Review Article

Current challenges in plant breeding to achieve zero hunger and overcome biotic and abiotic stresses induced by the global climate changes: A review

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Abstract

According Sustainable Development goals until 2030 we should have zero hunger and undernourished people in the world. But to achieve this goal plant breeders must improve plants in order to produce at least the double than is produced now. This is not an easy pathway because we have only few years, but considering that plant breeding programs normally take several years to produce improved genotypes, also the further improved plants should face with pest, disease and other abiotic factors that are increasing with the current climate changes. In this review we will discuss the situation of hunger in the world and the remaining available land to increase food production, point out effects of biotic and abiotic factors on the food production and present some ways that can be used to fastening plant breeding.

Introduction

Lately plant breeders have been warned that significant increase in agricultural production should be made to attend the increasing global demand for agricultural crops by the growing human population [1]. This will create the need to increase plant production in order to improve food security reducing the risk of hunger in the future. However, the increase of agricultural production is made even more difficult because of the reduction of available arable land and the current climate change and the environmental conditions where crops are grown. Until now, around 70% of the world land is currently used [2].

The projections of Tillman et al. [3] pointed out an increase of crop demand between 100 and 110% until 2050 considering the per capita consumption, and according to the projections of Ray, et al. [4] the annual increase in food production shows that it will be insufficient to attend the food demand in 2025 and in 2050 (Table 1). Therefore, extra available land will be necessary that yet doesn't exist, to cultivate the major crops and feed the world. All this is happening because the population in the world is increasing faster than crop production/area is achieved in the same time.

On the other hand, the last projections of the United Nations, DESA, Population Division, [5] pointed out that human growth will achieve 10 billion peoples in 2050 with continuous growth until the year 2100 when an equilibrium between birth and death is expected to happen (Figure 1). These continuous population increase makes crop genetic improvement of resilient species a priority [6].

Recent updates have shown that hunger in the world is slowly raising and that around 820 million people are nowadays feeling hunger, whereas the number of

Table 1: Mean of food production (ton/ha/year) according to Ray, et al. [4].

| Crop production | Yield in 2008 | Increase per year | Yield in 2025 | Required extra land | Yield in 2050 | Required extra land |
|-----------------|--------------|------------------|---------------|---------------------|---------------|---------------------|
| Maize           | 5.2 tons/ha/year | 84 kg/ha/year | 6.5 tons/ha/year | 15 millions ha | 8.6 tons/ha/year | 29 millions ha |
| Rice            | 4.4 tons/ha/year | 40 kg/ha/year | 4.9 tons/ha/year | 33 millions ha | 5.9 tons/ha/year | 67 millions ha |
| Wheat           | 3.1 tons/ha/year | 27 kg/ha/year | 3.4 tons/ha/year | 46 millions ha | 4.1 tons/ha/year | 95 millions ha |
| Soybean         | 2.4 tons/ha/year | 31 kg/ha/year | 3.0 tons/ha/year | 14 millions ha | 3.8 tons/ha/year | 28 millions ha |
levels which represent another important challenge in the future that until now was not considered.

The loss of 33% of the total production of maize, soybean, rice and wheat in the world is explained by climate variations [14]. These values are still important to take into consideration and deal with in plant breeding because abiotic stresses like high/low temperature, drought, salinity, between others, can offer big challenges to increase crop productivity and cultivation in several locations around the world. Currently, great advance was done in the knowledge of the genetic and physiological control of the abiotic resistance, with several genes identified in annual and perennial species [15], such as genes that control salin resistance [16] and drought tolerance [17]. Also, considerable advance was made in the knowledge of the epigenetic regulation in plant abiotic stress resistance [18], like salinity and drought resistance in plants [19].

On the other hand, according to Lucas, [20], losses produced worldwide, from 2001 to 2003, by weed, pests, pathogens and viruses were around 26.3% in soybean, 28% in wheat, 28.8% in cotton, 31.2% in maize, 37.4% in rice and 40.3% in potatoes. These values are considerable if we think that around one third of the total crop production will be lost. This persistent loss of crop production by pest and diseases is one of the major barriers to achieve global food sufficiency [21]. Fortunately, substantial advance was done in the knowledge of the genetic control of the development of biotic diseases and also in the strategies that can be used for their control [1].

