



## Use of Chitosan to Reduce SO<sub>2</sub> in Finished Wine

*Chateau Morrisette*

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### Summary

Grapes enter the winery covered in a host of microbes, some of whom contribute to complexity of flavor and aroma, while others may lead to spoilage. Chitosan has been used as an antimicrobial treatment at nearly every stage of winemaking. In this experiment, a single harvest lot of Chambourcin was split so that half received treatment with chitosan (Stab Micro M) while the other half did not. Both bins received 40 mg/L SO<sub>2</sub>. Resulting chemistry and microbial load were assessed post fermentation and after aging. There were no notable differences in wine chemistry, volatile acidity, or microbial load after treatment with chitosan during crush.

### Introduction

Grapes enter the winery covered in microbes from the vineyard. To date, 52 different species of yeast from 22 different genera have been identified on grapes including *Hanseniospora* (AKA *Klockera*), *Candida*, *Pichia*, *Hansenula*, *Metschnikowia*, *Sporobolomyces*, *Cryptococcus*, *Thodotorula*, and *Aurobasidium*<sup>1</sup>. The cast of characters changes as grapes ripen, with the greatest abundance of microbes present in the last few weeks<sup>1</sup>. Healthy grapes are generally inhospitable environments for any microbe because they are covered in plates of wax that form a cuticle to hold in nutrition and repel water. Microbes cluster around the stomata or next to cracks in the cuticle where seepage from the openings provides both water and nutrients. The overall microbial load on grapes depends on environmental factors such as climate, altitude, variety, age of grapes, disease pressure and vineyard practices. Fog, rain, and fruit damage (like that seen in wet vintages like 2018) quickly transform the microbial desert of grape skins into an oasis. Cells that are present in small numbers quickly multiply when given the chance<sup>1</sup>. For example, *Botrytis* infection can increase the overall abundance of microbes by 1000x. Grapes with sour rot have significantly higher microbial diversity and abundance<sup>2</sup>. Insect pressure will also increase abundance due to increased vectoring from diverse environments<sup>3</sup>. The overall inoculant of non-*Saccharomyces* yeast and bacteria coming into the winery from the vineyard on the grapes is often larger than the inoculant of selected *Saccharomyces* yeast added at the beginning of fermentation<sup>1</sup>.

Non-*Saccharomyces* yeast have several impacts on the wine, both positive and negative. *Klockera apiculata* (aka *Hanseniospora uvarum*) is a common member of the non-*Saccharomyces* yeast community found on grapes<sup>1,4</sup>. This yeast strain is easily identified under a microscope by its lemon shaped cells. It is tolerant to up to 100 mg/L SO<sub>2</sub>, can grow at low temperature (such as that found during cold soak), and can produce both acetic acid and ethyl

acetate (which smells like nail polish remover) under aerobic conditions<sup>3</sup>. Other offenders in the non-*Saccharomyces* yeast community include *Pichia guilliemondii*, a film forming yeast prevalent in warm conditions when fermentation is delayed. This yeast can form spores that become resident in barrels and produce ethyl acetate and 4 ethyl phenol (which can smell like band-aid, wet dog, horse sweat)<sup>1-3</sup>.

Many spoilage bacteria also come into the winery on grapes. Sour rot and *Botrytis* increase the prevalence of *Acetobacter*, *Gluconobacter* and *Gluconacetobacter*, all of which produce acetic acid. Several *Lactobacillus* species (*hilgardii*, *plantarum*, *casei*) and *Pediococcus (damnosus)* are also residents of mature grapes. These can produce acetic acid, mousy flavor and biogenic amines (which have names like putrescine and cadaverine...). They may also produce polysaccharides that lead to ropy texture<sup>1,3,5</sup>.

