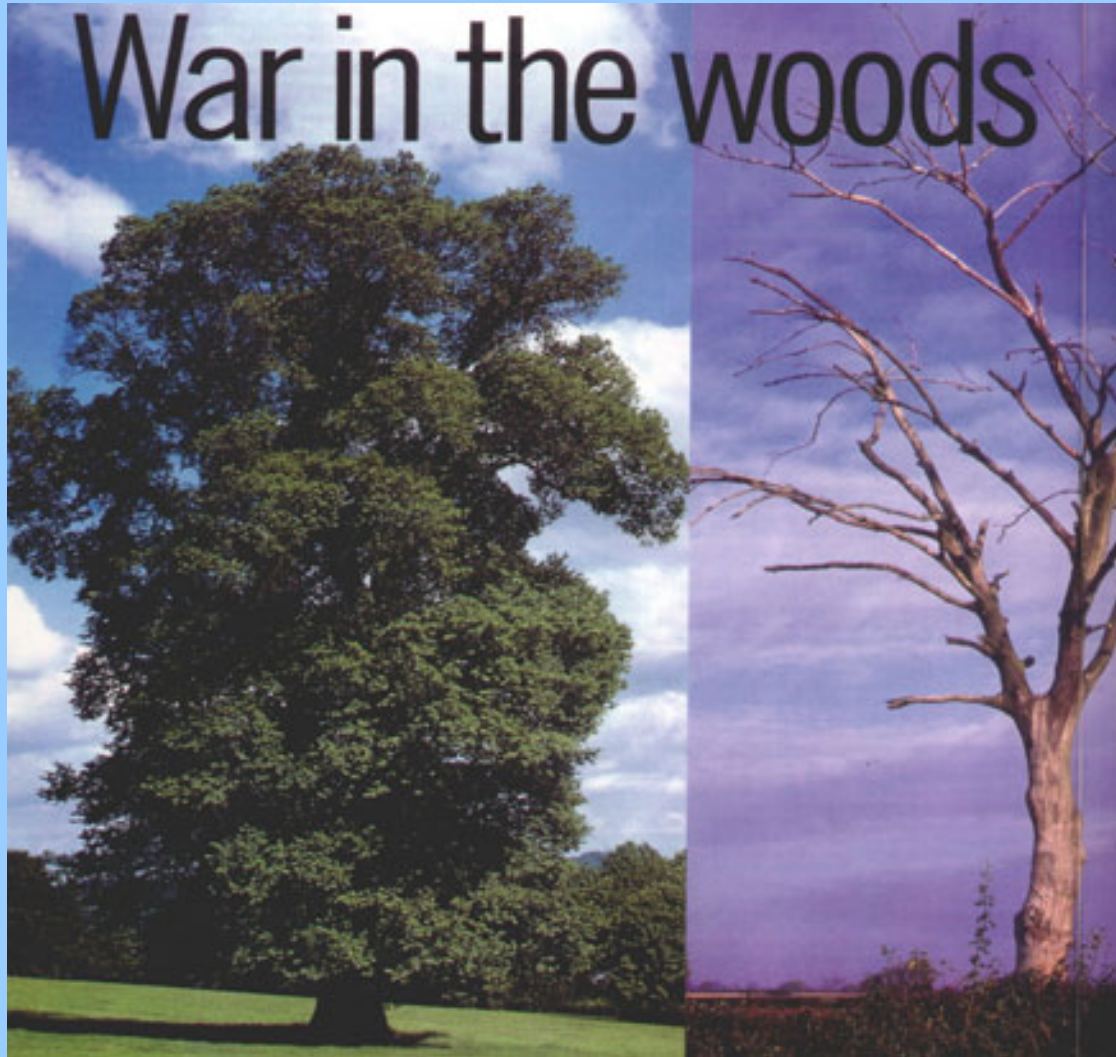


MANIPULATING ELMS: PROGRESS AND POTENTIAL

