# Saccharomyces cerevisiae Bio-Ethanol Production as an Alternative Source of Sustainable Energy

Bakare V.<sup>1</sup>, Benjamin B.<sup>2</sup> and Effiong TE<sup>3</sup>

<sup>1</sup>Department of Biological Sciences, Nigeria Defence Academy, P.M.B. 2109 Kaduna, NIGERIA
<sup>2</sup>DNA LABS Kaduna, 2A Kinkino Road, Off. Sultant Road, U/Rimi-Kaduna, NIGERIA
<sup>3</sup>Department of Biological Sciences, Nigeria Defence Academy, P.M.B. 2109 Kaduna, NIGERIA

<sup>1</sup>Corresponding Author: benthaibenjamin@gmail.com

### ABSTRACT

Current world energy demand is based on fossil fuels, which will vanish in coming decades. Renewable energy especially biofuels has attracted great interest as solutions to the current energy problem. Among available biofuel resources, bioethanol seems to be an efficient alternative thus, Saccharomyces cerevisiae a well-established organism for bioethanol production. However, during fermentation process, yeast cells experience various stress conditions and inhibitors hampering its efficacy for commercial bioethanol production. To overcome these yeast cells, adopt different signal transduction pathways. In this review, common and least explored carbon feedstock which can be readily converted into bioethanol are highlighted. The various protectants, genes, and pathways which can be tempered to engineer yeast strains are discussed. Thus, we have suggested strategies to utilize this lucrative alternative for sustainable bioethanol production.

Keywords- Biofuels, Fermentation, Feedstock and Yeast

## I. INTRODUCTION

Human population has dramatically increased in the past decades, stretching the finite fossil fuels resources. Fossil fuels, coal, natural gas, oils account for 90% of total energy demand in the world. However, fossil fuels are limited and are major contributors of greenhouse gasses. Renewable energy is an alternative and among available resources, biofuels seem to be an efficient and sustainable energy. Current biofuels for bioethanol and biodiesel production are based on sugar crops. However, food versus fuel dilemma jeopardizes its long-term usage. Non-food crops (switchgrass, poplar, and willow), algae and Genetically Modified Organisms (GMOs) are other sources for biofuel production. Till date, yeast -Saccharomyces cerevisiae is considered to be ideal microorganism for ethanol fermentation (Lam et al., 2014). It can utilize various feedstock for bioethanol production which are discussed below.

### **Bioethanol Production from Carbohydrates**

Carbohydrates present in starch, cellulose, and hemicellulose are used for bioethanol production. Sugarcane juices, molasses, and corn are primary feedstock used worldwide for bioethanol (Wilkie, 2000). Starch, a polysaccharide of glucose is obtained from corn, barley, wheat, rye, potato, sorghum, and cassava. For the production of bioethanol starch-containing feedstock must be first converted to sugar or dextrin by an enzymatic process and often enzyme used is amylase. Other complex sugars are converted to simple sugars by Saccharification process and are fermented to ethanol (Naik, 2010).

The biofuel produced from starch, sugars, animal fats and vegetable oils are referred as 'first generation biofuel'. Food and fuel dilemma however, jeopardizes its large-scale commercial production.

In coming decades, world population is expected to be about 9 billion and around 2.5 billion more people will be added by 2050 (Godfray *et al.*, 2010), thus, hampering sustainability of food crops for biofuel production. Moreover, insufficient supply of these crops hampers its long-term usage and its commercialization. As an alternative, lignocelluloses feedstock 'second generation biofuel', can be used (Kumar, 2009). Lignocellulose includes agricultural waste (straw of rice, wheat, corn, and sugarcane bagasse), nonfood plants like poplar, napiergrass, switches grass, paper waste, agro-industrial waste, water hyacinth and sawdust (Yasuda *et al.*, 2014). Being non-food crops it seems to be sustainable energy resource.

