



iFBM

THE REFERENCE IN THE BARLEY TO THE BEER



Last findings on T2/HT2 on malting barley and behaviour from malting barley to malt

Dr Régis Fournier, IFBM



Aim

- ⇒ ***Fusarium* contamination occurs in the field: control has to be setup within the field**
- ⇒ **Growth can occur during malting process (favourable humidity and temperature)**
- ⇒ **Better knowledge of Fusariotoxins from field to end products and by-products**
- ⇒ **Recommendation to reduce mycotoxin level in field and process**

The project

⇒ **BARSAFE** : *Fusarium langsethiae*, from barley culture (*Hordeum vulgare*) to the finished products (beer) and by-products: study of the biology and the epidemiology of the pathogen, the conditions of T2/HT2 toxins production, of their transfer, biological breakdown and toxicity, for a better sanitary risk management

PARTNERS

Institut Français de la Brasserie et de la Malterie (IFBM)

Arvalis Institut du Végétal

Laboratoire des Sciences du Génie Chimique

Laboratoire Génie chimique

Laboratoire de Pharmacologie-Toxicologie

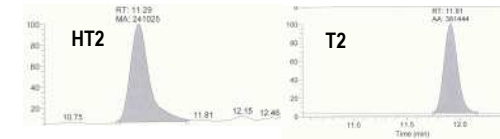
Laboratoire de Pathologie Végétale et Epidémiologie

- ⇒ **Study of *Fusarium* contamination & mycotoxins production in field**
- ⇒ **Impact of *Fusarium* strains viability during storage**
- ⇒ **Mycotoxins behaviour during malting & brewing process**
- ⇒ **Mycotoxins behaviour from feed to end products**

T2 / HT2 Toxins

Analysis method : barley & malt

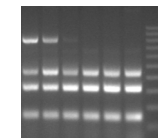
Multitrichothecene analysis : HPLCMSMS, LOD = 1 ppb)



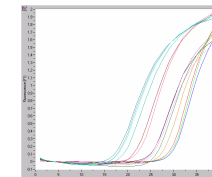
Isolation and identification of species by culture on agar media



Identification of species by specific primer DNA analysis



Quantification of species by real time PCR DNA analysis



Barley *Fusarium* species

Fusarium species actually found on brewing barley

F. graminearum

F. culmorum

F. langsethiae

F. tricinctum

F. avenaceum

F. poae

F. sporotrichioides (rare)

Mycotoxin

type B TCT (DON, NIV), zearalenone

TCT A, TCT B

TCTA

Fusaric acid, beauvericine, fusarine C

Moniliformine (*F. avenaceum*)

F. heterosporum

F. acuminatum

F. sambucinum

F. equiseti

F. lateritium

F. crookwellense (rare)

+ *Microdochium nivale*

Enniatines, Beauvericine

Other genera closely studied : *Penicillium*, *Alternaria*

Fusarium Identification

Fusarium spp
F. graminearum
F. tricinctum
F. culmorum
F. avenaceum
F. langsethiae
F. poae
F. sporotrichioides
F. heterosporum
F. solani
F. oxysporum
F. acuminatum
F. sambucinum
F. equiseti
F. lateritium
F. proliferatum
F. subglutinans
F. anthophyllum
F. verticillioides
F. crookwellense
F. chlamydosporum
F. scirpi
Microdochium nivale
Microdochium majus

Visual identification

Fusarium spp
F. langsethiae
F. sporotrichioides
F. graminearum
F. culmorum
F. poae
F. tricinctum
F. avenaceum
Microdochium nivale
Microdochium majus
F. heterosporum
F. solani
F. moniliforme
F. oxysporum
F. acuminatum
F. sambucinum
F. equiseti
F. lateritium

PCR identification

Fusarium spp
F. langsethiae
F. sporotrichioides
F. graminearum
F. culmorum
F. poae
F. tricinctum
F. avenaceum
Microdochium nivale
Microdochium majus

F. verticillioides
F. proliferatum
F. subglutinans

PCR quantification

Harvest analysis

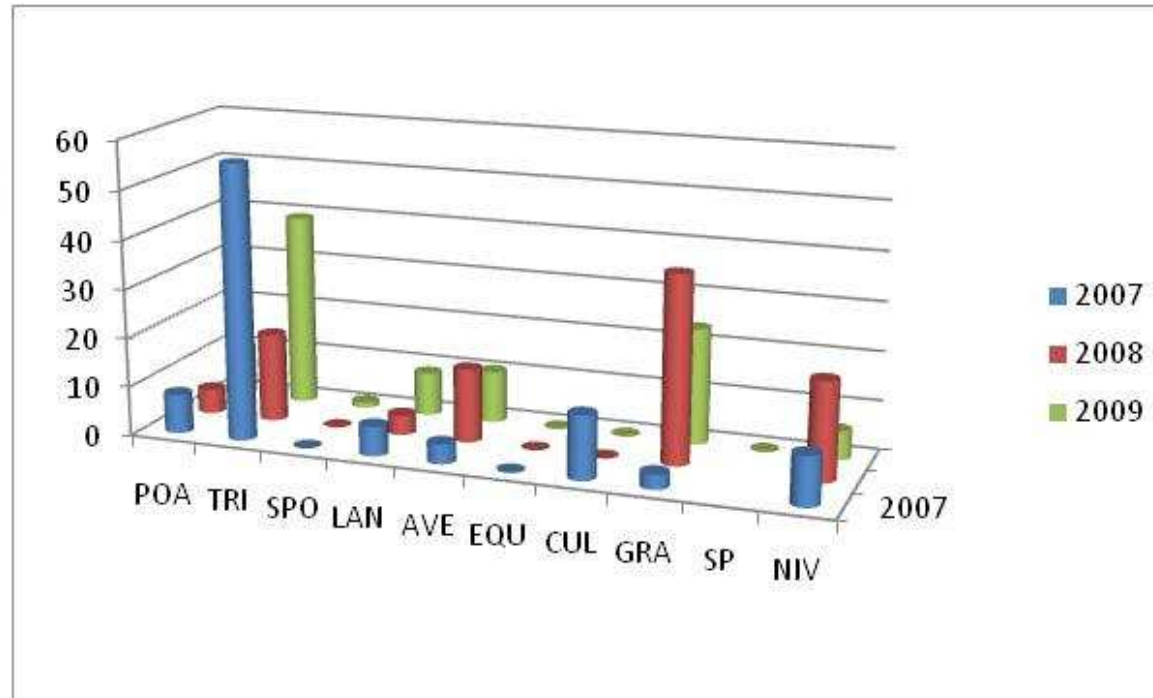
From 2003 to 2008, 150 to 200 malting barley samples are analysed every year

