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Investigating the Functional and Structural Adaptation Changes of Biofilm Communities Toward Better Azo-dye Wastewater Treatment

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Authors' contributions

This work was carried out in collaboration between both authors. Author MB provided valuable suggestions for the study design, ran the model, performed the data analyses and wrote the manuscript. Author YT designed the study, supervised the model runs and the data analyses and reviewed the manuscript. Both authors read and approved the final manuscript.

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ABSTRACT

Aims: The aim of this study was to investigate the functional and structural adaptation changes in three biofilm communities purifying azo-dye contaminated wastewater.

Study Design: Three lab-scale sand biofilters were constructed for treating wastewater with azodye. The dye amaranth was chosen as model toxicant and its concentration was gradually increased in the wastewater from 10 mg/L up to 55 mg/L. The biofilters functioned for 26 days.

Place and Duration of Study: Laboratory of Environmental biotechnology, Sofia University "St. Kliment Ohridski", 2011-2013 year.

Methodology: The residual amaranth concentration, efficiency and rate of its removal were monitored. The diagnostics of the functional structure of the biofilms is based on a cross analysis of data from culturable, culture-independent (fluorescent *in-situ* hybridization - FISH) and digital (digital image processing) techniques.

Results: The efficiency of the biofilters varied from 88.35% up to 95.97%. Three phases of the azo-detoxification process were distinguished.

In the early phase of functioning (0-191 h) culturable *Pseudomonas sp.* had key role in azodegradation as their part of the community was about 70% for the three biofilters.

In the late phase of functioning (191-455 h) the biofilters eliminated 2 times higher concentration of amaranth. The mean value for the part of the microorganisms from g. *Pseudomonas*, calculated on the base of FISH, remained unchanged (42%). Simultaneously the cultivation techniques showed

significantly decreased part of *Pseudomonas sp.* (4-10%). This suggest an important role of the unculturable *Pseudomonas sp.*

In the ending period (455-623 h) the rate of amaranth removal was increased with 20%. Wellformed zones with high concentration of bacteria from g. *Pseudomonas* were found in the biofilms which indicated formation of cooperative relationships.

Conclusion: This study shows three-stage mechanism of development of amaranth degradation potential. It includes successive importance of culurable, unculturable bacteria and cooperation in the g. *Pseudomonas*.

Keywords: Biofilm; FISH; digital analysis; azo-dye; unculturable; cooperative relationships.

1. INTRODUCTION

The hydrosphere, lithosphere, atmosphere and biosphere are the essential parts of our planet. They are our environment and their protection from pollution and deterioration is the main global problem and responsibility of the human society, science and technologies. In 21 century the humankind is facing many ecological problems related with the depletion of resources and pollution of the environment. Although the wastewater treatment technologies aroused in order to preserve the health of the people from the big cities, the goals of the contemporary biotechnologies are much broader and reached far beyond the human's health. They are also a solution for many problems concerning the "health" of ecosystems, water purification and minimization of the human impact.

The protection of water against pollution more and more are being recognized as main topics in the context of climate change. On the 21 Conference of the Parties (COP21), taking place in 2016 in Paris, new water related initiative "Climate is water" was announced [1]. With this the international community officially had recognized the sustainable water use as one of the most important factors affecting climate change. Most of the considered measures are related with water reuse, development of more efficient and sustainable use of water resources by better controlling the demand and increasing water availability [2]. In the same time among the most vulnerable industries is the textile processing industry because the use of large amounts of water and recalcitrant compounds. They are highly colored and difficult to eliminate in the same time [3,4]. Because of this special attention should be paid on the efficient treatment of dye-contaminated wastewater.

The annual production of the synthetic dyes is estimated at 280 000 t [3]. One of the most widely used synthetic dyes are the azo-dyes whose use globally increased from 28 000 t in 1992 year [5] up to 53 500 t in 2002 year. These are the second most used synthetic dyes for 2002 year [6].