### II. BIOETHANOL FROM AGRICULTURAL WASTE MATERIALS

Crops like corn, wheat, and sugarcane are primarily used for food. So, the sufficient production of these crops for fuel remains the major obstacle in the production of bioethanol (Cheng, 2011). The major agricultural wastes are straws of corn, wheat, and rice and sugarcane bagasse. These waste material don't have any nutritional value, easily available and are cheap. Moreover, it does not require separate agricultural land, water supply, fertilizers, and energy sources. Most of the agricultural waste materials are either left to rot in the field for composting or burnt in the fields. Rather than just disposing or burning these wastes it can be judiciously used as biomass for bioethanol production. Besides this feedstock like vegetable or fruit processing

# **International Journal for Research in Applied Sciences and Biotechnology**

#### www.ijrasb.com

wastes can also be used for bioethanol or biodiesel production.

#### Fermentation Stress Tolerance Mechanism

The yeast - S. cerevisiae is widely used in ethanol fermentation industry owing to its efficient conversion of sugars to ethanol (Fig. 1). However, during fermentation, it experiences numerous stress conditions. Stress conditions and an adaptive mechanism to overcome can be collectively called as 'Fermentation Stress Tolerance' (FST).

### Tolerance to Ethanol

cerevisiae ferments S. sugar, starch, lignocellulose to ethanol but when the ethanol https://doi.org/10.31033/ijrasb.7.6.27

accumulates above a threshold level, it inhibits growth, causes mitochondrial loss and eventually kills the yeast cells (Bai et al., 2004; Ibeas and Jimenez, 1997). Increased ethanol level affects membrane stability, damages protein and destroys cell membrane. There are several studies which have shown the major pathways and genes involved in ethanol stress tolerance. The knockout strains developed by You et al; showed tolerance to ethanol when supplemented with monounsaturated fatty acid (You KM, 2003). Inoue et al.(2000), demonstrated that strains lacking ergo sterol were sensitive to the moderate level of intracellular ethanol (Inoue et al., 2000).

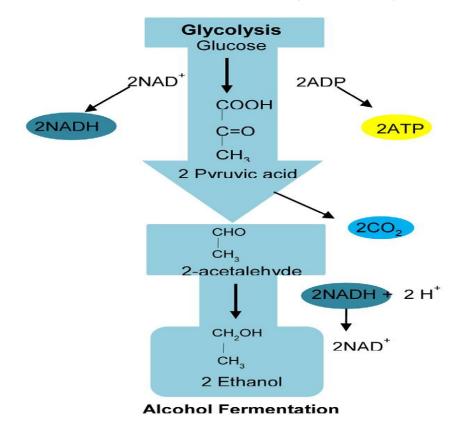


Figure 1: Ethanol fermentation (Inoue et al., 2000).

Fig.1. Ethanol fermentation: One molecule of glucose (C<sub>6</sub>H<sub>12</sub>O<sub>6</sub>) is converted into two molecules of pyruvic acid (C<sub>3</sub>H<sub>4</sub>O<sub>3</sub>) during the process of glycolysis. Pyruvic acid is further decarboxylated to generate two molecules of acetaldehyde (CH<sub>3</sub>CHO), which is reduced to ethanol (C<sub>2</sub>H<sub>5</sub>OH). During the process, there is a net gain of 2 molecules of ATP and one molecule of glucoseis converted to two molecules of ethanol and two molecules of carbon dioxide (CO<sub>2</sub>).

Yeast strains that over expressed genes like ARG4 and CAR1 responsible for the synthesis of arginine, showed to maintain the stability of cell wall and cell membrane (Cheng et al., 2016). Our group has also shown that over expression of RPI1 increases ethanol tolerance by over 50 fold compared to WT. RPI1 over

expression strain is highly resistant to cell wall lytic enzyme Zymolyase (Puria et al., 2009) suggesting; perhaps RPI1 improves the cell viability by strengthening the yeast cell wall.

A part from affecting plasma membrane, ethanol also denatures functional proteins and protein present in the cell membrane. In order to survive different environmental fluctuations, and to maintain the internal steady state homeostasis, cells have developed adaptive stress tolerance mechanisms. These cellular responses lead to change in gene expression and require signal transduction path ways to communicate from the sensors on the cell surface or cytoplasm to transcriptional machinery located in the nucleus to elicit a stress response, (Fig. 2) (Gasch and Werner-Washburne, 2002).

