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Genetic diversity of almond rootstocks. The INRA *Prunus* rootstock breeding program

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Abstract. Almond and peach seedlings and peach*almond hybrids are still the most grown almond rootstocks in the Mediterranean countries. However a lot of new almond rootstocks were released in the last years, with different vigour, disease and nematode resistance, tolerance to abiotic stress. The almond rootstock choose is become more important, but there are opportunities for new rootstocks adapted to certain soil types like heavy soils or adapted to new high density systems. It could be possible to create them by interspecific crosses between several *Prunus* species, checking first the graft-compatibility with almond varieties. The main objective of the Breeding rootstock program is to pyramid three root-knot nematode resistance genes by interspecific crosses almond*peach*myrobalan.

Keywords. Almond rootstock – Root-knot nematodes – Resistance gene – *Prunus* species.

Diversité génétique des porte-greffes de l'amandier. Le programme d'amélioration des porte-greffes *Prunus* à l'INRA

Résumé. Les semis d'amandier et pêcher, ainsi que les hybrides pêcher*amandier sont toujours les porte-greffes les plus utilisés dans les pays méditerranéens. Cependant beaucoup de nouveaux porte-greffes ont été inscrits dans les dernières années avec différentes vigueurs, avec des résistances aux maladies et aux nématodes, tolérants aux stress abiotiques comme l'asphyxie racinaire ou des porte-greffes adaptés à des systèmes haute-densité. Il est possible de créer de nouveaux porte-greffes par croisements interspécifiques entre les différentes espèces *Prunus*, mais en vérifiant l'incompatibilité au greffage avec les variétés d'amandier. Le principal objectif du programme porte-greffe INRA est de pyramider 3 gènes de résistance aux nématodes à gales par croisement interspécifique amandier x pêcher x myrobalan.

Mots-clés. Intensité de floraison – Persistance – Germoplasme – *Prunus*.

I – The place of the of almond rootstocks in the Redher *Prunus* classification

Rootstock allows a good adaptation of the almond trees to the soil and can be prevent the tree decline against biotic and abiotic stress. Many *Prunus* species are used as rootstocks for apricot, peach or almond crops. The wide genetic variability within the *Prunus* genus allows to breed a high diversity of rootstocks for accumulating the characteristics by intraspecific or interspecific crosses. In the Redher *Prunus* classification, Fruit *Prunus* species are grouped in the three sections *Cerasus*, *Prunophora* and *Amygdalus* (Figure 1). For almond rootstocks. most of them belong to the section *Amygdalus* because the species of this section have good graft-compatibility with almond varieties. For calcareous soils, peach*almond hybrids are preferred to peach seedlings because they have better tolerance and a higher vigour. 'GF677' is still widely used, but the peach*almond hybrids like 'Hansen' and 'Garnem', resistant to root-knot nematodes replace 'GF677' in the infected soils. In the heavy soils, plum rootstock or amygdalus*prunophora hybrids are the best good alternative. They are more tolerant to root diseases such as *Armillaria* than peach and almond. In California, the prunus Marianna clone 'M2624' is the most planted, but now there are new *Amygdalus***Prunophora*

hybrids clones like 'Krymsk', 'Atlas', 'Viking' that would be replaced 'M2624'. The INRA hybrids 'Myran' and 'Ishtara' are not very planted, but have shown good results in heavy soils, like also for some slow growing plums from *domestica* and *insititia* species that are graft-compatible with almond.

Microcerasus species like *Prunus besseyi* are interesting genitors to obtain dwarfing rootstocks. The new rootstock Rootpac®20 is well adapted for the super high density orchards. The horticultural characteristics of the main commercial almond rootstocks are summarized in the Table 1.

II – The INRA *Prunus* Rootstock breeding program

A lot of *Prunus* rootstock are available to cover all soil types (cf Table 1) and several of them are resistant to RKN nematodes (Reighard and Lorette 2008), however in prevision of the climatic changes, in all rootstock breeding programs, the RKN resistance is a required trait to introduce and face the risk of breaking resistance, it is important to build temperature-stable resistances. Three RKN resistance genes have been identified in three *Prunus* species, *Ma* in the myrobalan plum (*Prunus cerasifera*, clone P2980 and P2175), *RMia* in the peach rootstock 'Nemared' and *RMja* in the almond 'Alnem' variety. So pyramiding these three genes by interspecific crosses almond*peach*myrobalan is the main objective of the breeding rootstock program.

However, to ensure the presence of the three genes in a same rootstock, it has been necessary to develop molecular markers to detect genes in the 3 way-hybrids. The identification of intra-gene markers for the two nematodes resistance genes *Ma* and *RMia* has allowed to apply marker assisted selection for these two genes (Claverie *et al.*, 2011, Duval *et al.*, 2014). A new F2 almond population of Lauranne x Alnem, with more 1000 hybrids has been created in order to clone the *RMja* resistance gene of 'Alnem' and to find intragenic molecular markers for this gene. The *RMja* gene is localized in the linkage group 7 of the *Prunus* genome in the same region than the *Ma* gene (Van Ghelder *et al.*, 2010). After that, it will be possible to select in the hybrid population 'Alnem*Nemared)*myrobalan', the pyramided hybrids with the three nematode resistance genes.

