Enumeration of *Brettanomyces* in Wine Using Impedance

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Abstract: *Brettanomyces bruxellensis* is a wine spoilage concern in wineries around the world. In order to maintain wine quality during storage and ageing, it is imperative to control and monitor this yeast. Being a fastidious slow growing yeast, which requires 5 to 14 days of incubation for visible growth in agar plates, it is difficult to detect growth (colonies) by conventional agar plate count method. Yeast enumeration by impedance was investigated because previous research using other microorganisms has shown that it is potentially faster than plate counting. The relationship between plate counting and impedance detection times was investigated for *Brettanomyces* inoculated in red wine samples. A linear relationship between log plate count concentrations and impedance detection times was found. Incubation time was reduced from 120 h down to 0.9 and 57.7 h for samples with $6.7 \times 10^7$ and $1.8 \times 10^2$ cfu/mL, respectively, using the ‘indirect’ impedance method. The ‘direct’ method also reduced the incubation times to 9.5 and 81.9 h, for the same concentrations. The ‘indirect’ impedance method has the potential to be used by the wine industry to control and monitor the *Brettanomyces* numbers in wines.

Keywords: *Brettanomyces bruxellensis; Dekkera bruxellensis*; spoilage; microbial detection; analytical method; quality control

1. Introduction

In 2015, the average global wine production was approximately $2.8 \times 10^{10}$ L [1,2]. The yeast *Brettanomyces bruxellensis* (the anamorph *Dekkera bruxellensis*) is one of the major spoilage organisms faced by the wine industry, which leads to economic losses worldwide [3,4]. Concentrations as low as $10^3$ cfu/mL of this spoilage yeast have been known to cause unpalatable off-odors and flavors [5]. *Brettanomyces* spoilage is most frequently associated with red wines due, in part, to the increased use and efficiency of sulfur dioxide preservative in low pH white wines [3]. *Brettanomyces* contamination has been detected in wines and wineries from all major wine producing countries around the world [3,4]. The development of fast and effective methods for their detection and enumeration in the wine industry is therefore vital.

General plating media, including yeast extract peptone dextrose (YPD) and *Dekkera/Brettanomyces* differential medium (DBDM), can be used with the addition of 20–100 mg/L of cycloheximide to detect *Dekkera/Brettanomyces bruxellensis*. Due to reliability and low cost, plate counting is still the predominant method for detecting this yeast in the industry [5–7]. *B. bruxellensis* typically has plate incubation times of more than 72 h (3 days) at 25–30 °C [3,5]. Costly and time-consuming gas chromatography mass spectrometry (GC-MS) can also be used to detect these yeasts by measuring 4-ethylphenol concentrations. Real-time polymerase chain reaction (PCR) method has been shown to detect as little as 10 cells/mL in under 3 h, with highly specific detection and enumeration of targeted microorganisms [5,7]. One disadvantage of PCR is that the wine matrix is complex, complicating DNA extraction and causing amplification problems [8]. Microscopic examination of the distinctive morphology of *Brettanomyces* is often used in conjunction with
microscopic enumeration using Malassez or Thoma chambers [8,9]. Even though specific cell functions can be determined by staining cells, this is a time consuming and highly subjective method [8]. Malacrino et al. (2001) found that flow cytometry in combination with fluorescent dyes can be used to estimate yeast and bacteria counts in wines [10]. This is a rapid enumeration method, with high specificity and the ability to analyze the physiological state of cells [8].

Due to the lower growth rate of *Brettanomyces* compared to other yeasts, new technologies, including impedance, could have the potential to reduce detection and enumeration times. Impedance is the effective resistance of an electrical system to alternating electric current. The impedance enumeration method measures the change in impedance caused by microbial growth in a selective medium under an alternating current (AC) field. The lower molecular size charged compounds produced during microbial growth cause the change in impedance. The BacTrac™ system utilizes impedance splitting technology to measure the change in the impedance of the electrode (*E*-value) and in the medium (*M*-value). The impedance splitting technology measures the change in the impedance of the electrochemical double layer of the electrode in addition to the standard impedance signal (media impedance = *M*-value). Impedance detection time is defined as the time from the start of the measurement period until the signal exceeds the specified *M*- or *E*-value threshold. For both methods, impedance is evaluated based on the growth of microbial cells added to the BacTrac measuring vials. Therefore, more cells result in more growth and shorter impedance detection times. A calibration curve is then used to convert the detection times to colony forming units (cfu) [11–13].

