

Interactions between pesticides and microorganisms: The case of biodegradation of synthetic β -triketone herbicides

C. Calvayrac¹, S. Romdhane^{1,2,3}, M. Devers-Lamrani³, F. Martin-Laurent³ and L. Barthelmebs¹

¹ Laboratoire BAE-LBBM, USR 3549 CNRS UPMC, 52 Avenue Paul Alduy, Perpignan, France.

² CRIOBE, USR 3278 CNRS EPHE, 58 Avenue Paul Alduy, Perpignan, France

³ INRA, UMR 1347 Agroécologie, Pole EcoDur, 17 rue Sully, BP 86510, Dijon, France

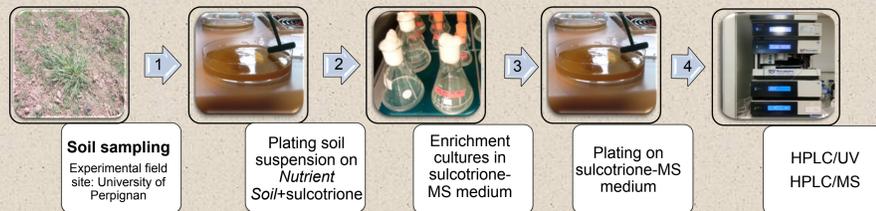
Introduction

Agricultural use of pesticides ensures a higher crop quality and production but it is also one of the major sources of diffuse pollution in the environment. Microbial degradation is considered as an important dissipation process limiting the accumulation of pesticides in the environment. In this context, two bacterial strains able to degrade sulcotrione, a β -triketone herbicide, were isolated from an agricultural soil previously exposed to this herbicide.

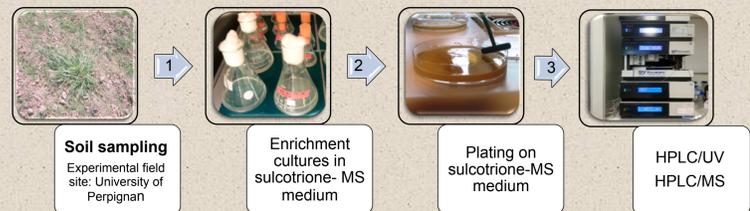


Bacterial isolation procedure: one soil, two strategies

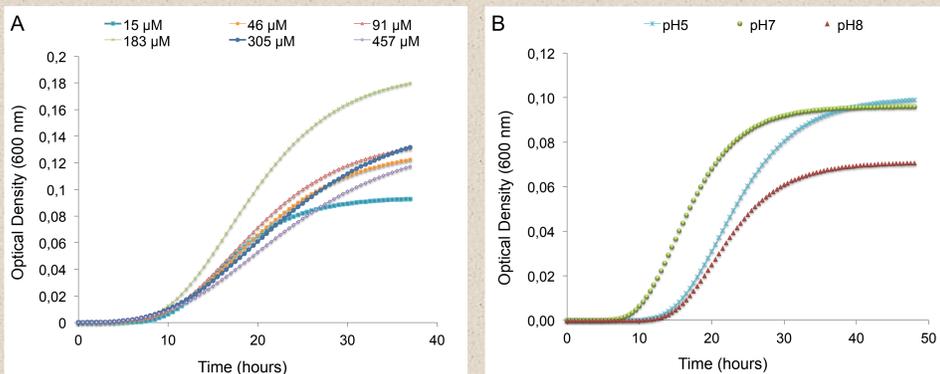
STRATEGY A



STRATEGY B

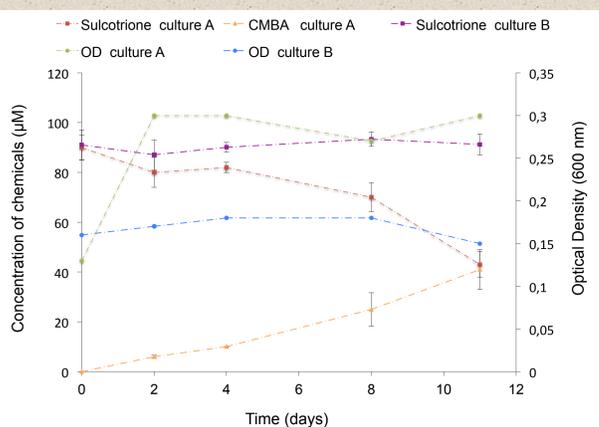


Characterization of *Pseudomonas* sp. 10P, a sulcotrione-degrading bacteria (Strategy A)