Despite the advances in crop species disease control, there are still several crop species affected with pest and diseases producing considerable impact and reduction on plant production [1]. To survive the species need continuously to produce food and for other purposes (Clay, 2011). Besides this, there is also the need to produce food through sustainable agricultural practices that is already endangered by several environmental factors [11] to reduce environmental impacts over soils yet highly depreated [3]. So, to avoid hunger in the future some studies have shown that global food production should double by 2050 to be able to cover the increasing projected food demand [3].

According to Rodriguez, et al. [12], extreme temperature, drought and salinity are between the major abiotic factors responsible for reducing plant productivity. It is expected that the changes in the environmental and climate conditions will promote the appearance of new pest and diseases that will reduce plant resistance [1]. Another problem recently detected by De Storme and Geellen [13] is that high temperature can induce meiotic restitution of the chromosome set during the male gamete formation. One consequence of this problem reported is be the increase of polyploid plants and, consequently, these individuals will have reduced fertility
of these factors change, also the degree of the severity of the disease will change at individual and population levels [22]. So, for a disease to occur the pathogens need to overcome the host defense by using sabotage or by changing its appearance [25].

The knowledge about the genetic inheritance of the resistance to biotic stresses was first reported by R.H. Biffen around 116 years ago. He described for the first time the inheritance mode of resistance to the yellow wheat rust fungus observed in the F2 progeny by crossing susceptible and resistant lines [25]. After that, H.H. Flor described the segregation of flax resistance against Melampsora lini that allowed him to establish the gene-to-gene theory. Flor’s studies have shown that for each resistance gene (R) in the host there was a gene present in the pathogen (Avr) [22], so it was proposed the gene-for-gene theory and summarized in the “zig-zag model” to explain the molecular activity behind the resistance behavior [26-28]. This theory improved our knowledge of the biochemical and genetic basis of the plant-pathogen interaction which later allowed the identification and cloning of the R resistance genes [28]. The cloning and identification of these R genes have shown for several classes of plant pathogens that the cell death produced around the place where intracellular plant pathogens are hosted in the plant species with “hyper sensibility” resistance is produced by a receptor of molecules of the R plant-specific gene that is responsible for recognizing the Avr genes produced by the pathogen [22]. Therefore, we can say that the R genes are plant elicitors that recognize the pathogen genes and that pathogen genes are effectors or Pathogen Associated Molecular Patterns (PAMPs) that are responsible for the plant attack [1,24,25]. To overcome the host defense many pathogens use effectors that suppress the signaling pathways via sabotage and make it easy the infection process by deviating nutrients, and optimizing their living condition in the host plant [29].

Nowadays, several R genes are identified, cloned, modified and transferred to different crop species by using conventional breeding techniques, molecular Marker Assisted Selection (MAS) and/or biotechnological tools, like OMICS (genomics, transcriptomics, proteomics, metabolomics, etc.) in order to fight against pest and diseases and achieve durable resistance or tolerance by combining them with quantitative genes [28]. Thus, in smart agriculture the use of traditional crop breeding techniques is considered unsuitable to increase food production and attend the growing population by sustainable environment. Hence, tools in tissue culture techniques, like micropropagation [30], gametic embryogenesis [31,32], somatic embryogenesis, cell suspension, protoplast fusion make possible fast large scale cloning of high value plants, to produce pure lines [33], select biotic and abiotic resistant plants and improved varieties [34]. And the new tools in molecular genetics, like: OMICS [35], transgenics, gene editing [36,37] and MAS make possible to put together different biotic resistance genes to obtain durable resistant lines in a short time [11]. Also, interference RNA technology can be used to activate and silence genes in order to achieve the desired phenotype [38-40].