Impedance technology has been used to enumerate *Dekkera bruxellensis*, *Saccharomyces cerevisiae*, *Candida stellata* and *Hanseniaspora uvarum* in a model wine, *Saccharomyces fragilis* in fermented goat milk, *Bifidobacterium lactis* in milk powders and *Escherichia coli* in simulated milk ultra-filtrate and molluscs [14–18]. As wine is a complex mixture of interacting chemicals, it is important to investigate the viability of impedance technology using a real wine instead of a model wine. In this study, filter sterile red wine samples were inoculated with only *B. bruxellensis* with the main objectives: (i) to investigate the relationship between plate counting and impedance detection times in red wine; and (ii) to validate and compare the ‘direct’ and ‘indirect’ impedance methods using a different red wine.

### 2. Materials and Methods

#### 2.1. Characteristics of Wines Used

For the calibration lines, Cabernet Sauvignon red wine (Australian, 2014 vintage, 13.5% v/v alcohol, 3.5 pH, 6.6 g/L total acidity, 30 mg/L free SO₂, 73 mg/L total SO₂) was used. Cabernet Sauvignon is the most popular table red wine [1,2]. As the focus was to solely enumerate *Brettanomyces*, the wine was filter sterilized (0.45 µm pore size) and the sterility of the wine was verified using plate counting. Then, inoculation of sterilized wine with different concentrations of *B. bruxellensis* was carried out. In addition, validation experiments were conducted using the 2015 vintage of the same wine.

#### 2.2. Yeast Propagation and Inoculation

*B. bruxellensis* yeast strain AWRI 1499 (Australian Wine Research Institute, Adelaide, Australia), was used in this study as it was the most resistant to non-thermal high pressure inactivation of three strains investigated previously by van Wyk and Silva (2017) [19]. According to the method used by van Wyk and Silva (2017), the yeast was first streaked onto YPD agar (adjusted pH between 5 and 6, autoclaved at 121 °C for 15 min) and incubated for 5 days at 28 °C. One colony was then transferred to autoclaved YPD broth (adjusted pH between 5 and 6). After incubation (30 °C, 120 rpm for 4 days) and at the end of the microbial exponential growth phase, the cells were harvested and the sterilized wine inoculated. To ensure that the yeast remained fresh and active, it was transferred to fresh YPD agars on a monthly basis [19]. The Difco YPD broth and agar media were obtained from Fort Richard Laboratories, Auckland, New Zealand.
Regarding wine inoculation, dilutions of *B. bruxellensis* in sterile Ringer’s solution, with concentrations ranging between $10^2$ and $10^8$ cfu/mL were prepared. Preliminary results showed that the use of Ringer’s solution instead of wine for dilutions gave more consistent results, with less variability. The dilutions were then centrifuged (5000 rpm, 15 min) and the yeast transferred to 10 mL of filter sterilized wine samples.

### 2.3. Enumeration of *B. bruxellensis*

#### 2.3.1. Plate Counting

The concentration of *B. bruxellensis* yeast in the wine samples were determined by spreading 100 µL of appropriately diluted samples onto duplicate YPD agar plates. Given this organism is a fastidious slow grower, 5 days at 28 °C were used for incubation. To ensure a uniform yeast suspension in the wine samples, a vortex mixer was used after each stage of dilution. Yeast colonies were then counted and *B. bruxellensis* concentration was expressed in colony forming units per milliliter (cfu/mL) of wine sample.

