

Monitoring of activated sludge settling ability through image analysis: validation on full-scale wastewater treatment plants

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Abstract In recent years, a great deal of attention has been focused on the research of activated sludge processes, where the solid–liquid separation phase is frequently considered of critical importance, due to the different problems that severely affect the compaction and the settling of the sludge. Bearing that in mind, in this work, image analysis routines were developed in Matlab environment, allowing the identification and characterization of microbial aggregates and protruding filaments in eight different wastewater treatment plants, for a combined period of 2 years. The monitoring of the activated sludge contents allowed for the detection of bulking events proving that the developed image analysis methodology is adequate for a continuous examination of the morphological changes in microbial aggregates and subsequent estimation of the sludge volume index. In fact, the obtained results proved that the developed image analysis methodology is a feasible method for the continuous monitoring of activated sludge systems and identification of disturbances.

Keywords Activated sludge · Image analysis ·
Aggregates · Filaments · Settling ability

Introduction

Biological processes are very sensitive to external and internal variations, affecting their optimum working conditions. In activated sludge systems, an adequate balance between the different types of bacteria is essential to ensure an efficient pollution removal, good settling ability of the sludge and low suspended solids effluent levels. After the oxidation of the organic matter in the aerated tank, the flocculated biomass is separated from the treated effluent by means of their settling ability in a settling tank. The settling phase is considered a critical stage of the process in which filamentous bulking and deflocculation processes are the most common problems, causing the decrease of the sludge settling ability and effluent quality deterioration [1–7].

The solid–liquid separation in wastewater treatment plants (WWTPs) is commonly assessed through the sludge volume index (SVI) evaluation, characterizing the sludge settling ability. However, as the SVI only provides a macroscopic evaluation of the activated sludge, the microscopic characteristics of the sludge should be also determined by examination under an optical microscope. Therefore, it comes as no surprise that microscopic image analysis procedures have been used, in recent years, to complement the wastewater treatment processes mainly on the morphological characterization and evaluation of aggregated and filamentous bacteria contents. As the activated sludge process efficiency is dependent on the aggregates physical properties, some basic parameters have been used to study the settling ability of the microbial aggregates such as the filamentous bacteria contents, aggregates contents, size, shape, density, porosity and settling velocity, as well as consistency and specific surface area [8–10]. In fact, image analysis procedures are,

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nowadays, considered to be a feasible method to characterize quantitatively aggregates and protruding filamentous bacteria, and subsequently used to relate to the settling abilities [11–16], assess biomass morphology changes and monitor bulking events in pilot and full-scale plants [14, 17, 18].

Following past studies [14, 18, 19], the present work aims to survey the filamentous bacteria and aggregates contents and morphology of the biomass collected from eight full-scale WWTPs during a combined period of 2 years. For that purpose, an image processing and analysis program was developed for bright field microscopy, providing the necessary data for monitoring the biological system. The results based on image analysis procedures will be related on one hand to the operating conditions, and on the other, with the reported events in the WWTPs.

Materials and methods

The biomass used in this study was collected from the aeration basins of eight WWTPs (Sobreposta, Frossos1, Cunha, Tadim, Ucha, Pousa, Oliveira and Frossos2), treating domestic effluents, located in the North of Portugal. Samples were taken to perform physical measurements, on one hand, and microscopic observations, on the other, in order to estimate the contents and morphology of the microbial aggregates and protruding filamentous bacteria by image acquisition and analysis. The biomass settling ability was measured through the determination of the SVI in a 10 L cylindrical column (15 cm diameter), with the sludge height variation monitored for 30 min. Total suspended solids were determined by weight, and used to determine the SVI parameter [20].

Image acquisition

For each sample, a volume of 25 μ L was placed on a slide and covered with a 20 mm \times 20 mm cover slip for visualization and image acquisition for all the surveyed WWTPs. The volume deposition was performed by means of a recalibrated micropipette with a sectioned tip at the end to a diameter allowing the passage of even the larger aggregates. For each monitored day three individual sample slides were obtained for each sample and screened in the upper, median and lower third each, with images grabbed for each four fields of view.