- Multitrichothecenes (LC-MS/MS)**
- Visual identification**
- Identification by PCR**

Fusarium : evolution from 2007 à 2009

Fusarium strains solated from calibrated French malting barley

Fusarium species proportion (%)



⇒ TCT A

- ⇒ *F. langsethiae*
- ⇒ *F. poae*
- ⇒ *F. sporotrichioides*

⇒ Autres

- ⇒ *F. tricinctum*
- ⇒ *F. avenaceum*
- ⇒ *M. nivale*

⇒ TCT B

- ⇒ *F. graminearum*
- ⇒ *F. culmorum*
- ⇒ *F. poae*

Fusariotoxins: Harvest 2006-2008

<i>Barley</i>		<i>T2+HT2</i>	<i>DON</i>	<i>NIV</i>
Harvest 2006				
Spring	mean	73	60	20
	max	708	508	135
Winter	mean	5	80	nd
	max	41	660	
Harvest 2007				
Spring	mean	59	121	83
	max	489	957	576
Winter	mean	2	26	8
	max	63,5	176	136
Harvest 2008				
Spring	mean	35	187	24
	max	146	1082	202
Winter	mean	12	122	7
	max	29	416	69
Harvest 2009				
Spring	mean	46	89	18
	max	687	409	229
Winter	mean	22	242	4
	max	306	1226	20

Toxinogenic Potential of *Fusarium Langsethiae*

Isolated from malting barley
Harvest 2005, 2006, 2007

	T2/HT2	TCT B
<i>Fusarium poae</i>	0-100 ppb	0-700 ppb
<i>Fusarium sporotrichioides</i>	59-668 ppb	None
<i>Fusarium langsethiae</i>	50 000 - 300 000 ppb	None

Harvest 2008

	T2/HT2	TCT B
<i>Fusarium sporotrichioides</i>	300 ppm	nd

Conditions: sterilised barley, 2 weeks at 25°C

T2/HT2 producer

Fusarium langsethiae



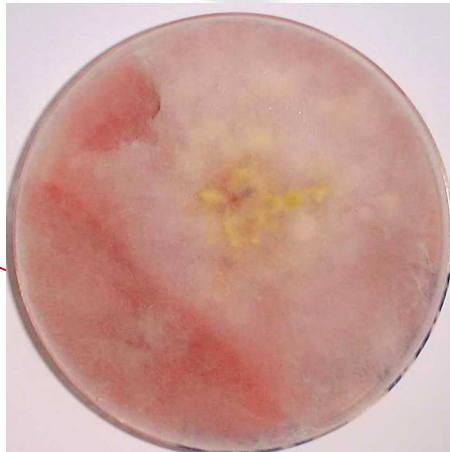
F. langsethiae on PDA



Microconidies of *F. langsethiae*

T2/HT2 producer

Fusarium sporotrichioides

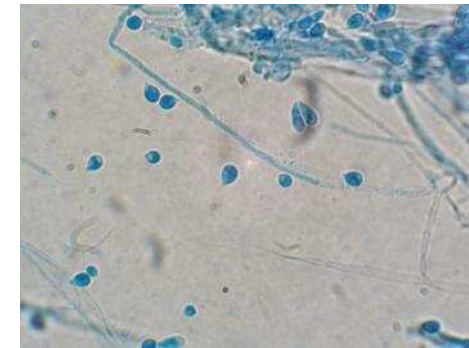


F. sporotrichioides on PDA



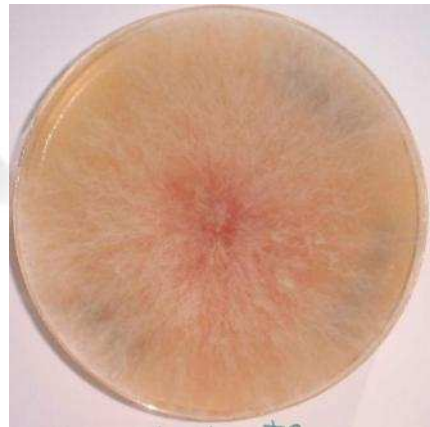
Microconidies of *F. sporotrichioides*

Fusarium poae



Strains	MYCOTOXINS								GUSHING
	Eniatin B (ppb)	Beauvericin (ppb)	T2triol (ppb)	T2 (ppb)	HT2 (ppb)	3-DON (ppb)	DON (ppb)	NIV (ppb)	
E-488 -08-088	0	1200	0	2	26	0	0	80	0
E-488 -0-089	0	450	0	0	0	0	0	480	0
E-488 -08-092	0	980	0	0	0	0	0	80	0
E-488 -07-094	0	1900	0	0	0	0	0	600	0
E-488 -07-008	0	1100	0	2	26	0	0	78	0