In the textile industry the azo-dyes are the most widely used types of dyes [4]. One of the characteristics of the textile coloring technologies is that much of the dye can't be absorbed from the fibers (between 2% and 50%). As result the wastewater becomes highly colored [7]. Some of the azo-dyes are visible in concentrations below 1 mg/L. Often the wastewater from textile processing contains dyes in concentration between 10-200 mg/L which makes them highly colored [7]. When such effluent is discharged in the surface waters, not only aesthetical problems occur but also the light penetration is reduced. This inhibits the photosynthesis and decreases the oxygen concentration in the receiving water body. The whole ecosystem becomes negatively affected [4,8,9,10]. Also because of their toxicity and carcinogenic properties many of the azodyes are hazardous for the human health [4,11,12,13,14,15,16].

Because of their adaptation potential and enzymatic diversity, the bacteria are most often used as azo-dyes biodegrading organisms. Some of the genera with high azo-degradation capacity are Shewanella, Proteus, Bacillus, Klebsiella, Micrococcus and Pseudomonas [17,18,19,20,21,22,23,24,25]. Single species of microorganisms may have a high decolorizing activity, and are suitable for studying the mechanisms of azo-dyes biodegradation but in practice wastewater treatment with singlespecies technology is impossible [17,26,27,28, 29,30,31,32]. The communities containing diverse microorganisms have advantages related with cooperation in pollutants biodegradation, availability of wide variety of enzymes ensuring complete mineralization [33], development of synergistic and co-metabolic relationships between biodegraders. For example one species can reduce the azo-bond and another species can metabolize the resulted aromatic amines [32]. Another important advantage of the multispecies communities is that the biodegradation processes are carried out with higher rate and efficiency (because of the established microbial relationships) [34,35,36,37].

In lab-scale processes the microorganisms are able to eliminate azo-dyes with concentrations about 50-200 mg/L for period of time from 2 h. up to 48 h. [38]. In the real practice the most often used technologies for treating textile wastewater are the classical bioreactor with activated sludge or SBR bioreactors [39]. Yet, in the scientific field, many technological designs have been proposed in order to achieve more efficient dye treatment. For example biological aerated filters [40,41]. up-flow anaerobic sludge blanket (UASB) bioreactors [42,43], membrane bioreactors [44,45,46]. immobilized microorganisms [47,48], combination of biological treatment and nanofiltration or adsorption [49,50,51,52], constructed wetlands [53,54].

Generally the scientific investigations are working concentrated in finding effective biodegraders or bioreactors for efficient purification of the textile wastewater. Outside of the focus remains the adaptation of microbial biofilm communities as a tool for optimal azo-dye wastewater treatment. That is why in the center of this study is put on the development of azodegradation capacity of three microbial biofilm communities. The adaptation of the communities is carried out under conditions of stepwise increase of the azo-dye concentration. The key technological and microbiological parameters are monitored. Special attention is paid on the data obtained with two non-culturable methods - FISH (fluorescent in-situ hybridization), combined with digital image processing.

2. MATERIALS AND METHODS

2.1 Design of the Experiment

Three model sand biofilters with volume 171 cm³ and depth of 3 cm functioned in the Laboratory of Environmental biotechnology in Sofia University for 26 days. The biofilters treated wastewater (520-790 mL/day) polluted with toxicant - the azo-dye amaranth. The concentration of the xenobiotic was stepwise increased from 10 mg/L up to 55 mg/L. The technological parameters

(residual amaranth concentration, decolorization efficiency, rate of amaranth removal) were monitored 2-3 times per day. Microbiological, FISH and digital analysis were made in three key points in the experiment: 191 h (early phase of functioning), 455 h (late phase of functioning), 623 h (end of the experiment).