#### 2.3.2. Impedance Technology

The BacTrac™ 4300 instrument (Sy-Lab, Neupurkersdorf, Austria), which consists of one incubator (64 measuring cells) and computer software, was used to perform impedance measurements. Figure 1 includes a detailed diagram of ‘direct’ and ‘indirect’ impedance methods. Selective media for yeast and mould BiMedia 510A and 501B nutrient broths were used for ‘direct’ and ‘indirect’ enumeration impedance methods, respectively. The ‘direct’ method measures the impedance of the nutrient broth-sample mixture in direct contact with the electrodes in the measurement cell. For the ‘indirect’ method, the nutrient broth-sample mixture is contained in an isolated smaller inner cell located inside the measurement cell. The bottom of the measurement cell contains a dilute solution of potassium hydroxide (KOH), which undergoes a change in impedance through the uptake of CO$_2$ from the microorganisms in the inner cell during metabolism. Therefore, there is no direct contact between the sample mixture and the electrodes. The following detection parameters were used for the ‘direct’ method: delay time 1 h, evaluation type E and threshold E-value 12%. The parameters for the ‘indirect’ method were: delay time 1 h, evaluation type M2 and threshold M-value –25%. For the ‘direct’ method, 1 mL of homogenized wine sample was inoculated into 9 mL of BiMedia 510A in prefilled cells. For the ‘indirect’ method, 1 mL of 0.2% KOH solution was added to the outer cell and 1 mL of homogenized wine sample was added to 5 mL BiMedia 501B in the inner vial. The cells were sealed using plastic screw caps before being inserted into the BacTrac incubator. Next, the samples were incubated at 30 °C for up to 90 h, during which cell growth occurred and thus changing the total impedance, which was recorded in 10 min intervals. The method was adapted from protocol H51 and V7.50.1 (supplied by Sy-Lab) for the ‘direct’ and ‘indirect’ methods, respectively. Previous research demonstrated that BiMedia 510 A and B supports *B. bruxellensis* growth [13].

### 2.4. Calibration Lines of Impedance versus Plate Counting

Ten wine samples, with yeast concentrations, ranging between $10^2$ and $10^8$ cfu/mL were prepared as previously described. Each sample was then analyzed by transferring 1 mL to the ‘direct’ and ‘indirect’ measuring cells containing 9 mL and 5 mL of growth medium, respectively. Five repeat measurements of each sample were performed using both the ‘direct’ and ‘indirect’ methods according to the procedure described. In addition, each wine sample was plated in duplicate on YPD agar according to conventional plate counting. Colonies were counted and the *Brettanomyces* concentration (N, N$_{PC}$) was calculated. Lastly, the log yeast concentration plate count results were plotted against the impedance detection times from the ‘direct’ and ‘indirect’ methods, generating two calibration lines. Linear regression analysis was performed using IBM SPSS Statistics (Version 23, Armonk, NY, USA).
Figure 1. Schematic of the ‘direct’ and ‘indirect’ impedance methods.

2.5. Validation of Calibration Lines Using a Different Wine

To validate the calibration lines, a wine from a subsequent vintage was used. Wine samples with five different concentrations of Brettanomyces were prepared and analyzed using the impedance methods and reference plate counting. The wine samples were prepared according to the method described, with concentrations ranging between $10^3$ and $10^8$ cfu/mL. The B. bruxellensis concentration of each sample was then determined using the plate counting and the ‘direct’ and ‘indirect’ impedance methods. For plate counting, samples were analyzed in duplicate, whereas for the impedance methods, five repeat measurements were made.

2.6. Statistical Analysis

The mean and standard deviation of all Brettanomyces concentrations ($N$) by standard plate counting ($N_{PC}$) and impedance detection ($N_I$) times were calculated using IBM SPSS. The results were then plotted as mean ± standard deviation (SD). Percentage error was calculated using:

$$\text{Percentage error} \% = \left( \frac{\log N_I - \log N_{PC}}{\log N_{PC}} \right) \times 100$$  \hspace{1cm} (1)

The yeast concentration for the ‘direct’ and ‘indirect’ impedance methods were calculated using the calibration equations (Equations (2) and (3), respectively). To validate the linear equations, the impedance and plate count results obtained were compared using Tukey tests followed by one-way analysis of variance (ANOVA), with a confidence level of 95% ($p < 0.05$).
3. Results and Discussion
3.1. Comparing the Relationship between B. bruxellensis Plate Counting and the ‘Direct’ and ‘Indirect’ Impedance Enumeration