In addition to outright spoilage, high levels of native flora may also cause nutrient depletion early in fermentations that limit nutrients available to *Saccharomyces*, potentially leading to stuck fermentations<sup>6,7</sup>. In a study of nutrient depletion by non-*Saccharomyces* yeast species, Medina et al (2012)<sup>6</sup> found that *Metchnikowia*, a non-*Saccharomyces* yeast strain present in potentially high numbers on grapes<sup>8</sup>, consumed YAN quickly within the first few days of fermentation. Mimicking what may be occurring in fermentations with cold soaking or delayed inoculation, sequential inoculation with *Metchnikowia* followed by *Saccharomyces* led to stuck fermentations that could be resolved with nutrient addition. In the same study, *Hanseniaspora*, another prevalent member of the grape microbiota, did not show large YAN depletion (90% of the YAN remained 3 days after inoculation with this species), however it did deplete thiamine, an essential vitamin for *Saccharomyces*. Excessive use of SO<sub>2</sub>, as would occur in vintages with high microbial load, also leads to reduction in thiamine, further increasing the potential for stuck fermentations.

Despite the risks, there are also some benefits to having a rich microbial community early in fermentation. Several non-*Saccharomyces* yeast species have been shown to produce positive compounds that add complexity to wine aroma such as esters, higher alcohols, glycerol, succinic acid and thiols. Proteases produced by non-*Saccharomyces* yeast have been shown to break down cells and add nutrients, ultimately making a more protein stable wine. Some produce glycosidases that help unmask aromas compounds that are bound to sugar molecules. Others produce enzymes to break down polysaccharides that would otherwise inhibit clarification and filtration. *Lachanacea thermotolerans* has been shown to consume acetic acid, reducing volatile acidity<sup>1,8,9</sup>. It is likely these are some of the mechanisms that occasionally lead winemakers to employ ambient fermentations.

Many winemaking decisions affect the abundance and diversity of the microbial community present at the beginning of fermentation. Mechanical harvesting and long transport times, especially at warm temperatures, can lead to a high microbial load<sup>8,10</sup>. As soon as the grapes are crushed, nutrients are released to feed the organisms that are present. *Klocker*

(*Hanseniaspora*) is often the most abundant species on the grapes, and remains prevalent until alcohol levels rise above 4-7% and oxygen is used up<sup>3,10</sup>. The low pH environment of the juice, rising alcohol, rising temperatures, and presence of phenolics tend to inhibit spoilage organisms in early fermentation. Harvesting wet grapes, prolonged cold soak, cool fermentation conditions, low inoculant of yeast, and lack of clarification (for white wines) can all lead to higher counts of yeast and bacteria in the fermentation<sup>3,11</sup>

In wet vintages such as 2018, the prevalence of damaged berries and wet grapes likely increased the inoculant of non-*Saccharomyces* microbes in fermentation and may have contributed to overall higher volatile acidity in wines that year. One approach to microbial management is to use higher than normal levels of SO<sub>2</sub>. Though SO<sub>2</sub> has efficacy against some microbial spoilage, many of these microbes (such as *Hanseniaspora*) have high tolerance to it. Much of the SO<sub>2</sub> added at crush is lost as it binds to grape solids that are prevalent in red wine fermentations, making it less effective. High SO<sub>2</sub> additions can also bind thiamine and slow down or halt fermentation, and may even select for SO<sub>2</sub> tolerant microbes that will cause spoilage during aging<sup>12,13</sup>. Still, fermentations that have some SO<sub>2</sub> added at crush do tend to have faster onset of fermentation (leading to lower potential for spoilage) and steadier kinetics (Egli et al 1998).

When SO<sub>2</sub> isn't enough, or when you want to limit SO<sub>2</sub> for other reasons, another option for combatting microbial spoilage during fermentation is chitosan. Chitosan is a naturally occurring molecule that can also be produced by the de-acetylation of chitin using NaOH or chitinase enzymes<sup>14</sup>. Chitin is the second-most common polymer found in nature (after cellulose), making up the cell walls of fungi and shells of crustaceans and thus readily available as a renewable resource<sup>5</sup>. In winemaking applications, chitin from *Aspergillus niger* is used as the source for chitosan. The effectiveness of any given formulation of chitosan depends on its molecular weight, deacetylation degree and the pH of the medium<sup>15</sup>. Lower molecular weight, higher degree of deacetylation is the favored formulation for antimicrobial purposes<sup>16</sup>. At juice and wine pH, chitosan is very positively charged, which increases efficacy<sup>16</sup>.