2.2 Biological System

The biological systems are biofilms formed on the sand carrier. Activated sludge (AS) from WWTP (wastewater treatment plant) of Sofia city was used as inoculation material. Before immobilization AS was disintegrated by an ultrasonic disintegrator (3×10 s). In this way the floculas in AS were destroyed and homogenous microbial suspension, suitable for immobilization of microbial cells, was obtained.

2.3 Synthetic Wastewater

The used wastewater was synthetic and contained 1/ salt solution $(NaH_2PO_4 - 3.5 \text{ g/L}, K_2HPO_4 - 5.0 \text{ g/L}, (NH_4)_2SO_4 - 2.5 \text{ g/L}, MgSO_4.7H_2O - 0.3 \text{ g/L}, FeSO_4 - 0.05 \text{ mg/L}, CuSO_4 - 0.01, ZnSO_4 - 0.005 \text{ mg/L}, CoCl_2 - 0.005 \text{ mg/L}, MgCl_2 - 0.005 \text{ mg/L}, CaCl_2 - 0.005 \text{ mg/L}, Na_2MOO_4 - 0.005 \text{ mg/L}), 2/ 3\%$ nutritious solution (NaCl 5 g/L; peptone - 10 g/L; yeast extract 5 g/L) and 3/ amaranth (from 10 to 55 mg/L).

2.4 Quartz Sand

As inert carrier for biofilters quartz sand was used. The sand particles size was between 0.08 and 0.16 cm and it was kindly provided by the Bistritza Drinking Water Treatment Plant (Sofia City, Bulgaria).

2.5 Amaranth

The model xenobiotic was supplied by Fluka Chemical Corp.

2.6 Technological Parameters

Residual amaranth concentration was determined spectrophotometrically (Utrospec 3000, PharmaciaBiotech), λ = 520 nm.

The efficiency of elimination of amaranth is calculated with the formula:

$$Eff = \frac{C_{in} - C_{eff}}{C_{in}} \cdot 100$$
, (1)

where Eff is the efficiency of amaranth removal in %, C_{in} is concentration of amaranth in the influent in mg/L, C_{eff} is the concentration of amaranth in the effluent in mg/L.

The elimination rate of amaranth is calculated with the following formula:

$$V = (C_{in} - C_{eff}).Q,$$
 (2)

where V is the rate of amaranth elimination in mg/h, C_{in} is concentration of amaranth in the influent in mg/mL, C_{eff} is the concentration of amaranth in the effluent in mg/mL, Q is the flow in mL/h.

2.7 Microbiological Analysis

The key microbial groups were studied by the plate count techniques [55]. The aerobic heterotrophs were cultivated on Nutrient Agar and bacteria from genus Pseudomonas were cultivated on Glutamate Starch Pseudomonas Agar. (Note: The samples are collected from 3 layers of the biofilters (upper, middle and bottom) and are analysed separately. Here only mean values presented). are Pseudomonas sp. ratio for culturable microorganisms was calculated on the base of aerobic heterotrophs.

2.8 FISH Analysis

The sample fixation, immobilization, dehydration and permeabilization were made according [56]. Oligonucleotide probe for microorganisms from genus Pseudomonas [57] was used to investigate the abundance and the spatial distribution of the target bacteria, key participants of amaranth biodegradation. The used probe was Cy3 5'-labelled 15-mer oligonucleotide with sequence 5-GCT GGC CTA GCC TTC-3 [58]. As a control non-sense probe was used - NON338 (5-ACT CCT ACG GGA GGC AGC-3) hybridization [57]. The was performed 20% formamide. After with hybridization samples the the were counterstained with DAPI (4,6-diamidino-2phenylindole) (AppliChem GmbH). The pictures taken with fluorescent microscope were Microsystems DFC310FX, at 400x Leica magnification.

2.9 Digital Image Analysis

The digital processing of the pictures was made by using the computer program *daime* [59]. The pictures are first segmented by using custom threshold criteria and then data for the area of interest was collected. *Pseudomonas* spp. ratio to the total quantity of the microorganisms was calculated on the basis of the pictures with DAPI and the pictures with the specific probe.