The calibration lines for the enumeration of Brettanomyces in wine using the ‘direct’ and ‘indirect’ impedance methods are shown in Figure 2. Clearly, as the B. bruxellensis concentration \(N\) (cfu/mL by plate counting) increased, the impedance detection time \(t\) (hours) decreased. A strong linear relationship between \(\log N\) and detection time \(t\) was found, as shown by the \(R^2\) values:

\[
\log N = -0.072t + 8.23 \quad R^2 = 0.91 \quad \text{‘Direct’ impedance (2)}
\]

\[
\log N = -0.091t + 7.55 \quad R^2 = 0.96 \quad \text{‘Indirect’ impedance (3)}
\]

Overall, the detection times of the ‘indirect’ method were lower than the ‘direct’ method, especially at lower yeast concentrations. For the ‘direct’ and ‘indirect’ impedance, the detection times were 9.5 and 0.9 h for samples containing \(6.7 \times 10^7\) cfu/mL (upper limit of detection). For a concentration of \(1.8 \times 10^2\) cfu/mL (low limit of detection), the detection time for the ‘direct’ impedance was 81.9 h. This is significantly longer than the corresponding detection time for the ‘indirect’ impedance, which was 57.7 h. Indirect impedance thus decreased the incubation and the Brettanomyces detection time in wine compared to 5 days (120 h) taken for conventional plate counting.
No previous research has been conducted on using impedance to detect and enumerate *B. bruxellensis* contamination in wine. However, a study with *B. bruxellensis* in beer, also resulted in a log-linear relationship between plate count concentrations and impedance detection times, and ‘indirect’ impedance detection times of 1.9 and 50.0 h, respectively, for yeast concentrations of $10^6$ and $10^4$ cfu/mL [13]. The ‘indirect’ method was also found to give satisfactory results when working with yeasts, including *Dekkera bruxellensis*, *S. cerevisiae*, *Candida stellata* and *Hanseniaspora uvarum* [15,20]. *B. lactis* in milk powders also exhibited a log-linear relationship using the ‘direct’ impedance, with microbial concentrations between $10^6$ and $10^9$ cfu/mL, resulting in detection times of 15 and 3 h, respectively [17].

The standard deviations of the impedance detection times (horizontal error bars) were higher than for plate count results (vertical error bars). While the standard deviation of plate count results never exceeded 0.2, the standard deviations of the impedance detection times were 0.0 to 2.1 and 5.3 to 7.4 h for $6.7 \times 10^7$ and $1.8 \times 10^5$ cfu/mL yeast concentrations, respectively. Overall, the ‘indirect’ impedance is more suited for use in the wine industry for the enumeration of *B. bruxellensis* due to significantly faster detection times compared to the ‘direct’ impedance over the same detection range (Figure 2). The ‘indirect’ impedance is potentially less influenced by the type of wine samples analyzed, as they are not in direct contact with the electrodes. However, for the ‘direct’ impedance, the ionic composition of the growth medium and wine sample mixture could be of greater influence. Since no two wines have the same composition or properties, this potentially decreases the suitability of using the ‘direct’ method in the wine industry.

### 3.2. Validation, Advantages and Limitations of the ‘Direct’ and ‘Indirect’ Impedance

Five red wine samples of a different vintage, with concentrations ranging from $10^3$ and $10^7$ cfu/mL, were analyzed. The enumeration of *Brettanomyces* in these samples was conducted using plate counting as a reference. Equations (2) and (3) were then used to calculate the ‘direct’ and ‘indirect’ impedance yeast concentration, respectively. A comparison between the three *B. bruxellensis* enumeration methods is presented in Figure 3. The ‘direct’ method was slightly more accurate with an average percentage error of 15.9%, compared to that of the ‘indirect’ method, which was 18.6% (Equation (1)). For samples with a *Brettanomyces* concentration above $10^4$ cfu/mL, the ‘indirect’ method generally proved to be more accurate, with an average percentage error of 8.2% compared to that of the ‘direct’ method (13.2%). Contrastingly, the ‘direct’ method was more accurate at concentrations below $10^4$ cfu/mL, with an average percentage error of 19.8% compared to 34.2% for the ‘indirect’ method. As expected, samples with a low yeast concentration are more difficult to quantify and thus presented results with higher variability. These results are in agreement with the results described earlier.