Not all abiotic factors can be reversed by plant breeding technologies. Some abiotic factors can be reduced by applying modern cultivation technologies like environmental management practices and link microbes to plants is it possible to promote biocatalyst against pathogens [41,42]. Also, the application of applying diethyl aminoethyl hexanoate, will increase plant resistance to abiotic stress, like cold [43,44] and salinity resistance [45]. And, using the proline aminoacid during plant acclimatization has shown to increase tolerance to salt stress [46] and to lower water condition [47].

Between the 17 goals of the Agenda for Sustainable Development adopted by the UN States members, the aim of the goal 2 is to achieve zero hunger, finish all malnutrition and double the crop productivity until 2030 [48]. To achieve this goal in the few years that remain to increase crop productivity, plant breeders need to deal hard not only producing improved genotypes for enhancing food production, but also producing resistant genotypes for the current and foreseen biotic and abiotic factors that the current change in climate will bring.

**So, what can we do to achieve the goal 2 of Sustainable Development?**

One of the biggest challenges to plant breeders’ scientists is to achieve novel solutions to increase food production in the current climate change [21]. So, in our opinion to reach the goal 2, plant breeders need firstly to reduce the impact of climate change on crop production by developing improved varieties genetically resilient or tolerant to biotic and abiotic factors [49], like drought and high temperature which will exert strong effect over the developmental time of plant life-cycle (Snowdon, et al. 2020) and affect greatly the plant-pathogen interactions [21].

Once several R resistance genes to biotic and abiotic stresses are known and available on genome data banks, such as NCBI, between others, plant breeders can use these sequences to design primers and use as molecular markers to search and identify species-specific resistance genes in the germplasm banks or at germplasm collections [50,51]. Identified resistant plants can be used to produce pure lines by traditional breeding approach, with subsequent selfing along 7 generations, or to make it faster by using in vitro gamete embryogenesis. The use of gametic embryogenesis is interesting because it allows producing in only one generation completely homozygous plants, reducing time and cost in pure line production. These pure lines can be used in crossings to produce hybrids by traditional breeding, reverse breeding, gene editing and make transgenic plants. Gametic embryogenesis is also interesting to induce mutation and produce transgenic plants before doubling the chromosome
number, so we will avoid hemizygosis, reducing time and costs with evaluation and conventional crossing to obtain homozygous plants. The time can also be reduced by using in vitro selection of resistant plants to biotic and abiotic factors, by using somatic embryogenesis and gamete fusion [34] to produce elite lines in a shorter time [52].

Genetic introgression also can be made by crossing wild relatives with elite plants and applying recurrent selection in order to increase the elite genome in resistant plants. If the resistance gene is found in the cytoplasm of the wild species it should be used as mother plant in the crossing and recurrent selection in order to conserve the resistance gene, otherwise it is independent. This process can be accelerated by using assisted selection with molecular markers. On the other hand, the resistance gene can be transferred by genetic engineering to the desired crop.

The integration of in vitro techniques, like gametic embryogenesis [31,32], plant cell and tissue culture, somatic embryogenesis, protoplast fusion [33] and mutagenesis to the modern biotechnology techniques, like: transgenic plants, synthetic biology, gene editing [6], interference RNAs [38-40] and other OMICS technologies can be used to speed the resistance in improved plants to biotic and abiotic factors and to modulate plant root depth or change plant architecture (Snowdon, et al. 2020). On the other hand, once susceptibility genes are identified, it is possible to use gene editing technologies in order to build resistant plants against different pathogens, and considering that QTLs are not race-specific and that we can combine them with R race-specific resistant genes to promote an effective and durable resistance in different environments [28], both strategies are highly recommended to deal with climate changes. Thus, according to Cobb, et al. [53], the public sectors should integrate new technologies to the Mendelian genetics and the principles of quantitative genetics in order to make vigorous change in crop productivity.

Another way to increase resistance is by using microbial biotechnology to enhance plant nutrition and/or promote biocontrol against pathogens [41,42], applying diethyl aminoethyl hexanoate to achieve resistance to abiotic stress [43-45], and use the prolineto increase salt tolerance [46] and low water condition [47].

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