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War in the woods



Cycles of *Ophiostoma novo ulmi* Spread

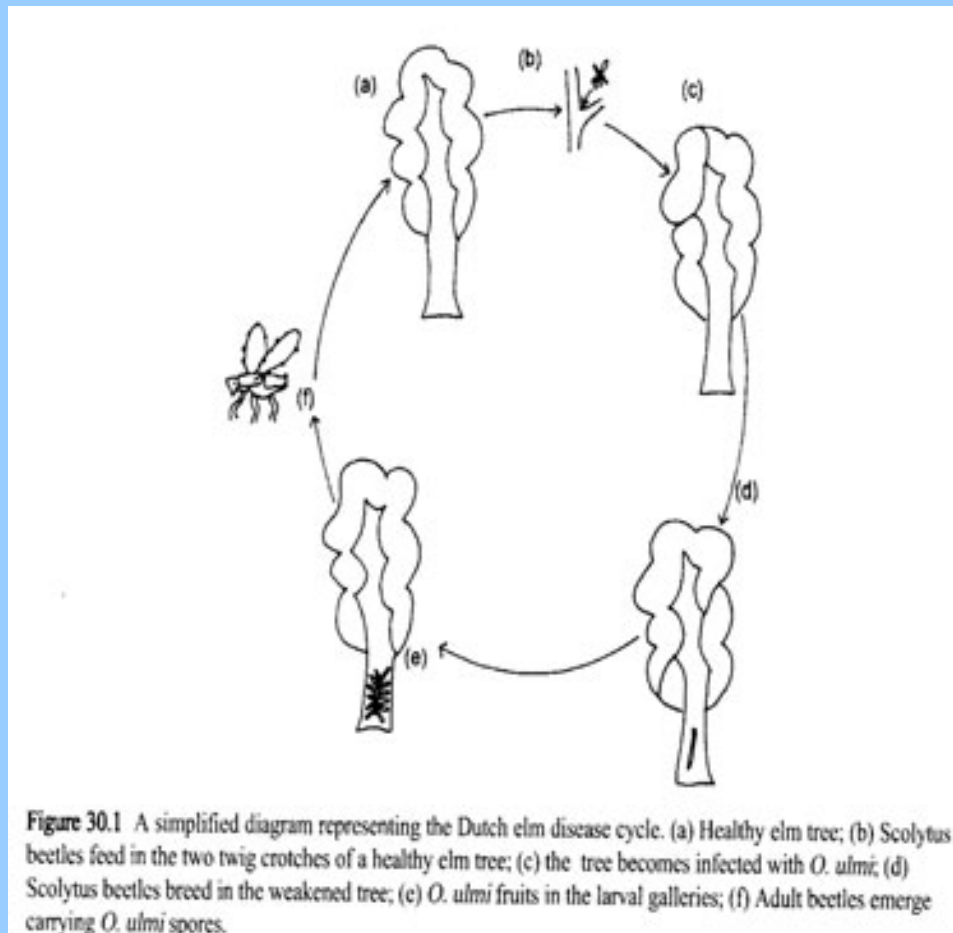
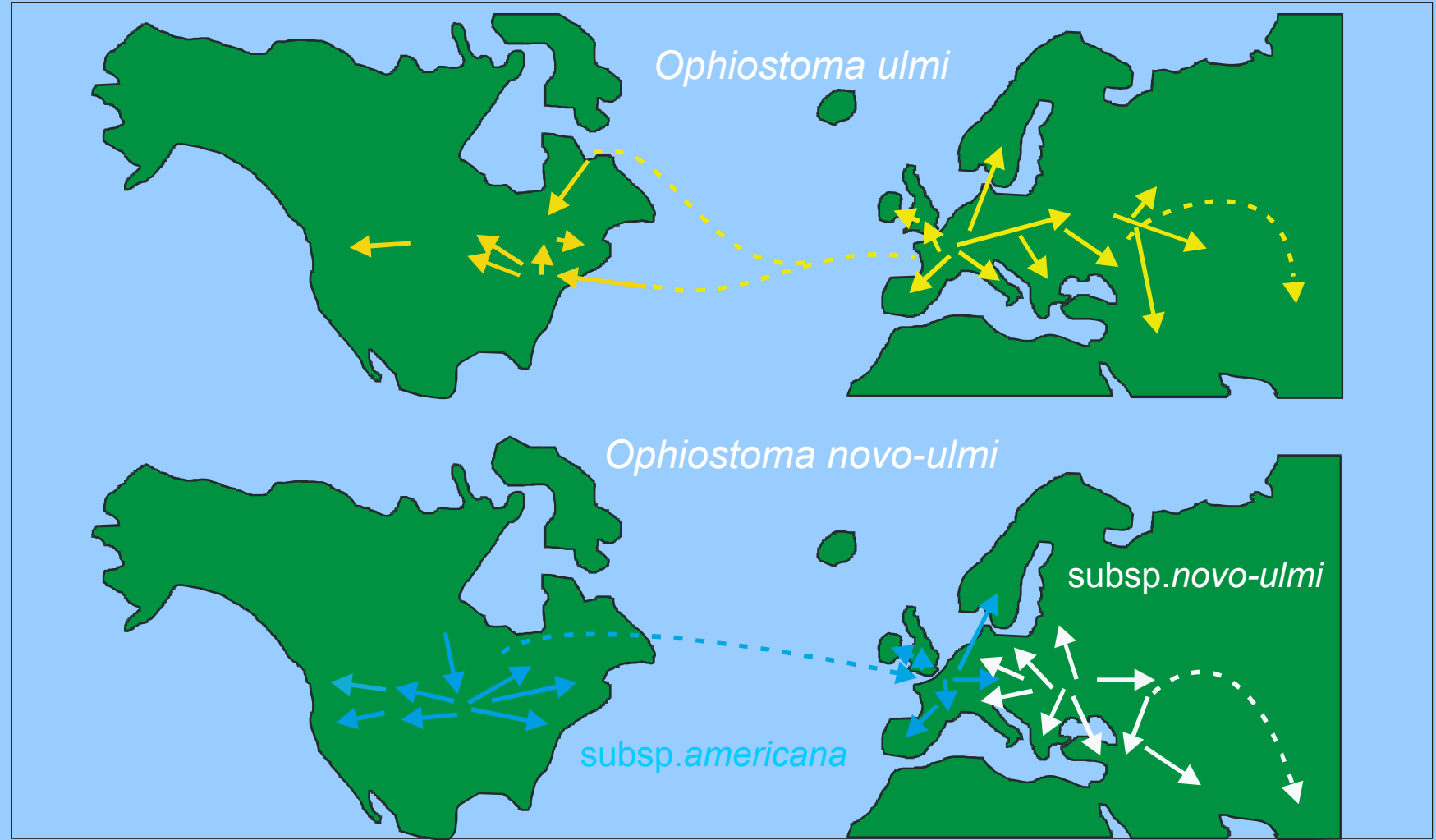