	Concentration of the herbicide in liquid medium	μ_{max} (h ⁻¹)	λ (h)	G (h)	OD _{max} (a.u.)
MSM + sulcotrione	15 μ M	0.0078 \pm 0.001	9	89.0	0.096 \pm 0.015
	46 μ M	0.0082 \pm 0.001	9	84.5	0.130 \pm 0.015
	91 μ M	0.0097 \pm 0.001	9	71.0	0.137 \pm 0.011
	183 μ M	0.0120 \pm 0.001	9	55.0	0.180 \pm 0.020
	305 μ M	0.0090 \pm 0.001	9	77.0	0.140 \pm 0.021
	457 μ M	0.0089 \pm 0.001	9	78.0	0.130 \pm 0.022
TS	—	0.1050 \pm 0.0024	5	7.0	—
	—	0.2779 \pm 0.0175	5	2.5	1.560 \pm 0.015

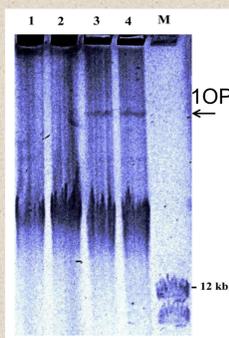
Growth parameters of the isolate *Pseudomonas* sp.10P. The maximum specific growth rate (μ_{max}), the lag time (λ), the generation time (G) and the maximum optical density (OD_{max}) values were determined for MSM+sulcotrione and TS media.



Degradation kinetics of sulcotrione, accumulation of CMBA and growth measurement of *Pseudomonas* sp.10P.

Culture A (MSM+sulcotrione) vs **Culture B** (10 repeated transfers from rich medium and then placed in MSM+sulcotrione).

No degradation was observed neither with CMBA nor tembotrione, another herbicide of the β -triketone family.

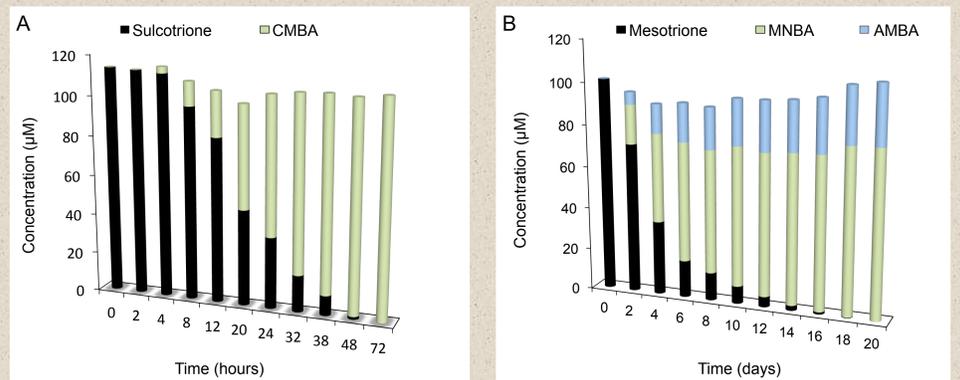


Genetic localization of sulcotrione degradation in *Pseudomonas* sp. 10P

Plasmid profiles from *Pseudomonas* sp. 10P obtained in MSM+sulcotrione or in absence of sulcotrione in a rich medium revealed the presence of a >12 kb plasmid. 10P plasmid was eliminated by curing experiment carried out on rich medium and sulcotrione-degrading ability was lost without pesticide selection pressure.