Different microbes bind chitosan to different degrees. Chitosan binding to cell walls is driven by chemical properties of the cell wall itself, with high degree of correlation to the hydrophilicity of the wall<sup>17</sup>. Gram negative bacteria are more susceptible to binding than gram positive<sup>17</sup>. Chitosan has been shown to have some efficacy against a wide range of grapevine and wine microbes including downy mildew<sup>18</sup>, powdery mildew<sup>19</sup>, *Phomopsis*<sup>20</sup>, *Lactobacillus*, *Oenococcus*, and *Brettanomyces*<sup>17</sup>. Due to its versatility as an antimicrobial agent, chitosan in various forms has been used worldwide at nearly every stage of wine production including vineyard applications, on grapes during transport and storage, at crush, after fermentation and during the aging of wine<sup>16,21,22</sup>.

Many different mechanisms of antimicrobial action have been proposed for chitosan in wine. Chitosan has been shown to physically bind to the cell walls of microbes<sup>15,17</sup>. Binding may

aid in sedimentation, leading to reduction in overall cell number with racking. Binding of the positively charged chitosan may also disrupt cell membranes, leading to leakage of ATP, potassium, and proteins, all essential components of cell function<sup>15,17</sup>. Leakage may therefore result in semi-viable cells or cell death. Other proposed mechanisms include the physical blocking of cell permeability by chitosan binding, chelation of survival factors such as copper, penetration of the cell membrane and binding to DNA<sup>15</sup>.

Regardless of the mechanism, chitosan has been shown to be quite effective in treating existing microbial infection of wine. When treating wine already inoculated with *Brettanomyces bruxellensis*, Taillander et al (2014) found that 85% of the population of *Brettanomyces* cells were dead within 20 hours of treatment. There was a dose effect in the rate of cell death, with 0.4 g/L treatment killing cells faster than 0.04 g/L. They also found differences in susceptibility based on the strain of *Brettanomyces* used, presumably due to genetic differences in cell wall components. Many cells initially compromised by chitosan treatment recovered, as evidenced by growth in the *Brettanomyces* population after 7 days. This rebound effect is a good argument for racking after treating an infection with chitosan. Other researchers tested the effect of 0.04 g/L chitosan on aging wine and found that, even at low dose, aging of wine on chitosan helped prevent infection by *Brettanomyces*<sup>23</sup>. Here, wine was not racked.

Most chitosan products are recommended for use in finished wine. Stab Micro M (Enartis) is a chitosan-based product specially formulated for use on juice and must to reduce the activity of a wide range of microbes, leading to reduction in volatile acidity, sulfide defects, volatile phenols and production of other off-flavors. It can be used in conjunction with SO<sub>2</sub> or as a replacement, depending on the condition of grapes and preference of the winemaker<sup>16</sup>. In this experiment, a single harvest lot of Chambourcin was split so that half received treatment with chitosan while the other half did not. Both bins received 40 mg/L SO<sub>2</sub>. Resulting chemistry and microbial load were assessed post fermentation and after aging.

## Methods

Chambourcin was hand harvested into slotted macro bins and randomly assigned to one of two treatments. Each lot was destemmed and pumped through a must pump into a variable lid tank to be inoculated for fermentation. Both tanks received 40 ppm SO<sub>2</sub>. Stab Micro M was sprinkled in layers into the treatment at the de-stemmer/must pump at alternative times with SO<sub>2</sub>. Color Pro (100 mL/ton) and 40 g/hL FT Rouge Soft and were added to both tanks on the following day. Must was inoculated with 30 g/hL Premier Red yeast rehydrated in 30 g/hL Startup two days after processing. Fermentations were chaptalized to 25°Brix the third day after processing along with the addition of 24 g/hL Superfood, 10 g/hL Cherry Tannin and 25 g/hL Color Max. Additional nutrients (a total of 36 g/hL DAP) were added at morning and evening pumpovers on the fourth day of fermentation.