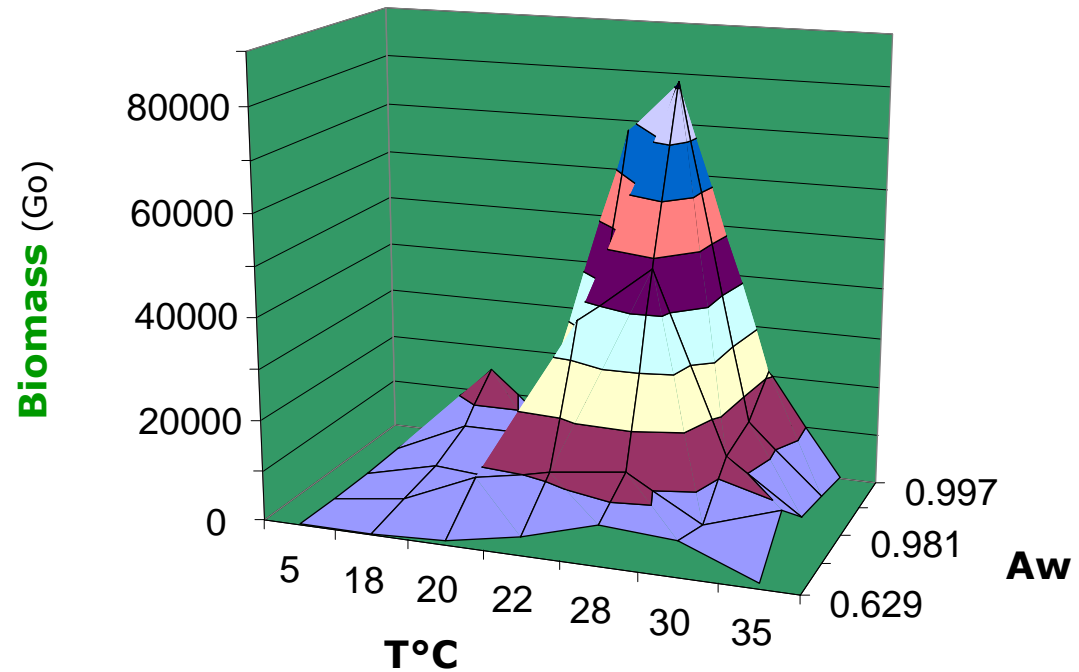
Fusarium graminearum



Strains	MYCOTOXINS								GUSHING
	Eniatin B (ppb)	Moniliformin (ppb)	T2triol (ppb)	T2 (ppb)	HT2 (ppb)	3-DON (ppb)	DON (ppb)	NIV (ppb)	
E-488 -07-071	0	0	0	0	0	0	100	0	0
E-488 -07-073	0	0	0	0	0	0	64	0	0
E-488 -07-075	0	0	0	0	0	0	189	0	0
E-488 -07-076	0	0	0	0	0	0	117	0	0
E-488 -07-079	0	0	0	0	0	0	90	55	0

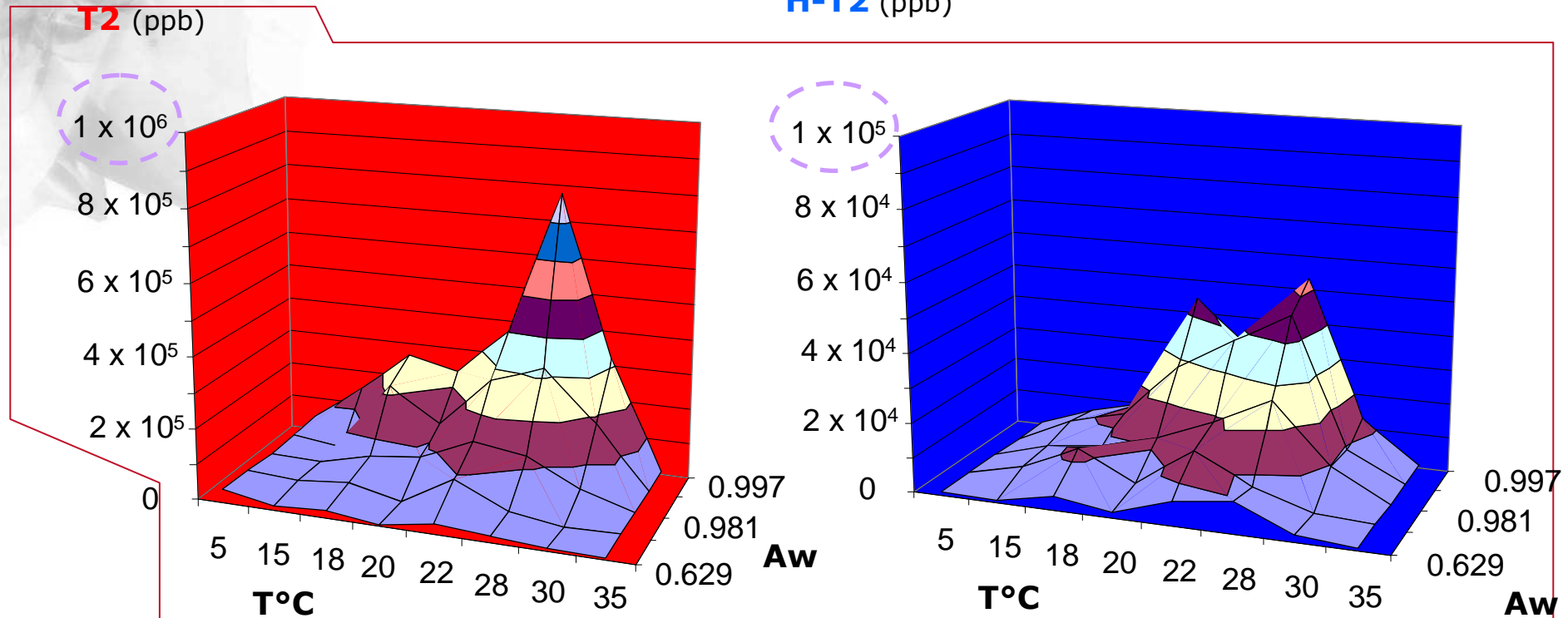
Ecotoxicogenesis conditions : Growth conditions