Except for Frossos2, around 200 images per sample were acquired in bright field microscopy to obtain representative information of the sludge. All the images were acquired in a Leitz Laborlux S optic microscope (Leitz, Wetzlar), with 100 \times magnification, coupled to a Zeiss AxioCam HRc (Zeiss, Oberkochen) camera. The image

acquisition was performed in 1,300 \times 1,030 pixels and 8-bit format through the AxioVision 3.1 (Zeiss, Oberkochen) software.

The image acquisition of the Frossos2 WWTP, had already been performed for an earlier study and, relied on 20 images per sample acquired in bright field microscopy in a SZ 4045TR-CTV Olympus stereo microscope (Olympus, Tokyo), with 40 \times magnification, coupled to a Sony CCD AVC-D5CE (Sony, Tokyo) grey scale video camera. The image acquisition was performed in 768 \times 576 pixels and 8-bit format by a Data Translation DT 3155 (Data Translation, Marlboro) frame grabber using the commercial software package Image Pro Plus (Media Cybernetics, Silver Spring).

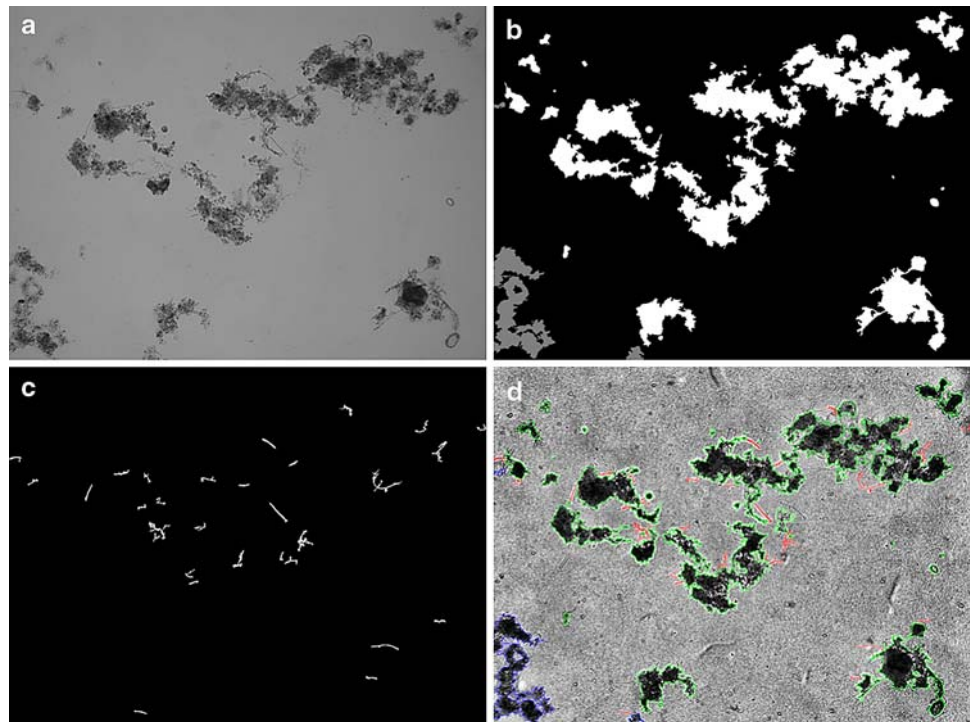
Image processing and analysis methodology

The image processing and analysis program for aggregates and filaments, was developed in Matlab 7.3 (The Mathworks, Inc., Natick) language, adapting a previous version developed by Amaral and Ferreira [18]. Primarily, the image processing step determined the binary images from the aggregated biomass and the protruding filamentous bacteria and thereafter, the morphological parameters were determined. Figure 1 shows an example of the original, binary and labeled images resulting of the main steps of the program, comprising the image pre-treatment, segmentation, and debris elimination whereas the image analysis program was oriented to the aggregates and filaments contents determination.