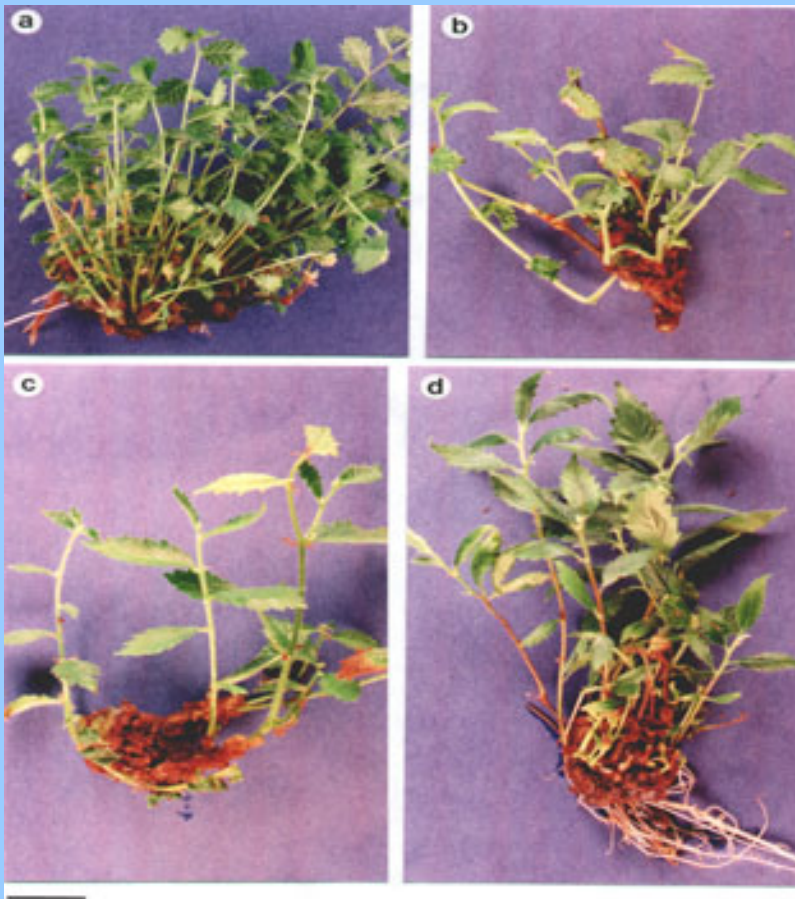
Figure 30.1 A simplified diagram representing the Dutch elm disease cycle. (a) Healthy elm tree; (b) Scolytus beetles feed in the two twig crotches of a healthy elm tree; (c) the tree becomes infected with *O. ulmi*; (d) Scolytus beetles breed in the weakened tree; (e) *O. ulmi* fruits in the larval galleries; (f) Adult beetles emerge carrying *O. ulmi* spores.