***F. langsethiae* (2008)**
Culture on autoclaved barley



- Optimal growth temperature : 28°C
- Optimal growth Aw 0.992
- No growth at 5 and 35°C

Ecotoxicogenesis conditions : Toxin production



- [T2] > [H-T2]
- Optimal conditions for production : 28°C ; Aw 0.997
- No production at 5 and 35°C

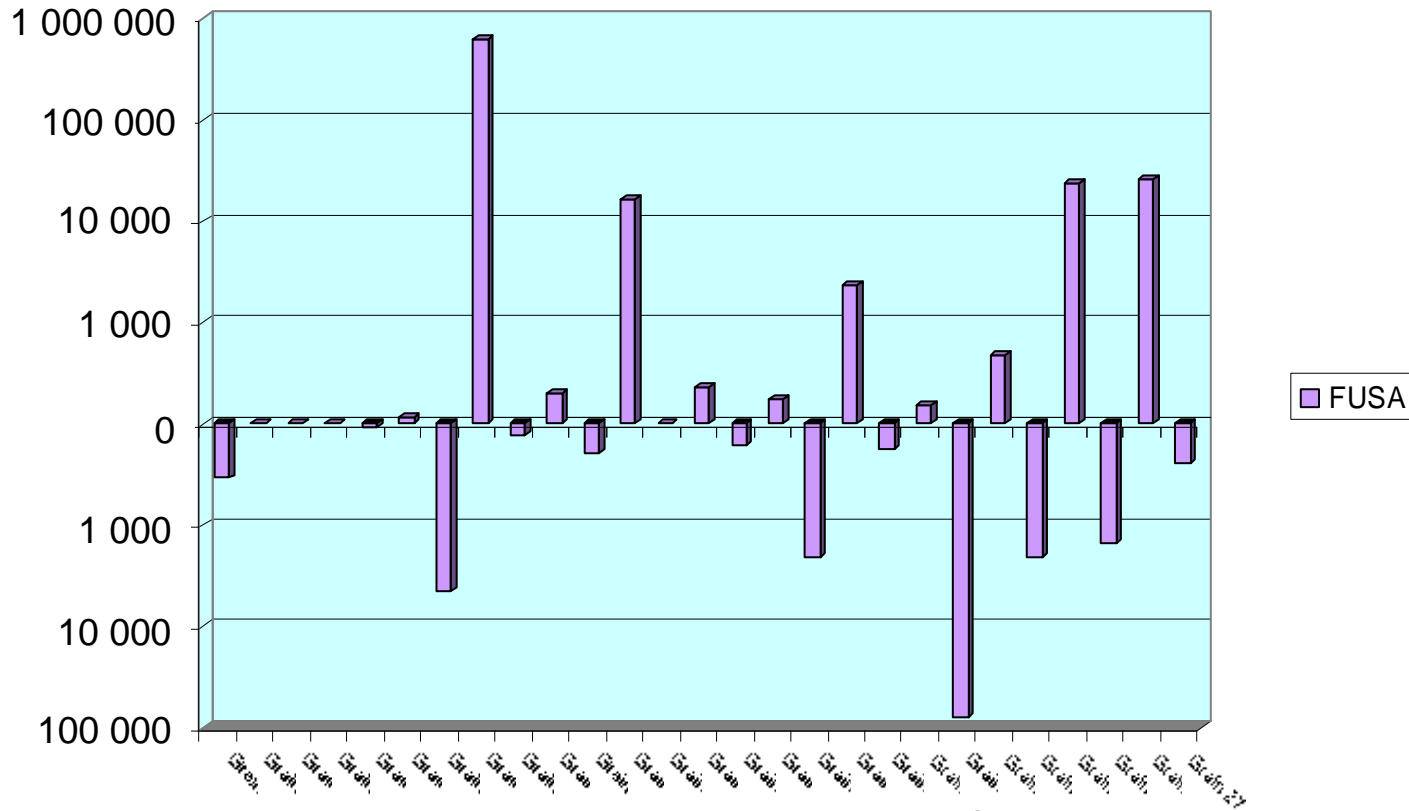
LGC Toulouse



Symptom infection