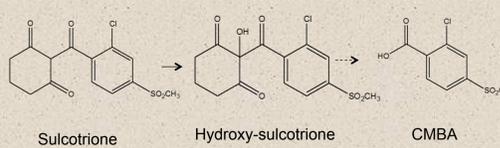
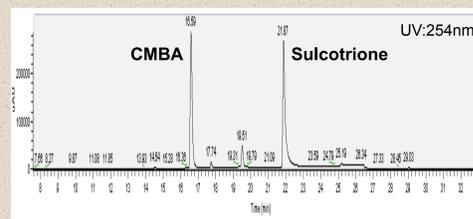
Lanes 1 and 2: strain cultivated in absence of sulcotrione in rich media. Lanes 3 and 4: strain cultivated in MSM in presence of sulcotrione (91 mM). Position of the plasmid is indicated by arrow.

Characterization of *Bradyrhizobium* sp. SR1, a sulcotrione and mesotrione degrading bacteria (Strategy B)



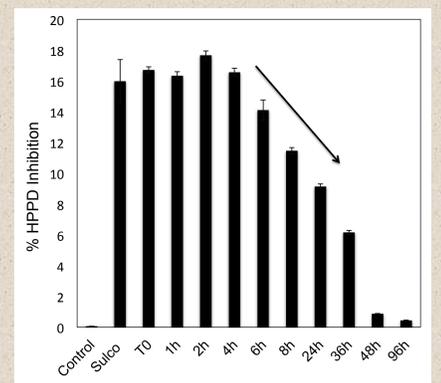
Degradation kinetics of (A) sulcotrione and (B) mesotrione by *Bradyrhizobium* sp. SR1 in resting cell experiments. Cumulative formation of metabolites during sulcotrione (CMBA) and mesotrione degradation is represented.

Study of sulcotrione metabolites



Concomitantly with the appearance of CMBA, a new metabolite in *Bradyrhizobium* sp. SR1 culture medium was detected and identified as hydroxy-sulcotrione, an intermediate metabolite of sulcotrione that finally gives CMBA.

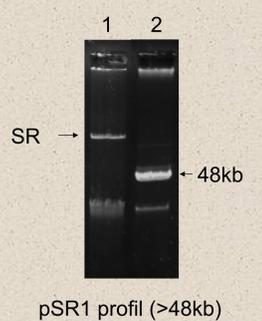
Microbial toxicity of sulcotrione and mesotrione metabolites



The toxicity of triketone molecules and their related metabolites was estimated by monitoring 4-hydroxyphenylpyruvate dioxygenase inhibition (HPPD). Toxicity due to HPPD inhibition was mainly linked to parent molecules, and not to the formed metabolites.

Genetic localization of sulcotrione degradation in *Bradyrhizobium* sp. SR1

Plasmid profiles from (1) *Bradyrhizobium* sp. SR1. The size of the plasmid is indicated in kb (2) *Sphingomonas* sp. SH., used as molecular weight marker. SR1 plasmid was not eliminated by curing experiment carried out on rich medium and that sulcotrione-degrading ability was maintained without pesticide selection pressure.



Conclusion: Two degrading strains exhibiting different genetic features

- *Pseudomonas* sp.10P was the first bacterial strain described in the literature as capable of degrading sulcotrione to 2-chloro-4-mesybenzoic acid (CMBA). Its growth performances have shown that initial neutral pH conditions and 183 μ M of sulcotrione seemed to be the best cultural environment for this strain. The isolate harboured a catabolic plasmid involved in sulcotrione biodegradation process.

- *Bradyrhizobium* sp. SR1 was able to biotransform two β -triketone herbicides, sulcotrione and mesotrione. The dissipation of sulcotrione and mesotrione led to the accumulation of different known metabolites, CMBA and MNBA/AMBA. A 14 000 Tn5 mutant library was constructed using a Tn5 mutagenesis approach conducted on *Bradyrhizobium* sp. SR1. Full sequencing of mutants are ongoing to identify possible degrading gene candidates.