ELM BIOTECHNOLOGY

Tissue Culture and Plant Regeneration

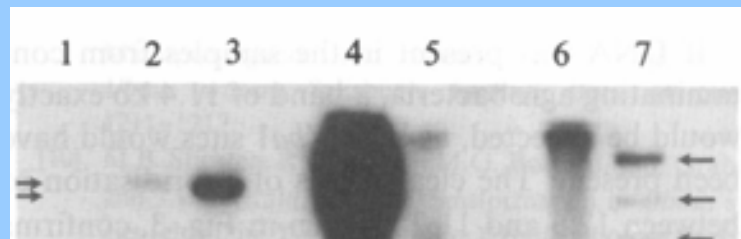


- ❖ *Ulmus procera* (a)
- ❖ *Ulmus glabra* (c)
- ❖ *Ulmus americana* (d)
- ❖ *Ulmus parvifolia* (b)

Agrobacterium tumefaciens



- ❖ Aberrant shoots regenerated
- ❖ Genomic blot confirmed transgenic
- ❖ Low efficiency transformation & regeneration





Agrobacterium rhizogenes

- ❖ SR4C1, SR4C2
- ❖ Dwarf shoots
- ❖ Reduced internode distance
- ❖ Reduced Leaf Area



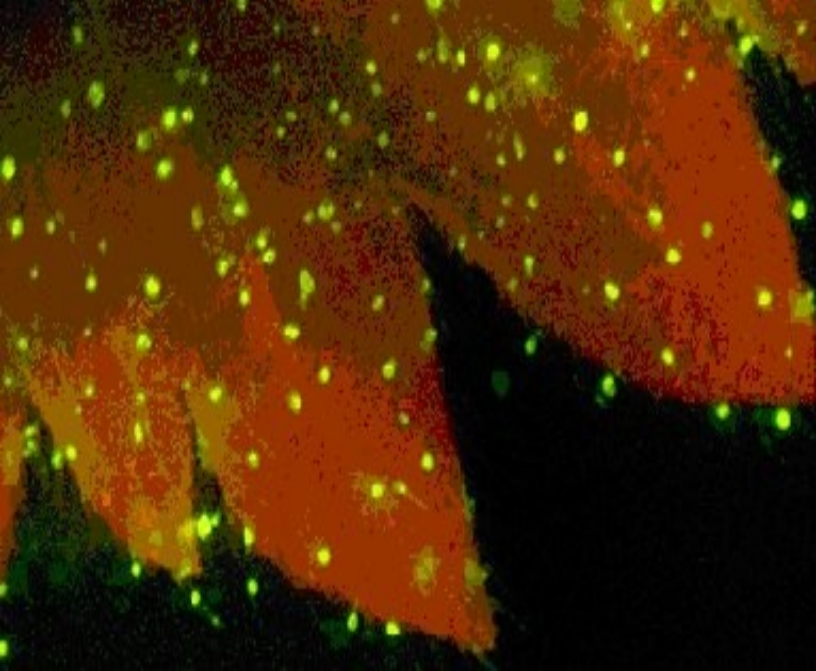
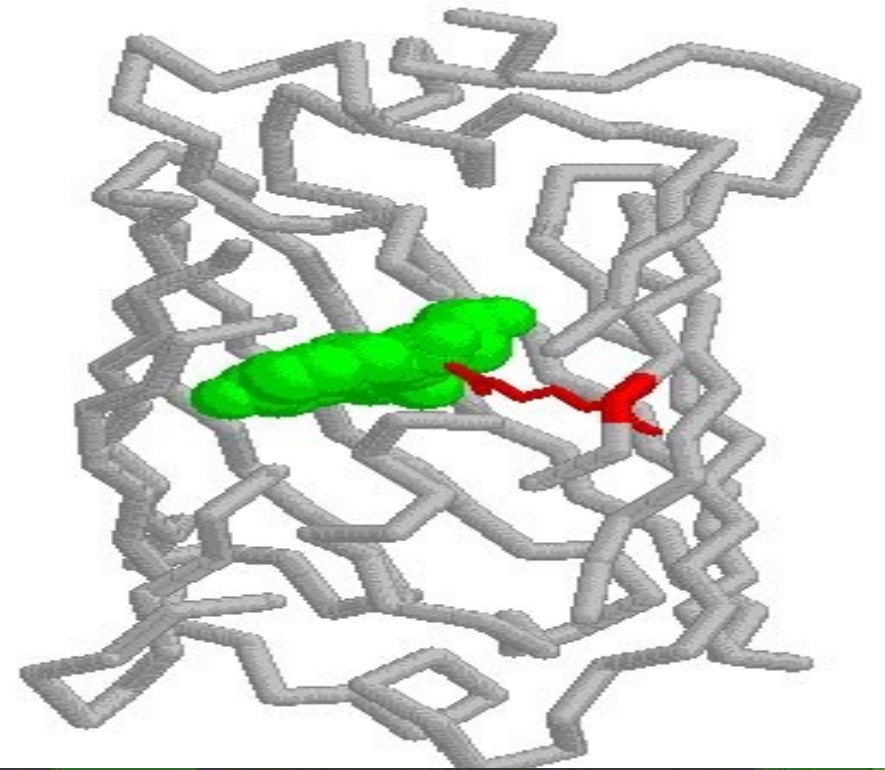


SR4C1 in Soil

Biolistics



- ❖ CaMV35S::Jellyfish Green Fluorescent Protein
- ❖ CaMV35S::GUS Intron
- ❖ Construct screening tool