3. RESULTS AND DISCUSSION

Based on the received data for the three biofilters, the azo-degradation process can be divided on three time periods: early phase (0-191 h), late phase (191-455 h) and ending period (455-623 h). In the early phase the amaranth concentration was low (10-30 mg/L) and the decolourization efficiency was high, almost without fluctuations (Table 1). The biofilm community was in its early phase of development and adaptation toward amaranth biodegradation. In the late phase the amaranth concentration was raised from 30 mg/L up to 45 mg/L. The biofilters were well adapted and they reached highest efficiency (Table 1). In the ending period (455-623 h.) the xenobiotic concentration got close to the critical one (from 45 mg/L up to 55 mg/L) which destabilized the functioning of the biofilters. This reflected as decreased efficiency of amaranth removal for two of the three biofilters (Table 1).

Several analyses were made on the biofilm community from the biofilters in order to study the adaptation changes during the detoxification processes. As it is described in "Materials and methods", the cultivation analyses were interpreted together with culture independent techniques as FISH and digital image analysis. Microorganisms of g. *Pseudomonas* were chosen as key amaranth biodegraders [60]. The data for these microorganisms as part of the biofilm community is shown on Fig. 1.

In the early azo-detoxification phase (0-191 h) main role played the culturable *Pseudomonas sp.*, which was between 67% and 70% for the three biofilters. For this period digital image analysis showed 37% of microorganisms of g. *Pseudomonas* for biofilters 1 and 3, and 53% for the biofilter 2. The FISH pictures from this period of time suggested a low abundance of the target bacteria for the three biofilters (Table 2, pink-red fluorescent signal).

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	Early phase				Late phase			Ending period	
	Residual	Rate of	Efficiency of	Residual	Rate of	Efficiency of	Residual	Rate of	Efficiency of
	concentration,	removal, mg/h	removal, %	concentration,	removal,	removal, %	concentration,	removal,	removal, %
	mg/L			mg/L	mg/h		mg/L	mg/h	
Biofilter 1	1.75	0.41	88.35%	4.04	0.99	90.60%	4.91	1.14	89.76%
Biofilter 2	1.33	0.42	90.92%	1.72	0.91	95.73%	2.90	1.09	94.92%
Biofilter 3	1.23	0.43	92.40%	2.83	0.94	93.35%		1.21	93.71%
а) 140%		b) 140	%		c) 140%			

Table 1. Key technological parameters for amaranth removal in the three biofilters



Fig. 1. *Pseudomonassp.* ratio for the three lab-scale biofilters (BF), calculated on the base of FISH pictures (*Pseudomonas sp.* (FISH)) and on the base of aerobic heterotrophs (*Pseudomonas sp.* (culturable)) for: a) Early phase; b) Late phase; c) Ending period of the detoxification processes





In the late phase of functioning of biofilters the inflow amaranth concentration was raised 2 times. That is why the residual concentrations in the three biofilters were slightly increased. Nevertheless the efficiency of amaranth removal was even higher in the all three biofilters. These results suggested development and adaptation of the biofilm community toward azo-dye biodegradation. The FISH images confirmed this. The fluorescent signal from g. Pseudomonas in the pictures in Table 2 (late phase) is significantly increased. In the same time the culture techniques showed significantly decreased part of the microorganisms from g. Pseudomonas (from 7 up to 18 times) (Fig. 1b). Because FISH gives information not only for culturable but also for unculturable target bacteria it can be suggested that unculturable microorganisms from g. Pseudomonas play significant role in azodegradation in the conditions of the increased toxicity.

