 65 h, continuous modeAerobic treatmentSoreactorreactorRefinery wastewaterInverse fluidized bed biological96HRT = 27.5–80 h, continuous modeAerobic treatmentSoRefinery wastewaterInverse fluidized bed biological96HRT = 27.5–80 h, continuous modeAerobic treatmentSo	14	Domestic wastewater	Inverse fluidized bed bioreactor		HRT = $6-40$ h, batch mode	Aerobic treatment	Haribabu and Sivasubramanian (2014)
Refinery wastewater Inverse fluidized bed biological 96 HRT = 27.5–80 h, continuous mode Aerobic treatment So reactor	15	Industrial wastewater	Inverse fluidized bed biofilm reactor	96	HRT = 65 h, continuous mode	Aerobic treatment	Sokol (2012)
	16	Refinery wastewater	Inverse fluidized bed biological reactor	96	HRT = 27.5 -80 h, continuous mode	Aerobic treatment	Sokol and Woldeyes (2011)

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S. no.	$U_{\rm g}~({\rm cm/s})$	\mathcal{E}_{g}	Correlation	References
1	0.34–0.41	0.0078-0.0109	$\varepsilon_{\rm g} = 0.00639 \ U_{\rm g}^{0.995}$	Present work
2	1-8	0.05-0.3	$\varepsilon_{\rm g} = 0.71 \ U_{\rm g}^{0.535}$	Ochieng et al. (2003)
3	0.1-1.8	0.003-0.045	$\varepsilon_{\rm g} = 1.45 \ U_{\rm g}^{0.8}$	Chisti (1989)
4	20-800	0.2–0.9	$\varepsilon_{ m g}=0.417~U_{ m g}^{0.44}$	Kito et al. (1976)

the present work was 0.98. It indicated strict bubble regime.

mimicked the actual textile effluent as far as the nutrient contents were concerned.

Effect of initial COD values

Mixed culture comprising *E. coli* and *P. aureofaciens* consumed more time in adapting to the chemical environment for textile effluent 1 (higher initial values of COD). However, with lower initial values of COD as in textile effluent 2, reduction of COD was achieved in comparatively shorter time.

Higher initial COD indicated higher organics in it. Higher COD required 28 h to degrade the organics content present for textile effluent 1 with sparger 3 whereas textile effluent 2 degraded maximum possible organics in 22 h with sparger 2 (both effluents exhibited maximum COD reduction at H/D ratio of 0.25 and 0.00343 m/s gas velocity). Adaptability of E. coli and P. aureofaciens was different for both the effluents. E. coli and P. aureofaciens had to adjust to the harsher environment of textile effluent 1 taking more 6 h to that in comparison with textile effluent 2 for removing maximum COD. However, more the value of initial COD of textile effluent; more duration required for degrading the COD. Organics of effluent 1 degraded over larger duration. Exactly opposite happened for textile effluent 2. Stoichiometry of available and required oxygen via aerobic digestion mode governed COD reduction. The supply of air was constant for textile effluent 1 and textile effluent 2 during the bioreactor operation. Flow regimes remained the similar over a period of degradation for textile effluent 1 and textile effluent 2 and nature of degradation was identical in regard to the growth pattern of E. coli as well as P. aureofaciens.

Both of these effluents comprised various nutrients onto which the growth of *E. coli.* and *P. aureofaciens* were dependent. Their growth decided the capacity to degrade the COD contents. Dextrose was the main nutrient for the growth of *E. coli* as well as *P. aureofaciens*. Urea and yeast extract served as nitrogen supplier to these microbial cultures. The composition of fabricated wastewater had typical pH content between 7 and 9 which was optimal for the growth of *E. coli* and *P. aureofaciens*. This composition

Conclusions

Textile effluents 1 and 2 reduce COD that it remains within permissible limit (<250 ppm) after treatment. Effluent 1 reduces in COD value from 1200 ppm to 23.14 ppm (98.07% reduction) with sparger 3 in 28 h. Effluent 2 degrades COD from initial 640 ppm to 22.4 ppm (96.5% reduction) with sparger 2 in 22 h. Gas spargers play a vital role in COD reduction process. Both samples accomplish maximum COD reduction at low gas velocity of 0.00343 m/s and low H/D ratio of 0.25. TPIFB is an effective bioreactor in treatment of textile effluent.

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Compliance with ethical standards

Conflict of interest This manuscript is the authors' original work and has not been published nor has it been submitted simultaneously elsewhere. All authors have checked the manuscript and have agreed to the submission.

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