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Trends in Plant Science



Q3 Spotlight

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- Drug Discovery forThirsty Crops
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Following virtual screening and structure-based ligand optimization, researchers have developed opabactin (OP), an abscisic acid (ABA)-receptor agonist with tenfold greater *in vivo* activity than ABA. This new ligand surpasses previous agonists for its potency and bioactivity on staple crops. OP leads a new class of agrochemicals designed to protect crops from drought.

ABA-Like Drugs

Among stresses, drought has the strongest impact on crop productivity and is occurring more frequently and intensely in a changing climate. ABA regulates plant growth and development and is crucial for adaptation to environmental stresses, including drought. In plants, ABA is perceived by the PYR/ PYL/RCAR family of ABA receptors that comprises 14 members in arabidopsis (Arabidopsis thaliana). PYR/PYL is the largest family of plant hormone receptors and is classified into three different subfamilies: subfamily I (PYL7-10), subfamily II (PYL4-6 and PYL11/12), and subfamily III (PYR1 and PYL1-3). Chemical compounds capable of activating ABA receptors (i.e., ABA-receptor agonists) hold promise for agriculture because their application could reduce yield losses due to drought. Thanks to the abundant crystallographic data gathered on ABA receptors, synthetic ligands found in chemical screenings can now be optimized to efficiently bind and activate PYR/PYL receptors. ABA's coordination in ternary complex with PYR/PYL receptors and PP2C co-receptors involves a network of water-mediated hydrogen

bonds and hydrophobic and electrophilic interactions. The 'Trp-lock' stabilizes ternary PYL-ABA-PP2C complexes by a series of hydrogen bonds that engage ABA's ketone with the gate and latch loops of the receptor and a key tryptophan residue of PP2C (Figure 1A). Additionally, a salt bridge links ABA's carboxylate with a conserved lysine of the receptor (Lys⁵⁹ in PYR1) essential for ABA binding (Figure 1A) [1].

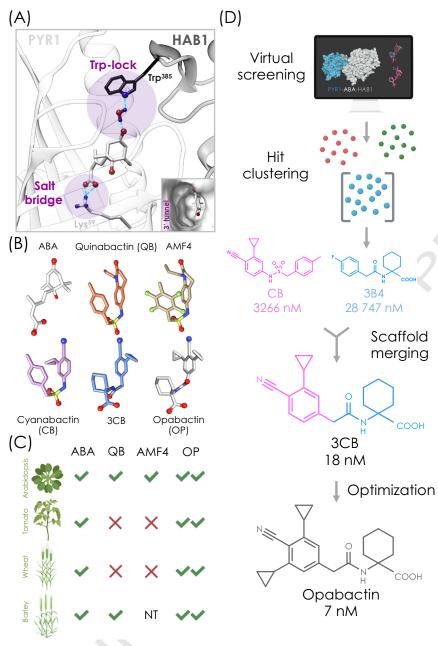
The sulfonamide quinabactin (QB) was the first synthetic ligand able to improve drought tolerance in plants [2]. Mutant analysis indicates that QB activates only the subset of dimeric ABA receptors PYR1, PYL1, and PYL2 in vivo. At the structural level, QB and ABA do not look alike (Figure 1B), but QB is able to fit into the receptor's pocket and mimic an important set of hydrophobic interactions and hydrogen bonds established by ABA [2]. Using crystallographic data on QB ternary complexes, several QB derivatives have been developed. AMF4 is a straightforward modification of QB where four fluorine atoms have been appended to the methylphenyl group, increasing the in vivo activity of the molecule and its persistence (Figure 1B) [3]. In subsequent work, QB's dihydroquinolinone was replaced by a benzyl ring decorated with a nitrile and a cyclopropyl moiety to generate cyanabactin (CB) (Figure 1B) [4]. CB's cyano group serves as hydrogen acceptor and engages the Trp-lock. CB's cyclopropyl group occupies the 3'tunnel, a hydrophobic cavity that interacts with ABA's 7' methyl group (Figure 1A) [5]. The interaction with the 3'tunnel increases CB's affinity for dimeric receptors [4].

Similar to the development of sulfonamidebased agonists like QB, the synthesis of ABA structural analogs has been instrumental in understanding ABA's structureactivity relationship [6]. Tetralone-ABA (tABA), where the vinyl methyl portion of ABA has been replaced with an aromatic ring, exhibits good *in vitro* activity and is likely to have a longer *in vivo* half-life than 49 ABA because it is unable to cyclize to the 50 catabolite phaseic acid [7].

From the First in Class to the Best in Class

ABA signaling is one of the most interesting 54 targets to improve plant drought tolerance. 55 Dozens of ABA-receptor agonists have 56 been protected with patents. However, 57 Vaidya et al. recently found that ABA-58 like molecules with sulfonamide linkers 59 (e.g., QB, AMF4, CB) show low activity on 60 important crops like wheat (Triticum aestivum) 61 and tomato (Solanum lycopersicum) [8]. To 62 find novel scaffolds to develop potent and 63 broad-spectrum ABA agonists, Vaidya 64 et al. performed virtual screening on mil- 65 lions of compounds. Docking experiments 66 were set up to identify ligands that retain 67 interaction with the conserved Lys of the 68 receptor. This constraint helped to identify 69 a set of substituted phenyl-amides whose 70 carboxylate might form a salt bridge with 71 the key Lys of the receptor, overcoming 72 the limitation of the sulfonamide-based 73 ABA agonists described so far. This group 74 of phenyl-amides is specific towards PYL8- 75 like receptors [9]. However, they lack a 76 properly positioned hydrogen acceptor to 77 interact with the Trp-lock water, resulting in 78 low agonist activity on family II and III recep- 79 tors [9]. The best molecule of this amide 80 group, 3B4 (Figure 1D), shows sub-81 micromolar activity towards subfamily I re- 82 ceptors but has poor IC₅₀ values on dimeric 83 receptors. To increase the ligand potency, 84 Vaidya et al. made use of a medicinal 85 chemistry trick. Playing with the compounds 86 as with LEGO® pieces, they merged the 87 amide of 3B4 with the cyclopropylphenyl 88 group of CB (Figure 1D). Thus, the 3B4's 89 carboxylate provides the key interaction 90 with the conserved Lys, while the cyano 91 group provides the interaction with the Trp- 92 lock water. This scaffold-merging exercise 93 resulted in a chimeric 3B4-CB hybrid, 94 named 3CB, a synthetic pan-agonist with 95 nanomolar activity for all arabidopsis and 96 wheat receptors tested [8,9].