Image analysis parameters

Supported on the previous study of Amaral and Ferreira [18], 36 parameters were determined either directly from the image analysis program either in association with the sludge physical properties for a total around 400 000 aggregates. Total aggregates number per sludge volume, total aggregates area per sludge volume, aggregates area, total filaments length per sludge volume, filaments length, aggregates length, aggregates perimeter and aggregates equivalent diameter were determined for all the samples collected. The morphological descriptors convexity, solidity, roundness, and eccentricity were also determined by the image analysis methodology. Furthermore, the aggregates characterization was subsequently divided in three classes in order to differentiate between small low settling structures (contained within the aggregates below 0.025 mm in equivalent diameter), normal sized good settling structures (between 0.025 and 0.25 mm in equivalent diameter), and large connected low settling structures (contained within the aggregates above 0.25 mm in equivalent diameter). The total filaments length per sludge

Fig. 1 Original image from the activated sludge system with $\times 100$ magnification (a), binary aggregates image (b), binary filaments image (c) and final labeled image (d)



volume, filaments average length per aggregates average area ratio, filaments average length per aggregates average equivalent diameter ratio and the total filaments length per total aggregated area ratio were determined alongside the total filaments length per volatile suspended solids ratio characterizing the filaments dynamics. A more detailed description of each parameter can be found in Amaral [19]. It should also be pointed out that a study [21] has also been conducted regarding the reduction of the parameters number for the best nine parameters, representing the free filamentous bacteria contents and characterization and aggregates contents, size and morphology, yielding satisfying results (reduction of only 0.034 on the correlation coefficient). Therefore, the use of the reduced data set may lead to a decreased correspondence between the predicted and observed SVI although not significantly far apart from the ones reported here.

Partial least squares

The collected data set (5,184 data points from 144 samples \times 36 parameters) was correlated by Partial Least Squares (PLS) regression, an iterative algorithm that extracts linear combinations of the essential features of the original X data while modeling the Y data dependence on the work set, being therefore well suited for multivariate calibration. PLS have been shown to be an efficient approach in monitoring complex processes since the high dimensional strongly cross-correlated data can be reduced

to a much smaller and interpretable set of latent variables [22]. To perform the PLS analysis from the data set, SIMCA 8.0 (Umetrics, Um ea) software package was used. SIMCA computes the variable importance in the projection (VIP) as the sum over all model dimensions (PLS components) of the variable influence. The parameters that are found to be the most important are the ones presenting a VIP value larger than 1. A more detailed description about this method can be found in Umetri [23].

Results and discussion

The PLS analysis allowed to predict the SVI values from the image analysis data with a reasonable correspondence as it can be seen in Fig. 2. Analyzing the results it was found that the multiple correlation coefficient (goodness of fit) attained for the SVI prediction was 0.947 (R^2 of 0.897), allowing to predict, at some extent, the SVI values from the obtained image analysis data.

It must be noticed that, above the 200 mL/g SVI threshold, there seems to be a higher dispersion regarding the observed versus the predicted results and, hence, the accuracy of the prediction will not be as high as for lower SVI values. However, this behavior may be explained by the fact the majority of the data points above 200 mL/g were acquired in a previous study with an acquisition of an average of 20 images per sample (acquisition for Frossos2 WWTP) instead of the average 200 images for the other

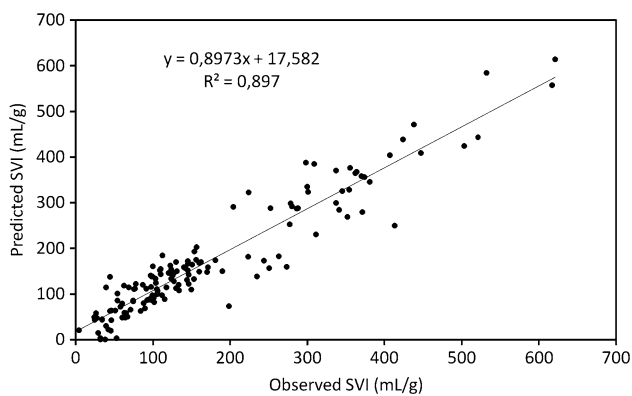


Fig. 2 Global correlation between the observed and predicted SVI data points

WWTPs. In accordance to this explanation, no significant correlations were found between the errors of the predicted SVI and the SVI values within the Frossos2 data set (R^2 of 0.0171) and between the errors of the predicted SVI and the SVI values within the remaining WWTP data set (R^2 of 0.0004).