# International Journal for Research in Applied Sciences and Biotechnology

www.ijrasb.com

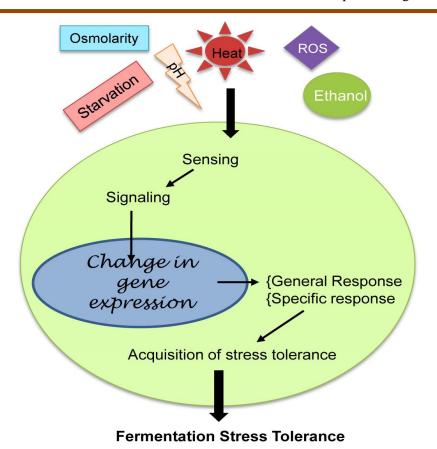


Figure 2: Fermentation Stress Tolerance Mechanism (Gasch and Werner-Washburne, 2002).

Fermentation Fig. 2 Stress Tolerance Mechanism: Yeast cells experience a plethora of stress conditions during fermentation namely high initial substrate concentration, nutrient deprivation, gradual accumulation of ethanol, the temperature rise of the fermentation medium, a decrease in pH, generation of reactive oxygen species (ROS). The cell sensing these signals by receptors over cell surface or intracellular receptors and transducer the signals for expression of certain genes under general or stress-responsive conditions. This expression of protective genes or detoxifier brings the adaptive stress responsive conditions called as fermentation stress tolerance.

### Ethanol Toxicity in Yeast

The cell wall of *S. cerevisiae* is made of about 85% polysaccharides and 15% proteins. The main functions of the cell wall are to stabilize osmotic homeostasis, protect cells against physical damage, maintain cell shape and act as a scaffold for glycoproteins (Klis *et al.*, 2006). The main targets of ethanol stress are yeast plasma membrane and hydrophilic and hydrophobic proteins (Stanley *et al.*, 2010). On exposure of yeast to ethanol, the fluidity of cell membrane increases and the stability of membrane decrease (Mishra, 1989). Thus, ethanol affects the structure and function of the cell membrane. Ethanol also denatures various proteins present in the plasma membrane. Ethanol concentration of 2-6% inhibits endocytosis across the plasma membrane

(Lucero *et al.*, 2000). Ethanol breaks proton motive force, which pumps protons across the plasma membrane (Cartwright, 1986). Exposure of ethanol to yeast cells affects the activity of Pma1 membrane protein, an H-AT Pase, necessary to preserve intracellular pH and membrane potential.

# III. GENETIC ENGINEERING TO IMPROVE YEAST STRAIN

### C6 and C5 Carbon Substrate

Yeast can efficiently metabolize glucose and have less affinity for other carbon sources such as galactose. Moreover, in presence of glucose other metabolic genes are repressed by a process called as 'glucose repression' (Le Borgne, 2011). In this regard various genetically engineered strains of S. cerevisiae have been developed. Improved galactose up take and yield of ethanol were obtained when the genes encoding phosphoglucomutase and positive regulator of Gal4p was overexpressed (Ostergaard et al., 2000). Overexpression of truncated TUP1 gene encoding repressor of transcription showed improved fermentation rates. Besides this lactose present in whey can be used to produce ethanol. But the commonly used S. cerevisiae for industrial ethanol production is not able to metabolize lactose.