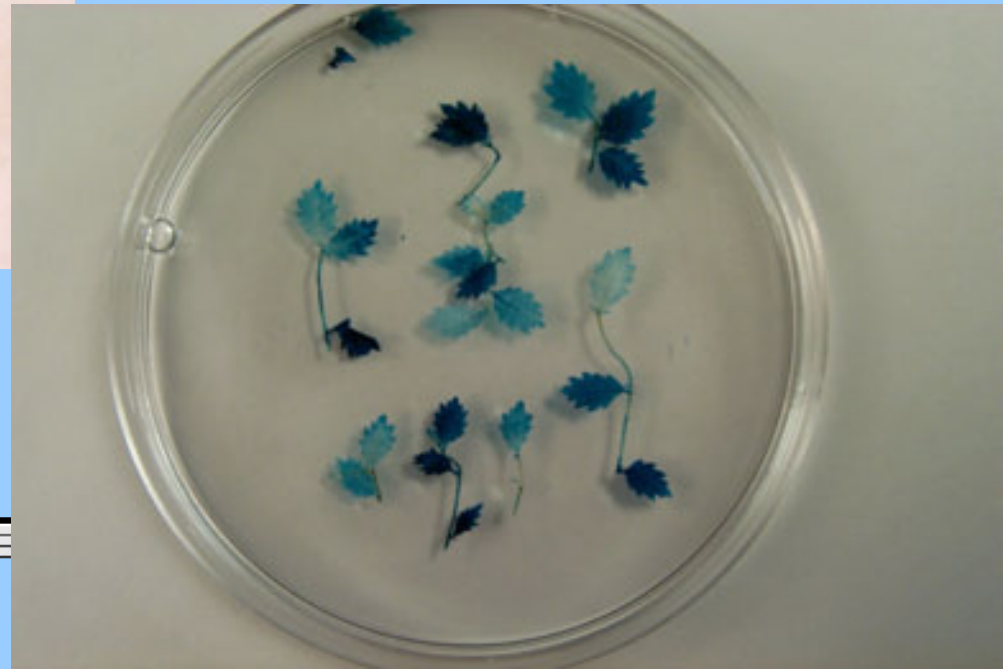
Gfp+ve Elm Growth



GM *U. procera* Phenotypically Normal Regenerants



- ❖ GUS expression in GM *U. procera* regenerants
- ❖ CaMV::35SGUS-Intron
- ❖ >100x increase GUS activity c.f. background

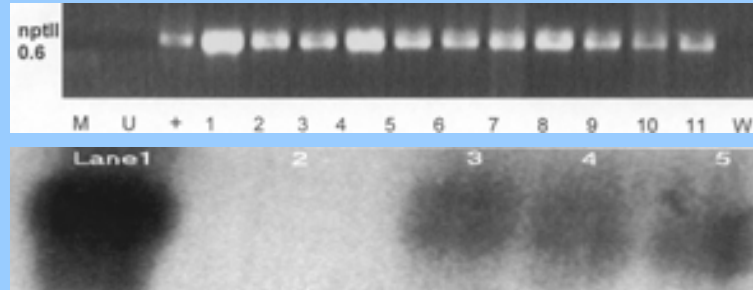


Variations in GUS Expression

Line	GUS Activity	+/-SD
2-5 Leaf	5.64	2.40
2-5 Stem	6.33	3.76
2-2 Leaf	6.31	3.71
2-2 Stem	7.52	4.39
3-7 Leaf	13.03	4.85
3-7 Stem	14.89	4.55
SR4 Leaf	0.00	0.02
SR4 Stem	0.00	0.02