In the ending period the amaranth concentration and the rate of its removal were additionally increased from 45 mg/L up to 55 mg/L for the amaranth concentration and from 0.91-0.99 mg/h up to 1.09-1.21 mg/h for the rate of amaranth removal. In this period of time the biofilm carried out azo-degradation communities process under strong toxic pressure but their efficiency hasn't been decreased significantly (Table 1). The cultivation techniques showed an increased part of Pseudomonas sp. for the biofilter 2 with 56.95% and for the biofilter 3 with 11.62%. The digital image analysis demonstrates an increased part of the target Pseudomonas sp. in biofilter 1 (with 17.82%) and biofilter 3 (with 8.99%). These results haven't been conclusive for the role of cultural and unculturable part of the microorganism from g. Pseudomonas. However the mean values for the three biofilters (culture technique - 22.11% increase; digital image analysis - 6.07% increase) suggested that the biofilm community additionally developed. FISH

technique demonstrated structural changes in the communities adapted towards amaranth biodegradation in these unfavorable conditions. From the pictures in Table 2 it is seen that the key biodegraders (Pseudomonas sp.) were found grouped in zones with high metabolic activities. Most probably the high toxicity stimulated adaptation microorganisms to the and concentration in "hot spots" for high azodegradation. These zones are indirect proof for established cooperative microbial relations such as synergistic, symbiotic and co-metabolic relationships. This represents adaptation on higher level, where the microorganisms aren't just raised their numbers but they become interdependent in the common process - the azo-degradation of the xenobiotic Amaranth.

4. CONCLUSION

The presented study showed that a cross analysis of data from plate count, culture independent and digital techniques could reveal some of the specific mechanisms for adaptation of azo-degrading microbial community. It was established that a biofilm adaptation process with stepwise amaranth increase for 26 days consisted of three stages with specific features: early phase when the culturable 1) Pseudomonas sp. had dominant role and increase their amount in the course of adaptation; 2) late phase when the community developed and the unculurable microorganisms from q. Pseudomonas had key importance: 3) end stage, when under the high toxicity pressure, zones with high concentration and high metabolic activity of the target bacteria were formed. From these results it is seen that the microorganisms from g. Pseudomonas developed cooperative relationships for maintain the high efficiency of azo-dye removal under the strong xenobiotic inhibition.

Obtained adaptation algorithm and mechanisms of the biological control will contribute to the increase the effectiveness and applicability of the wastewater treatment technologies. All these reveal the potential for acceleration of the detoxification technologies as well as to improve the quality of water in the water receivers – natural water bodies.

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COMPETING INTERESTS

Authors have declared that no competing interests exist.

REFERENCES

- 1. World Water Council. Water at COP21 #CLIMATE IS WATER. (Accessed 26 June 2016) Available:<u>http://www.worldwatercouncil.org</u>/fileadmin/world_water_council/documents/ <u>news/2015117 ClimatelsWater Initiative I</u> eaflet.pdf
- 2. Paris pact on water and adaptation. (Accessed 26 June 2016) Available:<u>http://www.riob.org/IMG/pdf/Pact</u> <u>e_Paris_Eng_version_Non_COP_v17.pdf</u>
- Jin X, Liu G, Xu Z, Tao W. Decolourisation of a dye industry effluent by aspergillus fumigatus XC6. Appl. Microbiol. Biotechnol. 2007;74:239-243.
- Van der Zee FP, Villaverde S. Combined anaerobic–aerobic treatment of azo dyes– A short review of bioreactor studies. Water Research. 2005;39:1425–1440.
- Reddig W. Book of Papers, Cottech Europe' 96 Conf., Basle, Switzerland 11– 13 November; 1996.
- 6. Foster S. 3Rs Conf.: Retailers, regulators and results, Manchester 26 June; 2002.
- Singh SN, Iyengar L, Pandey A. Bacterial decolorization and degradation of azo dyes. In: Singh SN, editor. Microbial Degradation of Xenobiotics, Springer-Verlag Berlin Heidelberg; 2012.
- 8. Weisburger JH. Comments on the history and importance of aromatic and heterocyclic amines in public health. Mutat Res. 2002;30:506–507.
- Rai HS, Bhattacharyya MS, Singh J, Vats P, Banerjee UC. Removal of dyes from the effluent of textile and dyestuff manufacturing industry: A review of emerging techniques with reference to biological treatment. Critical Rev Environ Sci Tecnhol. 2005;35:219–238.