# International Journal for Research in Applied Sciences and Biotechnology

### www.ijrasb.com

For metabolizing lactose Kluyveromyces fragilis (Guimaraes et al., 2008) or genetically engineered S. cerevisiae can be used. Starch rich materials are cheap and abundantly available; they can be used as feedstock for bioethanol productions. For fermentation of five carbon sources like xylose, yeasts with higher xylose fermentation rates like Pichia stipitis and Pachysolen tannophilus can be used (Jeffries, 1985; Jeffries et al., 2007). Xylose can be converted to xylulose by using enzyme xylose isomerase, xylose reductase and xylitol dehydrogenase (Klimacek et al., 2014). After conversion of xylose to xylulose, xylulose is phosphorylated to xylulose-5-phosphate and metabolized to ethanol. Klimacek et al. (2014) have developed an evolutionarily engineered strain of S. cerevisiae (IBB10B05) which can efficiently convert xvlose to ethanol (Klimacek et al., 2014). Konishi et al. (2015) developed a genetically modified strain of S. cerevisiae by using endogenous xylose digesting genes coding for sorbitol dehydrogenase, aldose reductase, and xylulose kinase to ferment xylose to ethanol (Konishi et al., 2015). Besides xylose, arabinose is also not utilized by yeast, for economical ethanol production from lignocelluloses feedstock; it has to be also channelized for ethanol production. Wisselink et al. (2007) developed a genetically engineered strain of S. cerevisiae expressing araA, araB and araD genes from bacteria Lactobacillus

*Saccharomyces cerevisiae* Bio-ethanol Production, A Sustainable Energy Alternative S205 plantarum to ferment arabinose to ethanol anaerobically (Wisselink *et al.*, 2007).

# IV. COMMERCIALIZATION AND FUTURE PROSPECTS

World energy consumption is increasing with the ever increasing population. Biofuels seem to be an efficient and sustainable energy resource. However, commercialization of biofuel is at infancy. The initial cost of investment, non-availability of arable land, seasonal nature of agricultural crops are some of the bottlenecks for commercialization. In this context algae considered to be the best option as it can thrive and profusely grow in non-arable land ranging from wasteland to aquatic ponds. Algal cell wall contents negligible lignin and intracellular stored starch granules can be readily converted to ethanol (Han et al., 2015). For the economical production of ethanol from algal biomass, it is necessary that all the carbohydrate content of the algal feedstock is converted to ethanol. Simultaneous saccharification and fermentation (SSF), a single bioreactor is an alternative for bioethanol production; it decreases method fermentation costs by reducing equipment requirements. If countries worldwide need to be self-sufficient and reduce the crude oil import then research should be focused on identifying improved harvesting and oil extraction processes, increasing the biomass of biofuel

https://doi.org/10.31033/ijrasb.7.6.27

crops. All of these challenges can be resolved by genetic, molecular, and ultimately synthetic biology techniques (Lee, 2010).

# V. CONCLUSION

Finally, looking into the near feature of renewable energy and demand and supply, more options needs to be explore as highlighted thus diversifying the sources. *Conflict of Interest* 

The authors contributed equally to the development of the manuscript and declare no conflict of interest.

# REFERENCES

Bai, F. W., Chen, L. J., Zhang, Z., Anderson, W. A., & Moo-Young, M. (2004). Continuous ethanol production and evaluation of yeast cell lysis and viability loss under very high gravity medium conditions. *Journal of Biotechnology*, *110*(3), 287–293. https://doi.org/10.1016/j.jbiotec.2004.01.017

[2] Cartwright, C. P., juroszek, J.-R., Beavan, M. J., Ruby, F. M. S., De Morais, S. M. F., & Rose, A. H. (1986). Ethanol Dissipates the Proton-motive Force across the Plasma Membrane of Saccharomyces cerevisiae. *Microbiology*, *132*(2), 369–377. https://doi.org/10.1099/00221287-132-2-369

[3] Cheng, J. J., & Timilsina, G. R. (2011). Status and barriers of advanced biofuel technologies: A review. *Renewable Energy*, *36*(12), 3541–3549. https://doi.org/10.1016/j.renene.2011.04.031

[4] Cheng, Y., Du, Z., Zhu, H., Guo, X., & He, X. (2016). Protective Effects of Arginine on Saccharomyces cerevisiae Against Ethanol Stress. *Scientific Reports*, *6*(1), 31311. https://doi.org/10.1038/srep31311