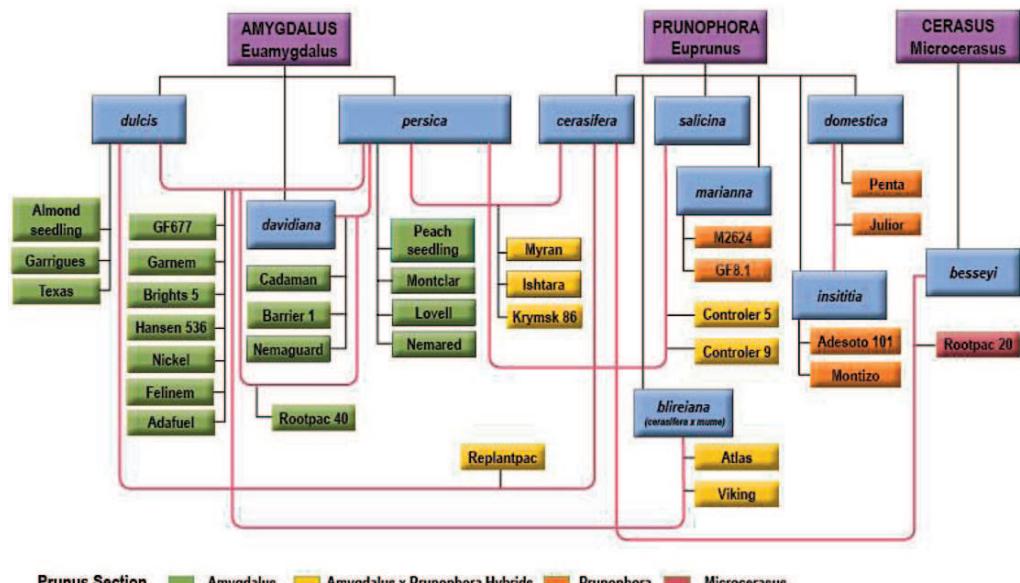


Fig. 1. Main almond rootstocks positioned in the Redher *Prunus* classification.

Table 1. Horticultural characteristics of commercial almond

Name	Rootstock	Vigor	Heavy soil	Chlorosis			Nematode		Armillaria
				RKN -Mi	RKN -Ma	RKN -Mj	Lesion		
Almond seedling	standard	poor	high	S	S	S	S	S	
Lovell	standard	fair	poor	S	S	S	S	S	
Nemared	standard	poor	poor	R	R	MS	S	S	
GF677	high	fair	high	S	S	S	MR	S	
Hansen 536	high	fair	high	R	R	MR	MR	S	
Brights 5	standard	fair	high	R	R	MR	MR	S	
Garnem (GN15)	high	fair	high	R	R	MS	MR	S	
Nemaguard	standard	fair	poor	R	R	MR	S	S	
Cadaman@avimag	high	fair	fair	R	R	MR	S	MS	
Barrier I	standard	fair	fair	R	R	MR	S	S	
Krymsk 86	semi-dwarf	good	poor	S	S	S	S	S	
Rootpac® R	standard	good	good	MR	MR	MR	S		
Ishtara®Ferciana	semi-dwarf	good	good	R	R	R	MS	MR	
Myran®Yumir	semi-dwarf	good	poor	R	R	R		MR	
Atlas	standard	poor	fair	R	R	R	S		
Viking	high	fair	fair	R	R	R	S		
Marianna 2624	semi-dwarf	good	good	R	R	R	S	R	
Penta	semi-dwarf	good	good	R	R	MR	S		
Julior®	semi-dwarf	good	fair	R	R	R	S		
Rootpac®20	dwarf	good	fair	R	R	R	R		

RKN : Root-Knot Nematode (Mi: Meloidogyne incognita, Ma: M arenaria, Mj: M javanica).

References

- Claverie M., Dirlewanger E., Bosselut N., Van Ghelder C., Voisin R., Kleinhenz M., Lafargue B., Abad P., Rosso M.N., Chalhoub B. and Esmenjaud, D., 2011. The Ma gene for complete-spectrum resistance to Meloidogyne species in Prunus is a TNL with a huge repeated C-terminal post-LRR region. In: *Plant Physiology*, 156, p. 779-792.
- Duval H., Hoerter M., Polidori J., Confolent C., Masse M., Moretti A., Van Ghelder C., Esmenjaud D., 2014. High-resolution mapping of the RMia gene for resistance to root-knot nematodes in peach *Tree Genetics & Genomes*, 10, p. 297-306.
- Reighard G.L. and Loreti F., 2008. Rootstock development. In: LAYNE, D. & BASSI, D. (Eds.) The peach: botany, production and uses.
- Van Ghelder C., Lafargue B., Dirlewanger E., Ouassa A., Voisin R., Polidori J., Kleinhenz M. and Esmenjaud, D., 2010. Characterization of the RMja gene for resistance to root-knot nematodes in almond: spectrum, location, and interest for Prunus breeding. In: *Tree Genetics & Genomes*, 6, p. 503-511.