- Öztürk A, Abdullah MI. Toxicological effect of indole and its azo dye derivatives on some microorganisms under aerobic conditions. Sci Total Environ. 2006;358: 137–142.
- 11. Rajaguru P, Vidya L, Baskarasethupathi B, Kumar A, Palanivel M, Kalaiselvi K. Genotoxicity evaluation of polluted ground water in human peripheral blood lymphocytes using the comet assay. Mutat Res. 2002;517:29–37.
- Pinheiro H, Tourad E, Thomas O. Aromatic amines from azo dye reduction: Status review with emphasis on direct UV spectrophotometric detection in textile industry wastewaters. Dyes Pigm. 2004; 61:121–139.
- Schneider K, Hafner C, Jager I. Mutagenicity of textile dye products. J Appl Toxicol. 2004;24:83–91.
- Umbuzeiro GA, Freeman H, Warren SH, Oliveira DP, Terao Y, Watanabe T, Claxton LD. The contribution of azo dyes to the mutagenic activity of the Cristais River. Chemosphere. 2005;60:55–64.
- Chen H. Recent advances in azo dye degrading enzyme research. Curr Protein Pept Sci. 2006;7:101–111.
- Puvaneshwari N, Muthukrishnan J, Gunasekaran P. Toxicity assessment and microbial degradation of azo dyes. Indian J Exp Biol. 2006;44:618–626.
- Khalid A, Arshad M, Crowley DE. Decolorization of azo dyes by Shewanella sp. under saline conditions. Appl Microbiol Biotechnol. 2008;79:1053–1059.
- Gopinath KP, Meera Sahib HA, Muthukumar K, Velan M. Improved biodegradation of congored by using *Bacillus sp.* Bioresour Technol. 2009;100: 670–675.
- Kalyani DC, Telke AA, Dhanve RS, Jadhav JP. Ecofriendly biodegradation and detoxification of Reactive Red 2 textile dye by newly isolated *Pseudomonas sp.* SUK1. J of Hazard Mat. 2009;163:735–742.
- Saratale RG, Saratale GD, Kalyani DC, Chang JS, Govindwar SP. Enhanced decolorization and biodegradation of textile azo dye Scarlet R by using developed microbial consortium. Bioresource Technology. 2009;100:2493–2500.
- Chougule AS, Jadhav SB, Jadhav JP. Microbial degradation and detoxification of synthetic dye mixture by *Pseudomonas sp.* SUK 1. Proc. Natl. Acad. Sci., India, Sect. B Biol. Sci. 2013;84(4):1059-1068.