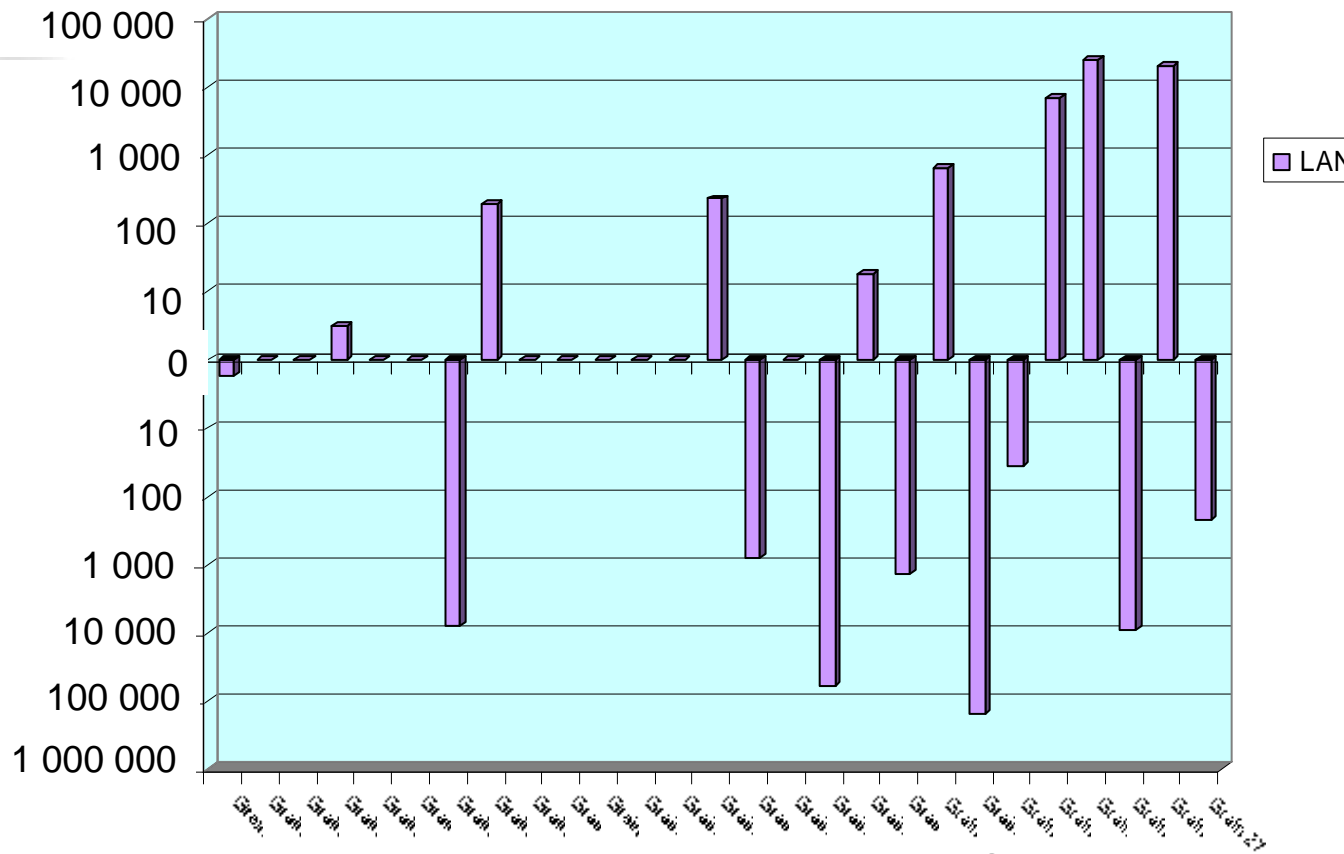
Fusarium genome amount /1000 barley genomes



Symptom infection



F. langsethiae
genome amount
/2500 barley
genomes



Symptom infection

⇒ **Other Fusarium species**

FIELD EXPERIMENTS

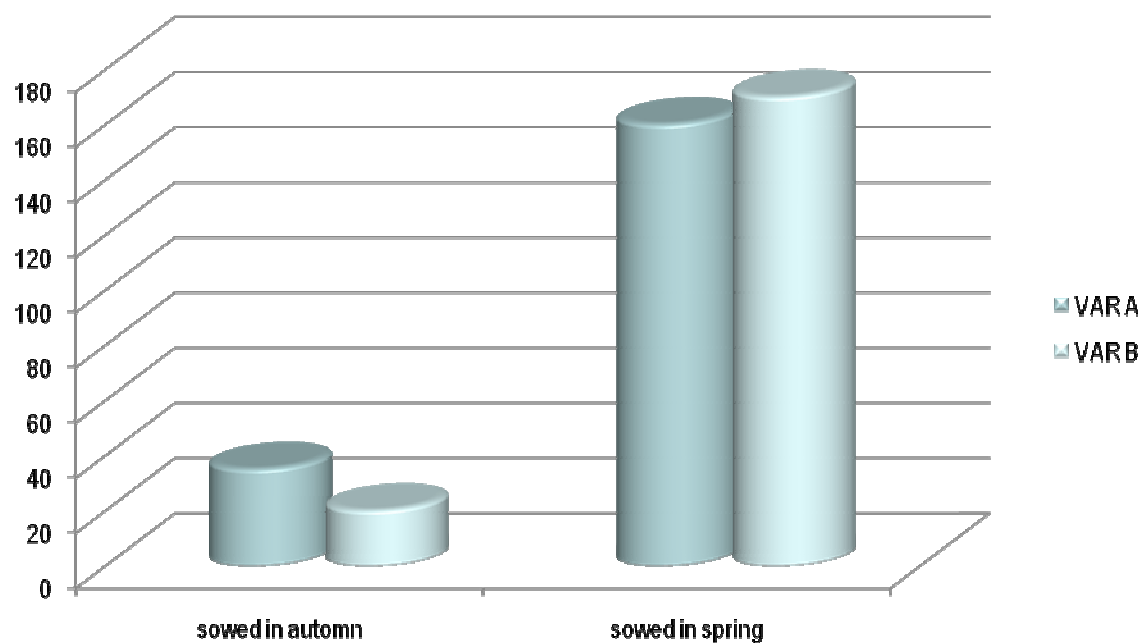


Influence of sowing date

YEAR 2007

Spring Barley : same site, different sowing date (autumn and spring)