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Figure 1. (A) Structure of the ternary complex PYR1-ABA-HAB1 (PDB: 3QN1) highlighting the 'Trp-lock' and the salt bridge. The 3'tunnel of PYL1 (PDB: 3JRS) is also depicted. (B) 3D structures of various abscisic acid (ABA)-receptor agonists. The opabactin (OP) 3D structure was obtained by docking using Maestro. (C) In vivo activity of ABA, quinabactin (QB), AMF4, and OP in various plant species. The in vivo potency of the different compounds is indicated with green ticks or red crosses if the compound was found not active (NT, not tested). (D) Scheme of the discovery process for OP indicating the IC_{50} values of the various molecules obtained through in vitro PP2C assays using PYL2 and HAB1 [8]. Some elements of this figure were created using BioRender (https://biorender.com).

own, structural analysis of the PYL10-

While 3CB is an exceptional agonist on its improved even further. In contrast to CB, 3CB's cyclopropyl group was not oriented 3CB complex suggested that it could be towards the 3'tunnel. To optimize 3CB, the

authors introduced a second cyclopropyl 101 substituent. In the newly synthesized mole- 102 cule, the second cyclopropyl is oriented 103 towards the 3'tunnel and improves the 104 activity even further, becoming the most 105 active ABA-receptor agonist described to 106 date. Using slang borrowed from video 107 gamers, the authors called this compound 108 'opabactin', for 'overpowered ABA recep- 109 tor activation'. This new compound is an 110 overpowered ligand with ten- and five- 111 times-stronger in vivo activity than ABA 112 in germination inhibition and stomatal 113 closure, respectively. Notably, the addition 114 of the second cyclopropyl group in OP re- 115 duces its in vitro activity on AtPYL8 but 116 increases it sixfold on TaPYL8, despite 117 having stronger in vivo activity than 3CB in 118 both plant species. 119

OP has strong in vitro activity over family II 120 and III ABA receptors in both arabidopsis 121 and wheat. However, the exceptional activity 122 of OP on dimeric receptors, five- to tenfold 123 higher than ABA, might be responsible for 124 the 'overpower' of OP. Genetic analysis 125 in arabidopsis revealed that OP's in vivo 126 activity is due to the activation of dimeric 127 PYR1, PYL1, and, especially, PYL2, con-128 firming the relevance of these receptors in 129 seed germination and stomatal closure 130 [2,8,10]. Furthermore, isothermal titration 131 calorimetry (ITC) experiments demonstrated 132 that the binding of OP to ABA receptors 133 is enthalpically driven, a characteristic com- 134 mon to best-in-class drugs [11]. Impor- 135 tantly, OP is able to activate stomatal 136 closure and to reduce transpiration not 137 only in arabidopsis but also in tomato, 138 wheat, and barley (Hordeum vulgare) 139 (Figure 1C). After the discovery of QB, the 140 first-in-class ABA-receptor agonist able to 141 improve drought tolerance, OP is currently 142 the best-in-class synthetic antitranspirant.

Still, Challenges Remain

QB was discovered 7 years ago. However, 145 the low activity of QB on staple crops like 146 wheat was not reported until recently [8]. 147 This highlights the importance of extending 148

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the characterization of ABA-receptor ago-149 nists from arabidopsis to crops or into 150 monocot crop models like Setaria viridis or 151 Brachypodium distachyon, closely related 152 to staple crops with C4 (maize, sorghum, 153 etc.) or C3 (wheat, rice, etc.) photosynthe-154 sis. Efforts in this direction have only re-155 cently started to be made [12]. However, 156 157 data obtained in laboratory setups will need to be confirmed in field trials to fully un-158 derstand the benefit of antitranspirants 159 under field conditions. We also propose 160 that understanding the chemical and ge-161 netic determinants for the bioactivity of these synthetic ligands, in different plant 163 species, will help in the development of the 164 next generation of antitranspirants. Addi-165 tionally, the combination of synthetic ligands 166 with plants expressing engineered recep-167 tors represents another layer of improve-168 169 ment to increase ligand potency and crop productivity while reducing agrochemical 170 input, making this alternative more environ-171 mentally friendly [3]. 172

The development of OP, an ABA-receptor 173 agonist with greater potency than the 174

endogenous hormone ABA, is a compelling example of the powerful combination of medicinal chemistry and plant biology and an exceptional advance in our mission to improve plant performance under stress conditions to improve global food security.

Acknowledgments

We thank Dr Laetitia Poidevin (IBMCP) and Jessica Toth (UCR) for comments on the manuscript. The editor and four anonymous reviewers are also acknowledged for their constructive suggestions. We apologize to authors whose work could not be cited due to space limitations. We also acknowledge Universidad Politécnica de Valencia for the grant 7. SP20180340 (PAID-06-18) to J.L-J. and Ministerio de Ciencia, Innovación y Universidades for the grant RTC-2017-6019-2 to P.L.R.

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https://doi.org/10.1016/j.tplants.2020.07.001

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