The observed SVI values were monitored throughout eight different WWTPs and are presented in Fig. 3, alongside the predicted SVI values. From the analysis of this figure it can be perceived that the predicted SVI values follow, at some extent, the observed SVI trends corroborating the findings of the PLS analysis results. Furthermore, it was found that the observed and predicted SVI values differ between each other around 31 mL/g in average. It was also found that the difference between the values is smaller for the lower SVI values, around 24 mL/g for the set of SVI values below 200 mL/g, and larger for higher SVI values, around 49 mL/g for the set of SVI values above 200 mL/g. Once again this behavior may be explained by the fact the majority of the data points above 200 mL/g rely on an average of 20 images per sample

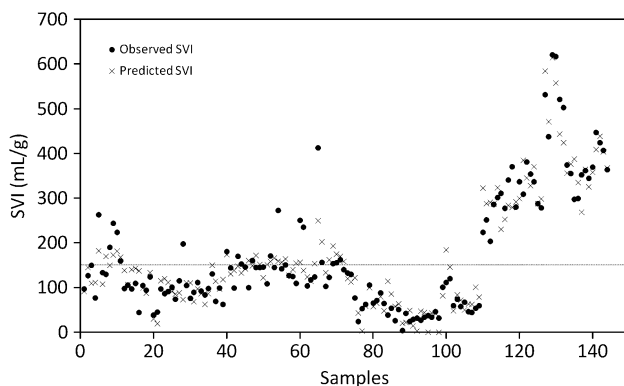


Fig. 3 Global observed and predicted SVI data points from the WWTP aerated tanks

instead of the average 200 images for the data points below 200 mL/g.

In most cases for the SVI below the bulking limit (150 mL/g according to Jenkins et al. [5]) there is a correspondence to the predicted SVI (presenting also values below 150 mL/g). An analogous trend is also found for the values indicating bulking phenomena with predicted SVI values above the 150 mL/g limit. In fact for the 89 data points that presented SVI values below 150 mL/g, this methodology correctly predicted 77 (86.5%) of these as non-bulking situations, whereas for the 55 data points presented SVI values above 150 mL/g, this methodology correctly predicted 49 (89.1%) of these as bulking situations. Given the 24 mL/g uncertainty for the SVI values below 200 mL/g and the fact that the bulking limit is set on 150 mL/g, a closer analysis of the values outside this uncertainty area (150 ± 24 mL/g) reveals that only 6 (6.7%) values below the lower limit and 3 (5.5%) values above the upper limit are incorrectly classified.

For an in-depth analysis of each of the WWTP, their individual SVI values (observed and predicted) are represented in Fig. 4. From this figure, it can be withdrawn that the eight aerated tanks have different operating conditions, and some with strong fluctuations in terms of the sludge settling abilities.