Wine was pressed after the completion of primary fermentation, 16 days after processing. After settling, both tanks were racked and transferred to barrel after inoculation with VP-1 bacteria. Tartaric acid (1.85 g/L) and SO<sub>2</sub> was added to each barrel after the completion of malolactic fermentation and maintained at a common target molecular SO<sub>2</sub> based on pH.

### Results

The initial juice chemistry for the two treatments was slightly different, with the Stab Micro lot having lower initial sugar and pH and higher TA (Table 1). This may be due to variation in the vineyard or inherent variation in sampling of non-fermenting red grapes. Fermentations were robust with no differences between treatments (Figure 1). Finished wine chemistry was also very similar with no difference in volatile acidity between the lots (Table 2, Figure 2). Likewise, the microbial load of wine after 4 months of barrel aging was very similar between lots (Table 3), with the Stab Micro treatment having only slightly lower levels of acetic acid bacteria. This treatment also had higher levels of some *Lactobacillus* microbes (Table 3).

Table 1: Juice chemistry for two treatments of Chambourcin (in-house data)

	Brix (deg)	pH	TA (g/L)	VA (g/L)	YAN (mg/L)
SO <sub>2</sub> only	22.1	3.67	5.8	0.18	271.64
Stab Micro	21.6	3.59	6.3	0.24	270.53

Figure 1: Fermentation kinetics for two treatments of Chambourcin (in-house data)

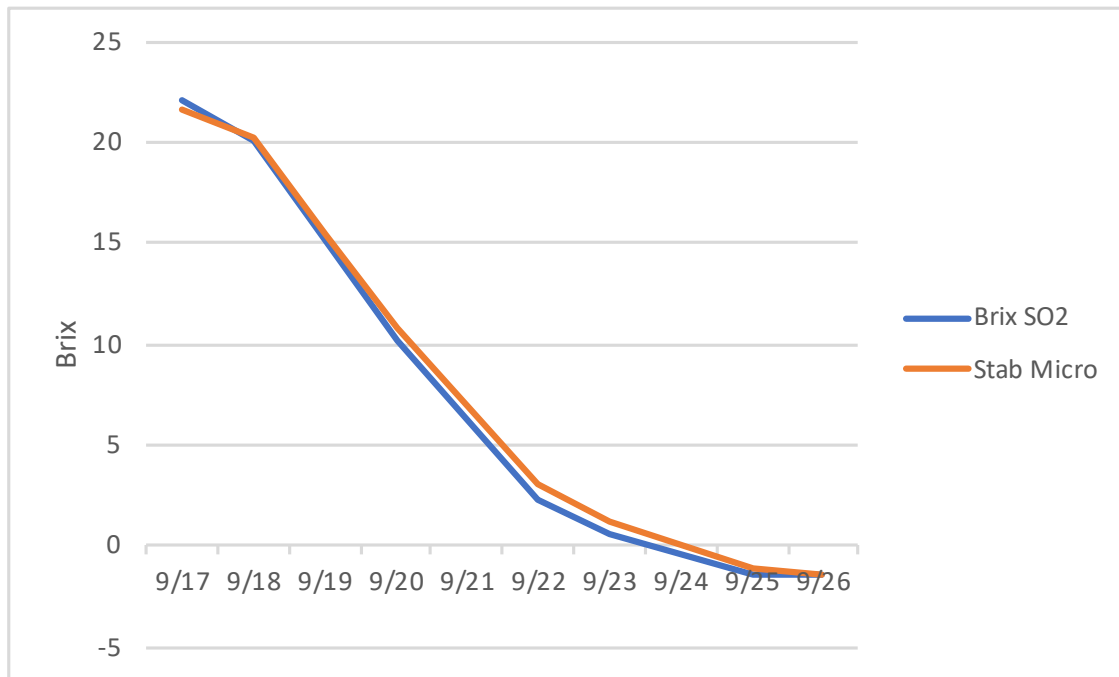


Table 2: Finished wine chemistry for two treatments of Chambourcin (ICV Labs)