T2+HT2 ($\mu\text{g}/\text{kg}$)



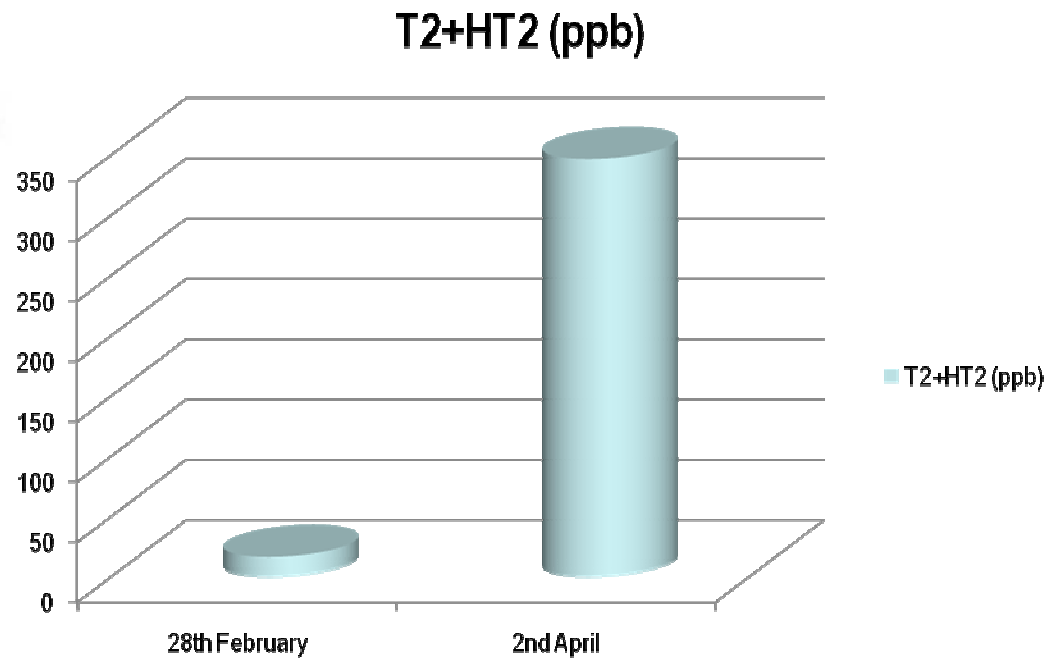
Spring Barley: the sowing date is important



Influence of sowing date

HARVEST 2008 Spring Barley

same variety, same site, different date (same year)