- Shanmugam S, Harinarayan P, Muthiah P. Response surface optimization for biodegradation of textile azo dyes using isolated bacterial strain *Pseudomonas sp.* Arabian Journal for Science & Engineering. 2013;38(9):2279-2291.
- Shah P. Microbial degradation of Azo dye by *Pseudomonas spp.* 2413 isolated from activated sludge of common effluent treatment plant maulin. Int J Environ Biorem & Biodegr. 2014;2(3):133-138.
- 24. Sriram N, Reetha D, Saranraj P. Biological Degradation of Reactive Dyes by Using Bacteria Isolated from Dye Effluent Contaminated Soil. Middle-East Journal of Scientific Research. 2013;17(12):1695-1700.
- 25. Franciscon E, Zille A, Garboggini FF. Microaerophilic-aerobic sequential decolourization/biodegradation of textile azo dyes by a facultative *Klebsiella sp.* Strain VN-31. Process Biochem. 2009;44: 446–452.
- Ghodake G, Jadhav S, Dawkar V, Govindwar S. Biodegradation of diazo dye Direct brown MR by Acinetobacter calcoaceticus NCIM 2890. Int Biodeter Biodegr. 2009;63(4):433-439.
- 27. Deng D, Guo J, Zeng G, Sun G. Decolorization of anthraquinone, triphenylmethane and azo dyes by a new isolated *Bacillus cereus* strain DC11. Int Biodeter Biodegr. 2008;62:263–269.
- 28. Mabrouk MEM, Yousef HY. Decolorization of fast red by *Bacillus subtilis* HM. J Appl Sci Res. 2008;4:262–268.
- 29. Liu YJ, Zhang AN, Wang XC. Biodegradation of phenol by using free and immobilized cells of *Acinetobacter sp.* XA05 and *Sphingomonas sp.* FG03, Biochem. Eng. J. 2009;44:187–192.
- Pavani V. Effect of physico chemical parameter on the biodegradation of azodye from *Pseudomonas aeruginosa*. World Journal of Pharmacy and Pharmaceutical Sciences. 2014;3(12):797-809.
- Kannan S, Dhandayuthapani K, Sultana M. Decolorization and degradation of azo dye
 Remazol Black B by newly isolated *Pseudomonas putida*. Int. J. Curr. Microbiol. App. Sci. 2013;2(4):108-116.
- 32. Shah MP, Patel KA, Nair SS, Darji AM, Maharaul S. Microbial degradation of azo dye by *Pseudomonas spp.* MPS-2 by an application of sequential microaerophilic & aerobic process. American Journal of

Microbiological Research. 2013;1(4):105-112.

- 33. Khehra MS, Saini HS, Sharma DK, Chadha BS, Chimni SS. Comparative studies on potential of consortium and constituent pure bacterial isolates to decolorize azo dyes. Water Res. 2005;39: 5135–5141.
- Moosvi S, Keharia H, Madamwar D. Decolorization of textile dye Reactive Violet 5 by a newly isolated bacterial consortium RVM 11.1. World J Microbiol Biotechnol. 2005;21:667–672
- Joshi T, Iyengar L, Singh K, Garg S. Isolation, identification and application of novel bacterial consortium TJ-1 for the decolourization of structurally different azo dyes. Bioresour Technol. 2008;99:7115– 7121.
- Saratale RG, Saratale GD, Kalyani DC, Chang JS, Govindwar SP. Enhanced decolorization and biodegradation of textile azo dye Scarlet R by using developed microbial consortium. Bioresource Technology. 2009;100:2493–2500.
- Moosvi S, Kher X, Madamwar D. Isolation, characterization and decolorization of textile dyes by a mixed bacterial consortium JW-2. Dyes Pigments. 2007; 74:723–729.
- Dave SR, Patel TL, Tipre DR. Bacterial degradation of Azo dye containing wastes. SN. Singh (ed.), Microbial Degradation of Synthetic Dyes in Wastewaters, Environmental Science and Engineering; 2015.
- Erkurt HA, editor. Biodegradation of azodyes. Berlin Heidelberg: Springer-Verlag; 2010.
- 40. Liu F, Zhao CC, Zhao DF, Liu GH. Tertiary treatment of textile wastewater with combined media biological aerated filter (CMBAF) at different hydraulic loadings and dissolved oxygen concentrations. J Hazard Mater. 2008;160(1):161-167.
- Ranganathan K, Karunagaran K, Sharma DC. Recycling of wastewaters of textile dyeing industries using advanced treatment technology and cost analysis— Case studies. Resour Conserv Recy. 2007;50:306-318.
- 42. Somasiri N, Li XF, Ruan NQ, Jian C. Evaluation of the efficacy of upflow anaerobic sludge blanket reactor in removal of colour and reduction of COD in real textile wastewater. Biores Technol. 2008;99:3692-3699.

