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Estimating seed dispersal at the landscape scale : the case of feral oilseed rape populations

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Introduction

Cutifica

- European agro-ecosystems are mosaic of biodiversity compartments : (fields, margins, fallows, woods, roadsides ...) where genes flows occur by pollen and seeds flows. Seed dispersal is responsible for the escape of cultivated species from crops (i.e. feral populations) and therefore could induce genetic pollution and modifications of road verge plant communities.
- This could enhance GMHT plants introduction consequences on natural, semi-natural and cultivated areas by a higher ability of these plants to colonise habitats under herbicide pressure.

We built a seed flow model adapted to cultivated species based on mating models (Burczyk *et al*, 2006) to estimate seed dispersal



An agro-ecosystem : photography of Selommes area, June, 2009

Methods I

Data obtention

- Survey of an agro-ecosystem area of 41 km² around Selommes town and its silo
- GPS mapping of all oilseed rape fields and populations from 2000 to nowadays
- Sample collections of leaves and seeds from 2002 to nowadays
- A cultivar database from seed producers (58 cultivars)
- Samples and cultivars were genotyped with 8 microsatellites markers :



- distances of oilseed rape associated with different types of seed sources at the landscape scale.
- We used dispersal kernels which are particularly effective tools in modelling dispersal processes because they allow accounting for the rare but so important long-distance dispersal events (i.e. the tail of the kernel), which notably govern the spread rate of populations and gene flow at long distances.



Feral population of oilseed rape growing along a roadside

2125 sample genotypes (2002-2005, sub area) 1480 cultivar genotypes

Samples assignation to cultivars with Likelihood method (Devaux et al., 2005) :





Preliminary Results



$L(\alpha, ms, a, i) = ms \times BSP + (1 - ms) \times \left(\alpha \times C_{x, v}^{2004} + (1 - \alpha) \times C_{x, v}^{2005}\right)$

	alpha	ms	а	logL
2005-2004	1.0	0.54	39.1	- 457.3

ms: contribution of the background seed pool alpha : estimated weight of mature fields in 2005 C: field seed contribution

Likelihood of observing all feral populations of 2005 taking account of contributions from 2004 and 2005 fields and BSP



a : estimated mean dispersal distance *logL* : log-Likelihood

Maximum Likelihood estimated parameters from the seed flow model

fields effect and long-distance dispersal (by mowers, for example) because of the scale of the study and the fact that only 5 feral populations were present in 2004.

We extended the seed model to 2003 and 2002 with the same construction type (exponential kernels, mean





	ms	а	logL
2003-2004	0.23	403.8	- 430.2
2002-2004	0.28	374.6	- 433.4

Maximum Likelihood estimated parameters from the seed flow model We extended the seed model without the contribution of the sown fields in 2005 but with 2002 and 2003 data in the area study :

ms fraction decreases to 0.23 and 0.28

* mean dispersal distance was multiplied by 10 !



2005 fields have no contribution on 2005 feral

39.1 m of mean seed dispersal distance

yopulations (*alpha* estimated to 1.0)



Mean distances between feral populations of the study area in 2005 and their origin fields in 2004 and 2003

Conclusions and prospects

- The seed dispersal model succeeds in estimating parameters for two years data and a sub-area of 7 km². But, 40 meters is shorter than we can expect for seed long distance dispersal and 400 meters is higher than expected.
- In order to have a more accurate estimation of long-term dispersal events, we project to take into account the whole region of

dispersal distance estimated for the years together ...):

Mean seed dispersal distance greatly improves to 400 m (versus 40 m) with 2003 data, which can be due to spatial configuration of the study area where 2004 fields were closer to relatives feral populations in 2005 than 2003 fields. As the distance estimated takes into account all year contributions together, it is still impossible to separate seed dispersal per year.

ms fraction decreases from 54% to 23% with 2003 data (16 fields of 2003 contribute to 2005 feral populations). ms fraction does not decrease with 2002 data (fields of 2002 do not contribute to 2005 feral populations).

Selommes (42 km²) and to include all populations and fields of 2002 to 2005 (239 populations in 2005 versus 38 in the sub-area).

The seed dispersal distance should be included in models estimating the escape of Genetically Modified Plants genes from fields. 40 meters is higher than ballistic abilities of dehiscent fruits, suggesting that dispersal is strongly linked with anthropogenic effects as vehicles traffic, mowing and harvesting.

This summer, we will conduce an in-situ experimentation to mesure seed spillage from trunks during oilseed rape harvesting period in Selommes area.

Litterature cited

J. Burczyk, W. T. Adams, D. S. Birkes, and I. J. Chybicki (2006). Using Genetic Markers to Directly Estimate Gene Flow and Reproductive Success Parameters in Plants on the Basis of Naturally Regenerated Seedlings. Genetics

Colbach N, Clermont Dauphin C & Meynard JM 2001b GENESYS: a model of the influence of cropping system on gene escape from herbicide tolerant rapeseed crops to rape volunteers - II. Genetic exchanges among volunteer and cropped populations in a small region. Agric Ecosyst Environ 83: 255-270

Garnier, A. (2006). Population dynamics and dispersal of a cultivated species escaped from fields : feral oilseed rape as a case study. Simulation models, statistics and genetics. PhD thesis Garnier, A., Pivard, S., Lecomte, J. (2008). Measuring and modelling secondary anthropogenic seed dispersal along roadverges for feral oilseed rape. Basic Appl. Ecol



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Further informations

http://www.ese.u-psud.fr/gmbioimpact/