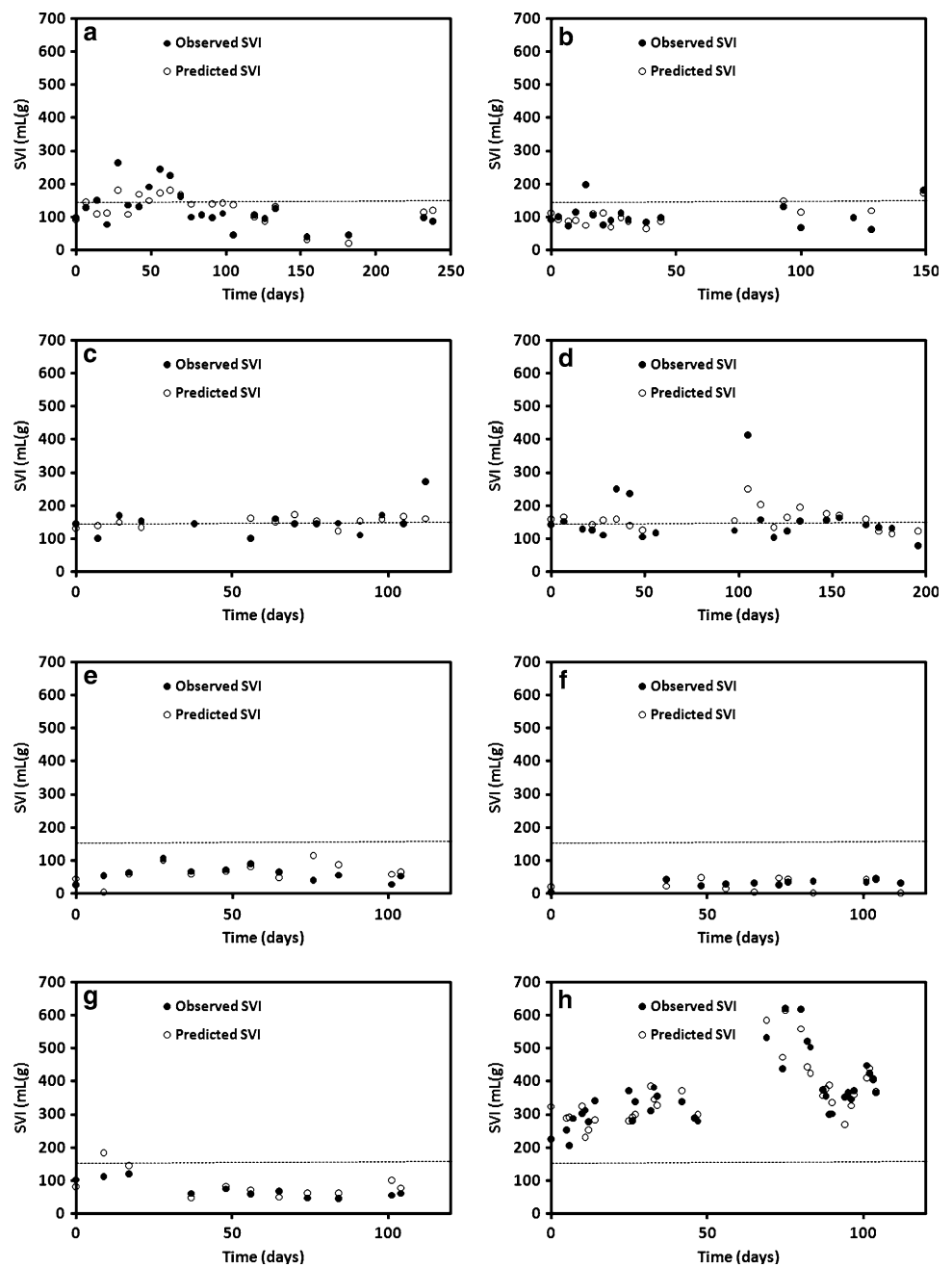
For Sobreposta WWTP, the SVI results, here depicted on Fig. 4a, presented values mainly below the 150 mL/g threshold limit for bulking events. Only 5 days (days 28, 49, 56, 63 and 70) presented SVI values higher than 150 mL/g (bulking phenomena), ranging from 160 to 265 mL/g. The predicted values for this WWTP were, for the most, in accordance with the observed SVI values with larger deviations occurring on days 28, 56 and 105 (all above 50 mL/g). With respect to the bulking prediction ability, only 2 days (out of 23) were incorrectly classified, with one of those days falling into the uncertainty area (150 ± 24 mL/g), thus meaning that only 1 day was effectively misclassified.

The SVI behavior in Frossos1 WWTP, depicted on Fig. 4b, revealed a non-bulking system with the only exceptions on days 14 and 149 with 198 and 181 mL/g, respectively. The predicted values for this WWTP were, for the most, in accordance with the observed SVI values with larger deviations occurring on days 14 and 128 (above 50 mL/g). With respect to the bulking prediction ability, only 1 day (out of 17) was misclassified, falling outside the uncertainty area (150 ± 24 mL/g).

Regarding the Cunha WWTP (Fig. 4c), the SVI values were always on the threshold of configuring a bulking phenomenon and in days 14, 21, 64, 98 and 112 were, in fact, higher than 150 mL/g. Furthermore, in the last monitored day, the settling properties of the biomass suffered a sudden deterioration with an SVI value of 273 mL/g.

