1	Supporting Information for				
2	High permeation rates in liposome systems explain rapid glyphosate biodegradation				
3	associated with strong isotope fractionation				
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31 Supporting experimental section

32 Chemicals and media composition

The following chemicals were used in the liposome permeation study: glyphosate (98%, Sigma Aldrich), D₂O (99.9%, Sigma Aldrich), sodium hydroxide (97%, Fisher scientific), 1-palmitoyl-2-oleoylsn-glycero-3-phosphocholine (99%, Avanti Polar Lipids), praseodymium(III)-chloride (99.99%, Fisher scientific), and 3-(trimethylsilyl)-2,2,3,3-tetradeuteropropionic acid (98 atom % D, Sigma Aldrich).

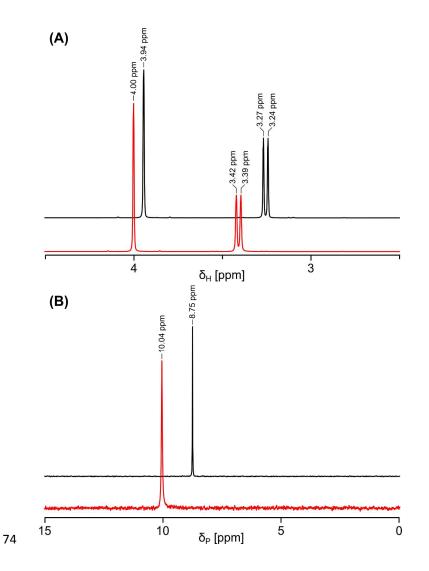
37 The following chemicals were used for the isolation of Ochrobactrum sp. FrEM and for glyphosate 38 degradation: glyphosate (99%, Sigma Aldrich), ammonium acetate (99%, Sigma Aldrich), sodium 39 glutamate (99%, Sigma-Aldrich), sodium peroxodisulfate (99%, Fluka), potassium hydroxide (99%, 40 Fluka), phosphoric acid (99%, Fluka), , potassium dihydrogenphosphate (99%, AppliChem), sodium 41 hydrogenphosphate (98%, Sigma Aldrich), sodium hydroxide (99%, Sigma Aldrich), potassium sulfate 42 (99%, Merck), ammonium chloride (99.5%, Sigma Aldrich), magnesium sulfate heptahydrate (99%, 43 Fluka), calcium chloride hexahydrate (94%, Roth), boric acid (99.8%, Merck), manganese sulfate 44 monohydrate (99%, Sigma Aldrich), zinc sulfate heptahydrate (99%, Fisher Scientific), nickel chloride 45 hexahydrate (99%, Merck) sodium molybdate dihydrate (99.5%, Merck), and iron sulfate 46 heptahydrate (98%, Sigma Aldrich).

The medium contained (in gL⁻¹): NH₄Cl, 2.0; MgSO₄ × 7 H₂O, 0.2; K₂SO₄, 0.5; as well as trace elements
(in mgL⁻¹): FeSO₄×7H₂O, 2.5; CaCl₂×6H₂O,10.0; CuSO₄×5H₂O, 2.0; H₃BO₃, 0.06; ZnSO₄ × 7 H₂O,20.0;
MnSO₄×H₂O, 1.0; NiCl₂×6H₂O, 0.05; Na₂MoO₄×2H₂O, 0.3.

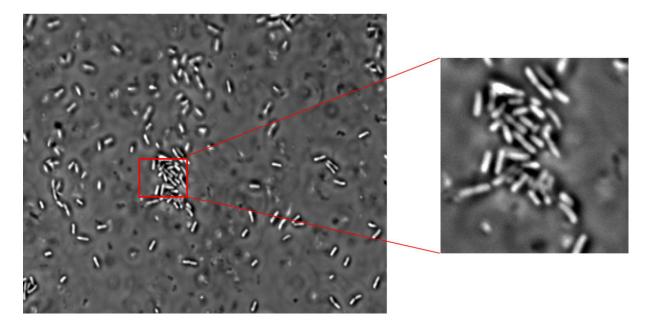
50 Bacterial isolation and characterization

The soil samples were collected from different plots (and later combined) on a vineyard site in northern France (Agricultural and Viticultural College of Rouffach - Rouffach soil) where glyphosate was the most frequently used herbicide with a yearly application of between 18 and 61 kg·ha⁻¹.¹ Soil samples from each plot location were thoroughly mixed in sterile bottles, sealed, transported to the laboratory and stored at 4 °C until use.