[5] Gasch, A. P., & Werner-Washburne, M. (2002). The genomics of yeast responses to environmental stress and starvation. *Functional & Integrative Genomics*, 2(4–5), 181–192. https://doi.org/10.1007/s10142-002-0058-2

[6] Godfray, H. C. J., Crute, I. R., Haddad, L., Lawrence, D., Muir, J. F., Nisbett, N., Pretty, J., Robinson, S., Toulmin, C., & Whiteley, R. (2010). The future of the global food system. *Philosophical Transactions of the Royal Society B: Biological Sciences*, 365(1554), 2769–2777.

https://doi.org/10.1098/rstb.2010.0180

[7] Guimarães, P. M. R., Teixeira, J. A., & Domingues, L. (2008). Fermentation of high concentrations of lactose to ethanol by engineered flocculent Saccharomyces cerevisiae. *Biotechnology Letters*, *30*(11), 1953–1958. https://doi.org/10.1007/s10529-008-9779-1

[8] Han, S.-F., Jin, W.-B., Tu, R.-J., & Wu, W.-M. (2015). Biofuel production from microalgae as feedstock: Current status and potential. *Critical Reviews in Biotechnology*, 35(2), 255–268. https://doi.org/10.3109/07388551.2013.835301

193

[9] Ibeas, J. I., & Jimenez, J. (1997). Mitochondrial DNA loss caused by ethanol in Saccharomyces flor yeasts. *Applied and Environmental Microbiology*, *63*(1), 7–12. https://doi.org/10.1128/AEM.63.1.7-12.1997

[10] Inoue, T., Iefuji, H., Fujii, T., Soga, H., & Satoh, K. (2000). Cloning and Characterization of a Gene Complementing the Mutation of an Ethanol-sensitive Mutant of *Sake* Yeast. *Bioscience, Biotechnology, and Biochemistry*, 64(2), 229–236. https://doi.org/10.1271/bbb.64.229

[11] Jeffries, T. (1985). Emerging technology for fermenting -xylose. *Trends in Biotechnology*, *3*(8), 208–212. https://doi.org/10.1016/0167-7799(85)90048-4

[12] Jeffries, T. W., Grigoriev, I. V., Grimwood, J., Laplaza, J. M., Aerts, A., Salamov, A., Schmutz, J., Lindquist, E., Dehal, P., Shapiro, H., Jin, Y.-S., Passoth, V., & Richardson, P. M. (2007). Genome sequence of the lignocellulose-bioconverting and xylose-fermenting yeast Pichia stipitis. *Nature Biotechnology*, *25*(3), 319–326. https://doi.org/10.1038/nbt1290

[13] Klimacek, M., Kirl, E., Krahulec, S., Longus, K., Novy, V., & Nidetzky, B. (2014). Stepwise metabolic adaption from pure metabolization to balanced anaerobic growth on xylose explored for recombinant Saccharomyces cerevisiae. *Microbial Cell Factories*, *13*(1), 37. https://doi.org/10.1186/1475-2859-13-37

[14] Klis, F. M., Boorsma, A., & De Groot, P. W. J. (2006). Cell wall construction inSaccharomyces cerevisiae. *Yeast*, 23(3), 185–202. https://doi.org/10.1002/yea.1349

[15] Konishi, J., Fukuda, A., Mutaguchi, K., & Uemura, T. (2015). Xylose fermentation by Saccharomyces cerevisiae using endogenous xylose-assimilating genes. *Biotechnology Letters*, 37(8), 1623–1630. https://doi.org/10.1007/s10529-015-1840-2

[16] Kumar, P., Barrett, D. M., Delwiche, M. J., & Stroeve, P. (2009). Methods for Pretreatment of Lignocellulosic Biomass for Efficient Hydrolysis and Biofuel Production. *Industrial & Engineering Chemistry Research*, 48(8), 3713–3729. https://doi.org/10.1021/ie801542g

[17] Lam, F. H., Ghaderi, A., Fink, G. R., & Stephanopoulos, G. (2014). Engineering alcohol tolerance in yeast. *Science*, *346*(6205), 71–75. https://doi.org/10.1126/science.1257859