Fig. 4 Observed and predicted SVI data points from the eight WWTP aerated tanks.

a Sobreposta, **b** Frossos1, **c** Cunha, **d** Tadem, **e** Ucha, **f** Pousa, **g** Oliveira, **h** Frossos2



Again, the predicted values for this WWTP were, for the most, in accordance with the observed SVI values with larger deviations occurring on days 56 and 112 (above 50 mL/g). With respect to the bulking prediction ability, 8 days (out of 14) were incorrectly classified, but six of those days fell into the uncertainty area (150 ± 24 mL/g), thus meaning that only 2 days were effectively misclassified.

The SVI behavior of Tadem WWTP is depicted on Fig. 4d, showing values around the threshold limit for bulking events although such was not the case for a few periods of the survey. In fact, for day 7, day 112 and from day 133 to 154 the plant presented mild bulking

phenomena, and for days 35 to 42 and day 105 considerable bulking problems. It was found a good correspondence between the predicted and observed SVI values for this WWTP, with larger deviations occurring only on days 35, 42 and 105 (above 50 mL/g). With respect to the bulking prediction ability, 6 days (out of 21) were incorrectly classified, with two of those days falling into the uncertainty area (150 ± 24 mL/g), thus meaning that 4 days were effectively misclassified.

Regarding the Ucha WWTP (Fig. 4e), low SVI values predominated throughout the survey, as the SVI remained below the bulking threshold (150 mL/g) during all the

monitoring period. Such was also the case of the Pousa WWTP (below 50 mL/g) and Oliveira WWTP (between 40 and 120 mL/g), depicted on Fig. 4f, g, respectively. Once more, the predicted values for these WWTPs were in accordance with the observed SVI values with a larger deviation (above 50 mL/g) occurring solely on day 9 of the Oliveira WWTP. With respect to the bulking prediction ability, only 1 day (out of 34) was misclassified, falling outside the uncertainty area (150 ± 24 mL/g).

The SVI behavior in Frossos2 WWTP, depicted on Fig. 4h, revealed a bulking problem, throughout all the monitoring period, with values ranging from 204 up to 617 mL/g. It was found a reasonable correspondence between the predicted and observed SVI values for this WWTP with larger deviations (above 50 mL/g) occurring on 12 of the 35 days (days 1, 6, 11, 14, 25, 32, 69, 80, 82, 83, 89 and 94). Again it should be emphasized the fact that this WWTP data points rely on an average of 20 images per sample instead of the average 200 images for the remaining WWTP, hence, explaining the poorer correspondence between the predicted and observed SVI values. However, and with respect to the bulking prediction ability, not a single day (out of 35) was incorrectly classified. Furthermore, it should also be stressed that, the predicted SVI followed the observed SVI behavior throughout all the monitoring period.

In a global analysis, it should also be noticed that, the two WWTP that were chronically on the threshold of bulking events (Cunha and Tadim) accounted for 14 of the 18 days incorrectly classified, and taking into account the uncertainty area (150 ± 24 mL/g) for 6 of 9 days effectively misclassified.

Conclusions

The PLS analysis of the image analysis data allowed to predict the SVI values with a reasonable correspondence as it can be inferred by the 0.947 multiple correlation coefficient obtained between the observed and predicted SVI values. It should be kept in mind, however, that although the obtained correlation holds true for the SVI range presented in the surveyed WWTPs (up to 700 mL/g) no direct extrapolations can be performed for higher SVI values regarding severe bulking phenomena.

It was also found that, for the current methodology, the observed and predicted SVI values distanced from each other an average 24 mL/g for the set of SVI values below 200 mL/g, which comprises the bulking threshold limit. This fact allowed the establishment of an uncertainty area (150 ± 24 mL/g) regarding the effectiveness of classifying bulking/non-bulking situations. That being the case, the results revealed that only 9 out of 144 (6.3%) outside the

uncertainty area were, in fact, incorrectly classified according to this methodology.

In conclusion, the obtained results showed that the developed image analysis and PLS methodology proved to be adequate for continuous examination of sludge settling properties, in terms of the SVI. Furthermore, this methodology was able to predict, for the most part, the disturbances within the aerated tank of eight different WWTPs.

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