One drawback of plate counting is the prolonged incubation periods of 120 h (5 days) required for observing *Brettanomyces* colonies on the plates. Using the ‘indirect’ impedance, incubation times are reduced to 0.8 to 58 h. The reduced incubation time could be explained by the accelerated yeast proliferation rate in liquid broth compared to semi-solid agar medium, conventionally used for plate counting. Moreover, the principle of detection is different in the semi-solid agar medium contained in plates, where every cell is counted by itself, and in the liquid broth, where all cells grow together. Avoiding prolonged incubation times also decreases the chance of sample contamination during the enumeration process. Accelerated detection and enumeration times are especially important for *Brettanomyces* in connection with the wine industry. This would enable faster implementation of preventative actions in the wine industry, such as the addition of more SO₂ and cleaning of equipment, to reduce the negative effects of *Brettanomyces* contamination and hence wine quality degradation. The risk of large-scale wine dumping due to *Brettanomyces* growth would also be reduced, in turn reducing economic losses.
Figure 3. Enumeration of *Brettanomyces bruxellensis* contamination in wine: a comparison between plate counting and impedance technology; \( N \) is the concentration of *B. bruxellensis* in red wine (cfu/mL); different letters indicate that the average \( \log N \) (logarithm of yeast concentration) are significantly different according to Tukey’s test (\( p < 0.05 \)).

One other important aspect is that standard plate counting is more labor intensive and possibly more time consuming, due to the multiple operations required for sample preparation. For the ‘direct’ and ‘indirect’ impedance methods, the sample preparation time can be significantly reduced from approximately 1 h (plate counting) to less than 5 min (impedance methods) per sample. Therefore, impedance technology has the potential to increase the throughput of enumeration analyses thereby providing faster responses to wine producers. Figure 3 and percentage error calculations (Equation (1)) demonstrated that the ‘indirect’ impedance method successfully (average error of 18.6% against plate counting as a reference) detected and enumerated the *Brettanomyces* yeast in wine. Lastly, even though the initial capital and material costs for the impedance technology is higher, this is offset by the reduction of economic risk, labor and turnaround time of samples analyzed.

This study is focused on wine samples inoculated only with *B. bruxellensis*. In addition, only exclusively fresh cultures of *B. bruxellensis* were used in the experiments. Since the age and viability of yeasts (and other microorganisms) are different in real wines, future research on the enumeration of *B. bruxellensis* in unfiltered wines should be conducted, including the identification of the impact of the age/condition of the yeast cells. Unfiltered wines contain the main fermenting organism and contaminant microorganisms (e.g., other yeasts and lactic acid bacteria) that can interfere with impedance detection. This could involve pre-treating the wine samples before analysis or adding antimicrobial agents to the detection media directly. Cycloheximide, to which *Brettanomyces* is known to be highly resistant, has been used to isolate this yeast from real wine samples, as *S. cerevisiae*’s (the predominant fermenting yeast) is susceptible to this antimicrobial additive [21].
4. Conclusions

‘Direct’ and ‘indirect’ impedance were able to detect \( B. \text{bruxellensis} \) concentrations as low as \( 1.8 \times 10^2 \) CFU/mL. The ‘indirect’ impedance has a greater potential for use in the wine industry for the enumeration of \( B. \text{brettanomyces} \), being able to decrease the processing time of wine samples containing \( B. \text{brettanomyces} \) by between 62 h and 119 h (at \( 1.8 \times 10^2 \) CFU/mL and \( 6.7 \times 10^2 \) CFU/mL, respectively), depending on concentration. For \( B. \text{brettanomyces} \) concentration above \( 10^4 \) CFU/mL, the ‘indirect’ method was more accurate overall compared to the ‘direct’ impedance. The ‘indirect’ impedance, in addition to offering faster detection times, is potentially less influenced by the type or composition of the wine analyzed. Therefore, the same calibration lines can possibly be used for different vintage year red wines, saving time. In conclusion, the ‘indirect’ impedance has the potential to be an alternative option for the enumeration of yeasts in the wine industry because it offers faster preparation time, high throughput and has the potential to reduce economic losses. Further research is recommended using different wines and unfiltered wines that contain other microorganisms typically present in wines.

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