- 43. Wijetunga S, Li XF, Jian C. Effect of organic load on decolourization of textile wastewater containing acid dyes in upflow anaerobic sludge blanket reactor. J Hazard Mater. 2010;177:792-798.
- 44. Lubello C, Caffaz S, Mangini L, Santianni D, Caretti C. MBR pilot plant for textile wastewater treatment and reuse. Wat Sci Tech. 2007;55(10):115–124.
- 45. You SJ, Teng JY. Anaerobic decolorization bacteria for the treatment of azo dye in a sequential anaerobic and aerobic membrane bioreactor. Journal of the Taiwan Institute of Chemical Engineers. 2009;40:500–504.
- Deowan SA, Galiano F, Hoinkis J, Figoli A, Drioli E. Submerged membrane bioreactor (SMBR) for treatment of textile dye wastewater towards developing novel MBR process. APCBEE Procedia. 2013;5:259– 264.
- Georgiou D, Hatiras J, Aivasidis A. Microbial immobilization in a two-stage fixed-bed-reactor pilot plant for on-site anaerobic decolorization of textile wastewater. Enzyme Microb Technol. 2005;37(6):597-605.
- Sandhya S, Sarayu K, Swaminathan K. Determination of kinetic constants of hybrid textile wastewater treatment system. Bioresource Technology. 2008;99: 5793–5797.
- 49. Sen S, Deminer GN. Anaerobic treatment of real textile wastewater with a fluidized bed reactor. Water Research. 2003;37: 1868–1878.
- 50. Sirianuntapiboon S, Sadahiro O, Salee P. Some properties of a granular activated carbon-sequencing batch reactor (GAC-SBR) system for treatment of textile wastewater containing direct dyes. J Environ Manage. 2007;1:162-70.
- Sahinkaya E, Uzal N, Yetis U, Dilek FB. Biological treatment and nanofiltration of denim textile wastewater for reuse. Journal of Hazardous Materials. 2008; 153(3):1142-1148.
- 52. Haroun M, Idris A. Treatment of textile wastewater with an anaerobic fluidized bed reactor. Desalination. 2009;237(1–3): 357-366.
- 53. Bulc TG, Ojstrsek A. The use of constructed wetland for dye-rich textile wastewater treatment. Journal of Hazardous Materials. 2008;155:76-82.
- 54. Ong SA, Uchiyama K, Inadama D, Yamagiwa K. Simultaneous removal of

color, organic compounds and nutrients in azo dye-containing wastewater using upflow constructed wetland. Journal of Hazardous Materials. 2009;165:696–703.

- Kuznetzov SI, Dubinina GA. Methods for investigation of waters microorganisms. Acad. Sc. USSR, Moskow; 1989.
- Nielsen H, Daims H, Lemmer H. FISH Handbook for Biological Wastewater Treatment: Identification and Quantification of Microorganisms in Activated Sludge and Biofilms by FISH. IWA Publishing: London; 2009.
- Loy A, Maixner F, Wagner M, Horn M. probebase – an online resource for rRNAtargeted oligonucleotide probes: New features 2007. 2006;35(Database issue): D800–D804. DOI: 10.1093/nar/gkl856
- Schleifer KH, Amann R, Ludwig W, Rothemund C, Springer N, Dorn S. Nucleic acid probes for the identification and *in-situ* detection of pseudomonads. In: Galli E, Silver S, Witholt B, editors. *Pseudomonas*: molecular biology and biotechnology. American Society for Microbiology: Washington DC; 1992.
- 59. Daims H, Lucker S, Wagner M. daime, a novel image analysis program for microbial ecology and biofilm research. Environ Microbiol. 2006;8:200-213.
- Grekova-Vasileva M, Popov I, Vassilev D, Topalova Y. Isolation and Characterisation of Microbial Strain AZO29 Capable of AZO Dye Decolourization. Biotech& Biotech Eq. 2009;23:(Sup1):318-322. DOI: 10.1080/13102818.2009.10818428

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