The later the sowing takes place, the higher the contamination

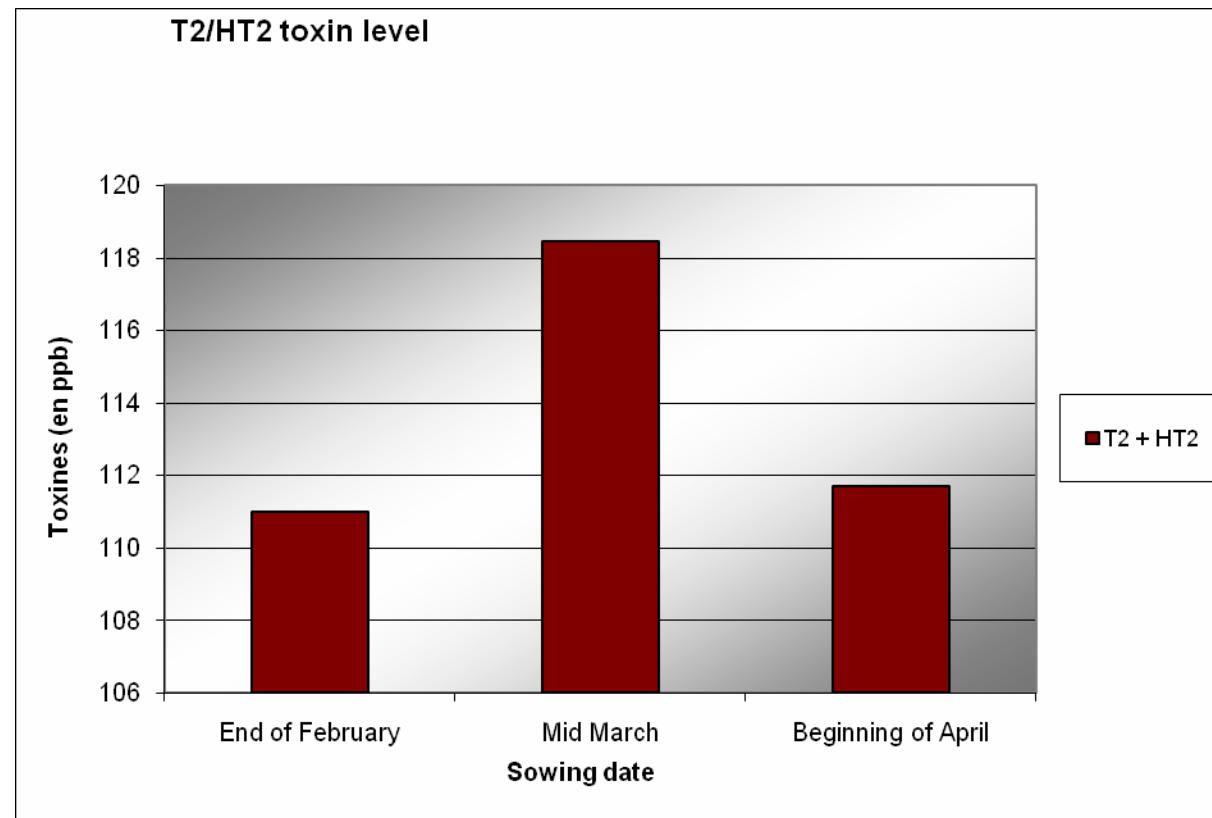
Sowing date trial

⇒ 3 sowing dates of a Spring Barley (Tipple):

- End of february/ Mid March / beginning of April

⇒ 3 repetitions

- No difference observed for T2 + HT2 level
- Climatic influence



MALTING PROCESS

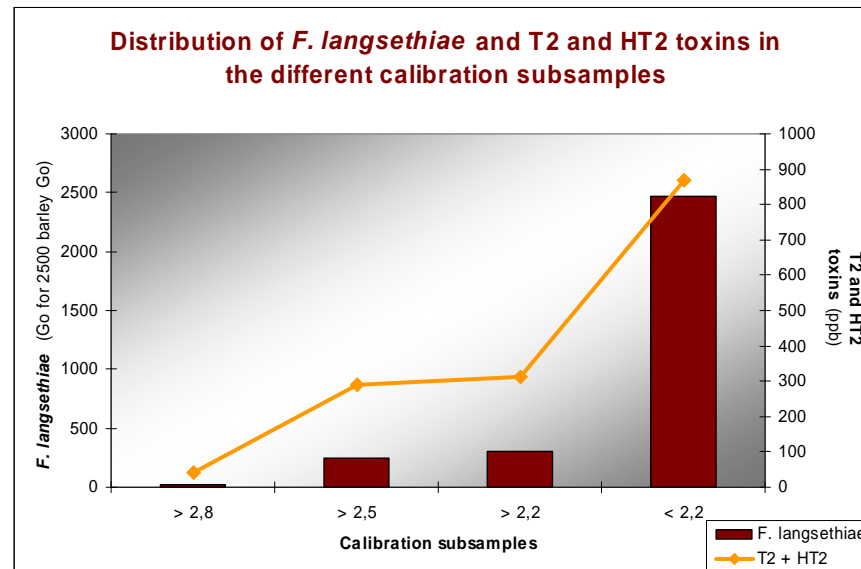


T2+HT2 toxin ($\mu\text{g}/\text{kg}$)

Barley	>2.8	2.8-2.5	2.2-2.5	<2.2
HT2	74	136	1103	5000
T2	26	61	379	917
HT2+T2	100	197	1482	5917

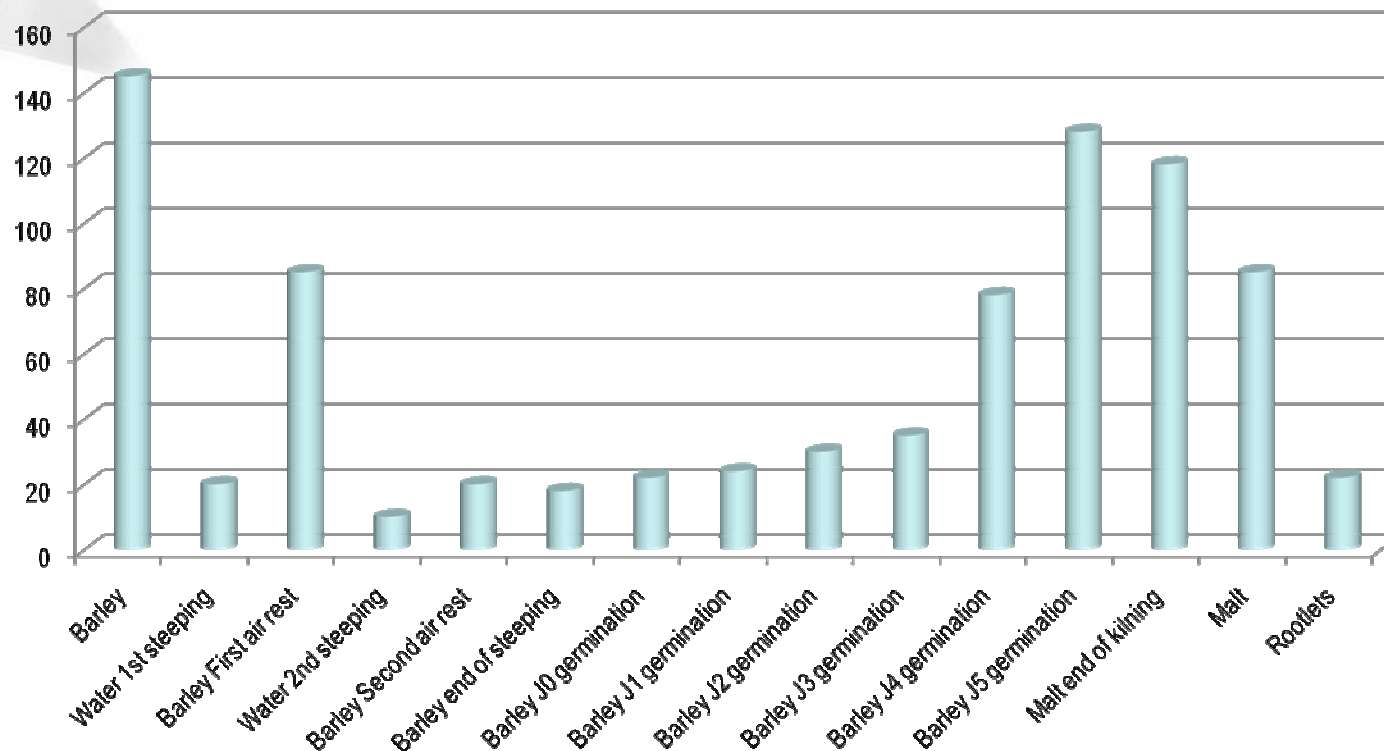
Calibration: impact for the malting industry