[18] Le Borgne, S. (2012). Genetic Engineering of Industrial Strains of Saccharomyces cerevisiae. In A. Lorence (Ed.), *Recombinant Gene Expression* (Vol. 824, pp. 451–465). Humana Press. https://doi.org/10.1007/978-1-61779-433-9\_24

[19] Lee, D. H. (2011). Algal biodiesel economy and competition among bio-fuels. *Bioresource Technology*, *102*(1), 43–49.

https://doi.org/10.1016/j.biortech.2010.06.034

[20] Lucero, P., Peñalver, E., Moreno, E., & Lagunas, R. (2000). Internal Trehalose Protects Endocytosis from Inhibition by Ethanol in Saccharomyces cerevisiae. *Applied and Environmental Microbiology*, *66*(10), 4456– https://doi.org/10.31033/ijrasb.7.6.27

4461. https://doi.org/10.1128/AEM.66.10.4456-4461.2000

[21] Mishra, P., & Prasad, R. (1989). Relationship between ethanol tolerance and fatty acyl composition of Saccharomyces cerevisiae. *Applied Microbiology and Biotechnology*, 30(3).

https://doi.org/10.1007/BF00256221

[22] Ostergaard, S., Olsson, L., Johnston, M., & Nielsen, J. (2000). Increasing galactose consumption by Saccharomyces cerevisiae through metabolic engineering of the GAL gene regulatory network. *Nature Biotechnology*, *18*(12), 1283–1286. https://doi.org/10.1038/82400

[23] Puria, R., Mannan, M. A., Chopra-Dewasthaly, R., & Ganesan, K. (2009). Critical role of *RPI1* in the stress tolerance of yeast during ethanolic fermentation. *FEMS Yeast Research*, 9(8), 1161–1171. https://doi.org/10.1111/j.1567-1364.2009.00549.x

[24] Naik, S. N., Goud, V. V., Rout, P. K., & Dalai, A. K. (2010). Production of first and second generation biofuels: A comprehensive review. *Renewable and Sustainable Energy Reviews*, 14(2), 578–597. https://doi.org/10.1016/j.rser.2009.10.003

[25] Stanley, D., Bandara, A., Fraser, S., Chambers, P. J., & Stanley, G. A. (2010). The ethanol stress response and ethanol tolerance of *Saccharomyces cerevisiae*. *Journal of Applied Microbiology*. https://doi.org/10.1111/j.1365-2672.2009.04657.x

[26] Wilkie, A. C., Riedesel, K. J., & Owens, J. M. (2000). Stillage characterization and anaerobic treatment of ethanol stillage from conventional and cellulosic feedstocks. *Biomass and Bioenergy*, *19*(2), 63–102. https://doi.org/10.1016/S0961-9534(00)00017-9

[27] Wisselink, H. W., Toirkens, M. J., del Rosario Franco Berriel, M., Winkler, A. A., van Dijken, J. P., Pronk, J. T., & van Maris, A. J. A. (2007). Engineering of Saccharomyces cerevisiae for Efficient Anaerobic Alcoholic Fermentation of 1-Arabinose. *Applied and Environmental Microbiology*, 73(15), 4881–4891. https://doi.org/10.1128/AEM.00177-07

[28] Yasuda, M., Nagai, H., Takeo, K., Ishii, Y., & Ohta, K. (2014). Bio-ethanol production through simultaneous saccharification and co-fermentation (SSCF) of a low-moisture anhydrous ammonia (LMAA)-pretreated napiegrass (Pennisetum purpureum Schumach). *SpringerPlus*, *3*(1), 333. https://doi.org/10.1186/2193-1801-3-333

[29] You, K. M., Rosenfield, C.-L., & Knipple, D. C. (2003). Ethanol Tolerance in the Yeast Saccharomyces cerevisiae Is Dependent on Cellular Oleic Acid Content. *Applied and Environmental Microbiology*, *69*(3), 1499–1503. https://doi.org/10.1128/AEM.69.3.1499-1503.2003