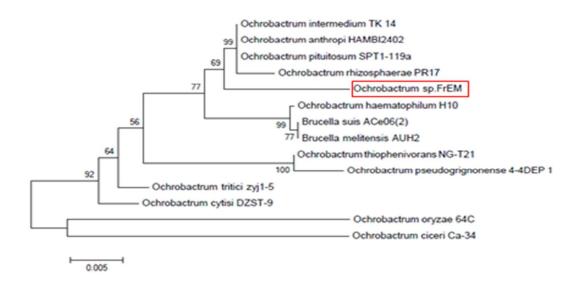
56 For bacterial isolation from soil, medium (see above) containing 60 mM sodium glutamate as carbon 57 source was used. Ammonium chloride was used as nitrogen source, and glyphosate was the sole 58 phosphorous source. Soil samples (10 g) were first sieved (> 2 mm). Then, 5 g of soil were suspended 59 in 10 mL sterile water and centrifuged. A 1 mL aliquot of the supernatant was used to inoculate 50 60 mL medium containing 3 mM glyphosate and incubated at 30 °C at 160 rpm for 24 h. Several 61 transfers were made and later streaked on agar plates containing 3 mM glyphosate. The single 62 colonies formed were inoculated on agar plate to represent the pure isolated strain. The isolated 63 bacteria were identified using 16S rRNA gene sequencing. For 16S rRNA gene amplification, the 64 chromosomal DNA was isolated using a bacterial DNA extraction kit (Roche Applied Science, 65 Germany) following the protocol of the manufacturer. PCR amplification was performed using 66 universal primers (forward 27f and reverse 1492r). Standard PCR conditions was carried out in a 50 67 μ L volume containing 1× PCR buffer, 1.5 mM MgCl₂, 2 mM dNTP mixture, 1 μ M primers, 1 μ M of Pfu 68 DNA polymerase (Fermentas, St. Leon-Rot, Germany), and 2 ng of template DNA. DNA was purified 69 from a gel using an Agarose Gel Extraction kit (Roche Applied Sciences, Germany) and sequenced. 70 Sequence homologies were evaluated using BLAST software (version 2.2.12). ClustalQ software was 71 used to align the sequences. A neighbour-joining tress was constructed using Molecular Evolution 72 Genetic Analysis (MEGA) software (version 6.0).



SI Figure S1. (A) ¹H with solvent suppression and (B) ³¹P{¹H} spectra of glyphosate without (black) and with addition of 1 mM PrCl₃ (red). The NMR spectra were obtained glyphosate in D₂O without pH adjustment where glyphosate dissociates into the monoanionic and the zwitterionic form.² The chemical shifts of glyphosate depend on pH. Therefore, the chemical shifts in these spectra differ from those presented in Figure 1 and Figure 2.

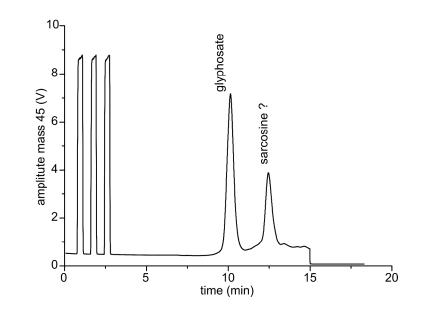


- 80
- 81 SI Figure S2. Micrograph of Ochrobactrum sp FrEM cells by light microscope (Axioskop Plus2, ZEISS,
- 82 Germany (×100 resolution oil emulsion), AxioVision 4.1



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SI Figure S3. Phylogenetic position of the strain FrEM within *Ochrobactrum* species. Neighbourjoining tree based on partial 16S rRNA sequence. The bar indicates 0.005 substitutions per nucleotide.



SI Figure S4. LC-IRMS chromatogram of ongoing glyphosate biodegradation. The three peaks between 0 s and 200 s are the CO₂ peaks of the reference gas. Glyphosate elutes around 10 min and is separated (resolution 2.3) from the peak around 12.5 min which could be the metabolite sarcosine. The CO₂ background increases after 11 minutes, probably due to overloading the column with the carbon source glutamate which elutes after 900 s when the split to the IRMS is already closed.

experiment	transmitter	relaxation	spectral	acquisition	number
	frequency	delay	width	time	of scans
	offset	(s)	(Hz)	(s)	
¹ H Standard	3165.1 Hz	1.0	9973.4	2.855	8
³¹ P{ ¹ H} Standard	10028.7 Hz	2.0	81521.7	0.99	16
¹ H with solvent suppression (watergate W5)	HOD freq.	1.5	7978.7	2.4009	128
${}^{1}\text{H}\{{}^{31}\text{P}\}$ with solvent suppression prior PrCl_{3} addition	HOD freq.	1.5	10000.0	1.499	16
$^{1}\text{H}\{^{31}\text{P}\}$ with solvent suppression after PrCl_{3} addition	HOD freq.	1.5	10000.0	1.499	64

SI Table S1.Summary of NMR spectra collection parameters

93 Supporting Information References

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 vineyard catchment. *International Journal of Environmental and Analytical Chemistry* 2010, *90*, (3-6),
 406-420.

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