- 1st step = Elimination during harvest : smallest grains including those presenting *Fusarium langsethiae* symptoms are not recovered and a great part remains in the field
- 2nd step = Elimination due to calibration : according to *Fusarium langsethiae* symptom described above, the most contaminated fraction for both type A trichothecenes and *Fusarium langsethiae* is the smallest fraction < 2,2. Calibrating over 2,8 allows a maximal elimination of both mycotoxins and fungi.



T2+HT2 toxins are more concentrated in the smallest fraction (dust, small kernels, broken kernels...)

T2+HT2 ($\mu\text{g}/\text{kg}$) follow up in a 2008 sample malted in November 2008

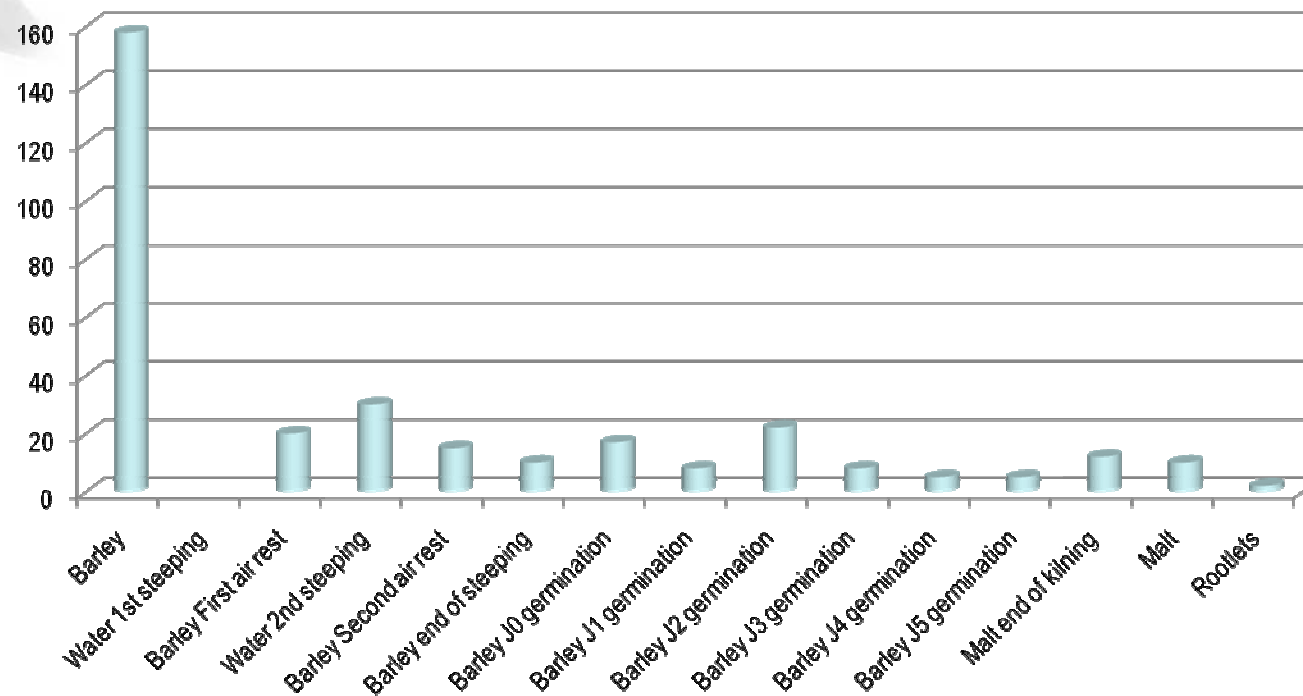


Reaccumulation of T2+HT2 toxins during the germination steps



Malting process

T2+HT2 ($\mu\text{g}/\text{kg}$) follow up in a 2008 sample malted in March 2009



No re-accumulation of T2+HT2 toxins during the germination steps
F. Langsethiae did not survive during storage

General conclusion

- ⇒ The *Fusarium langsethiae* and T2/HT2 toxin contaminations are still present in 2009
- ⇒ *Fusarium sporotrichioides*, high T2/HT2 producer, has appeared in 2009 in French Malting Barley
- ⇒ Harvest calibration, barley calibration prior to malting process and steeping step during malting process allow a consequent elimination of the toxins

General conclusion

⇒ Thanks for your attention