	VA (g/L)	pH	TA (g/L)	Alcohol (%)	Glucose/Fructose (g/L)	Lactic Acid (g/L)
SO <sub>2</sub> only	0.65	3.66	5.1	13.67	2.9	1.6
Stab Micro	0.66	3.63	5.16	13.87	2.8	1.58

Figure 2: Color metrics for two treatments of Chambourcin (ICV labs)

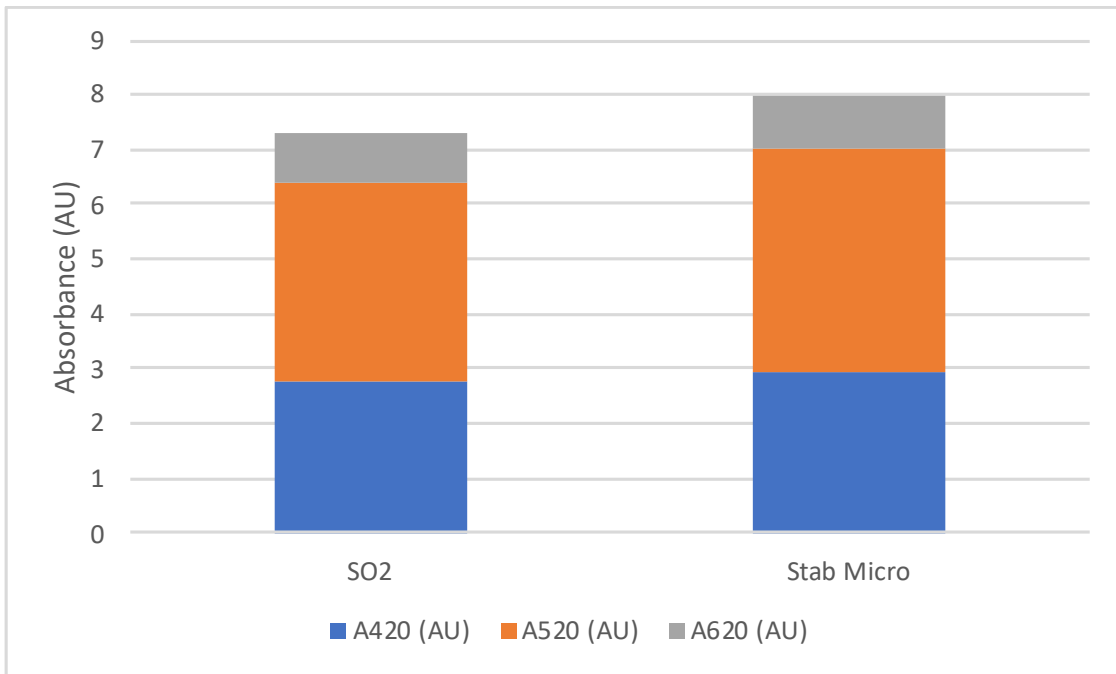


Table 3: Microbiology for two treatments of Chambourcin (ETS Labs)

Microbe	SO <sub>2</sub> only	Stab Micro
<i>Saccharomyces cerevisiae</i>	3900	2380
<i>Oenococcus oeni</i>	1 x 10 <sup>7</sup>	1 x 10 <sup>7</sup>
<i>Acetic Acid Bacteria</i>	8.8 x 10 <sup>4</sup>	6.4 x 10 <sup>4</sup>
<i>Brettanomyces bruxellensis</i>	0	0
<i>L. brevis/hilgardii/fermentum</i>	30	30
<i>Lactobacillus kunkeei</i>	1430	900
<i>Lactobacillus plantarum/casei/mali</i>	2.96 x 10 <sup>4</sup>	3.19 x 10 <sup>4</sup>
<i>Pediococcus Species</i>	10	0
<i>Zygosaccharomyces Species</i>	0	0

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