GUS Act.: nMol MU/ug Protein/h; Mean 9 replicates

Characterisation GM Elms



- ❖ PCR analysis
- ❖ DNA-DNA Hybridisations
- ❖ Rooting on Kanamycin

Transfer to Soil

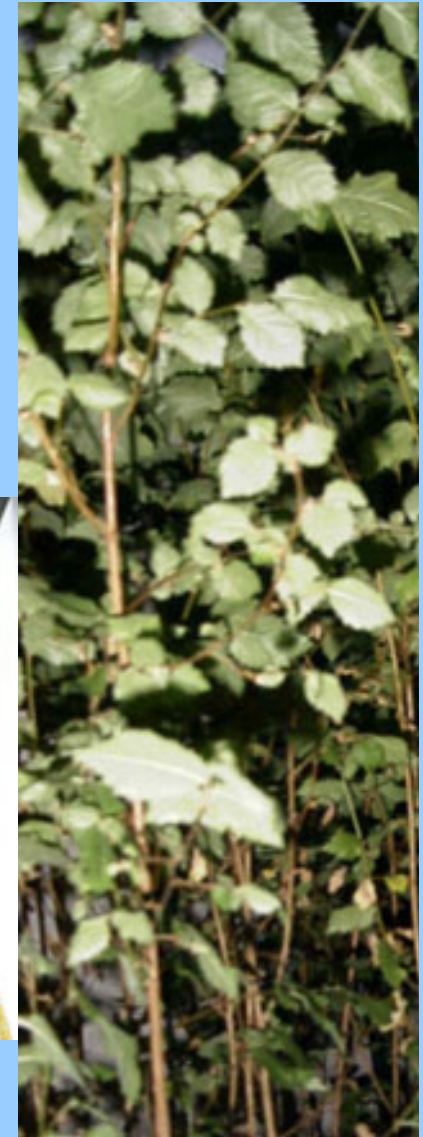
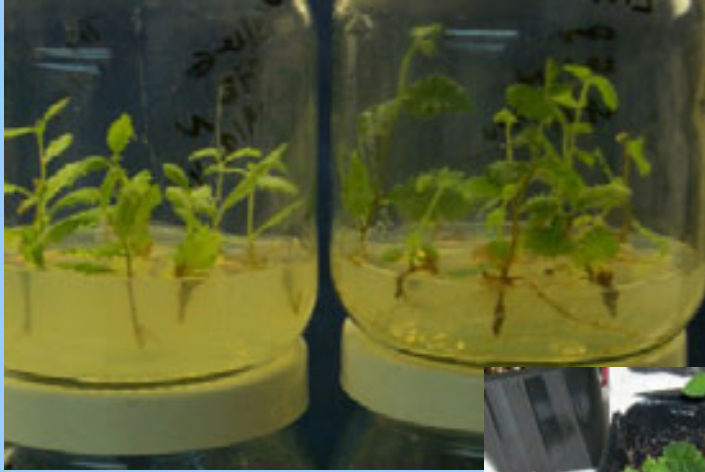
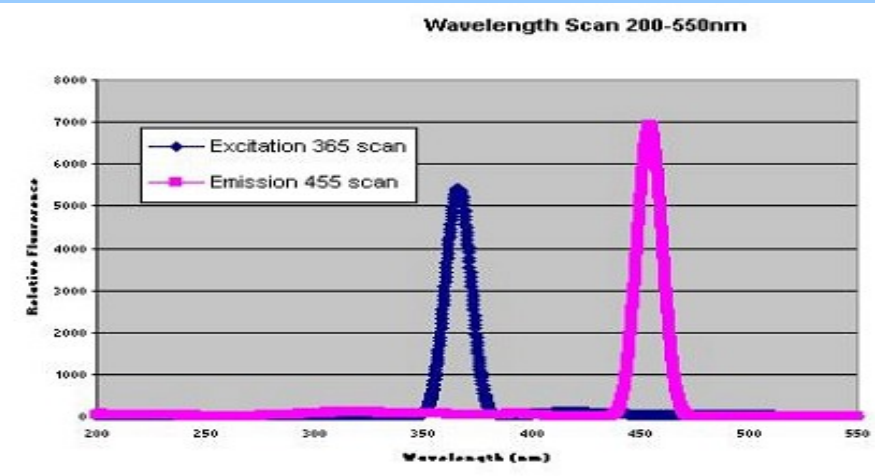
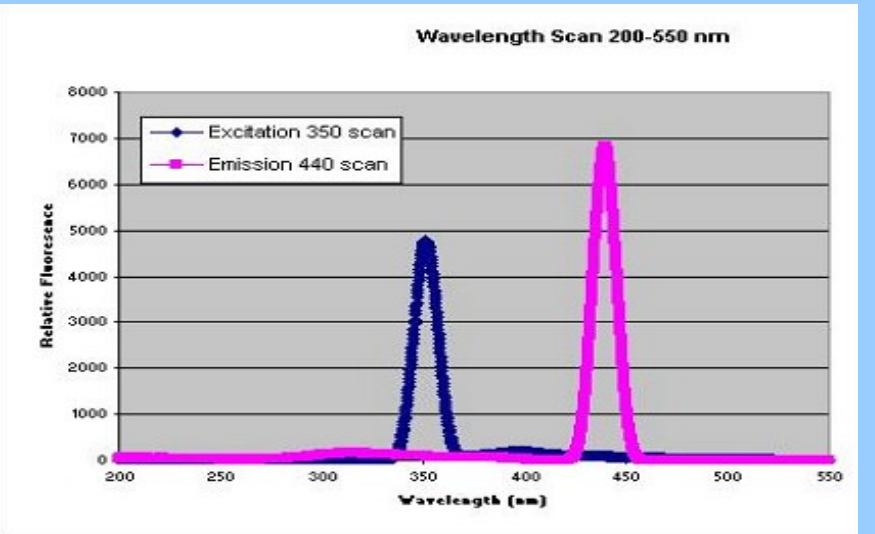


Figure 10 NaMU Fluorescence Spectra



a) Using a Conventional Fluorimeter



b) using RealTime ALEXAFilter

Figure 12 Real Time GUS analysis ZH4 and untransformed *Ulmus procera* SR4

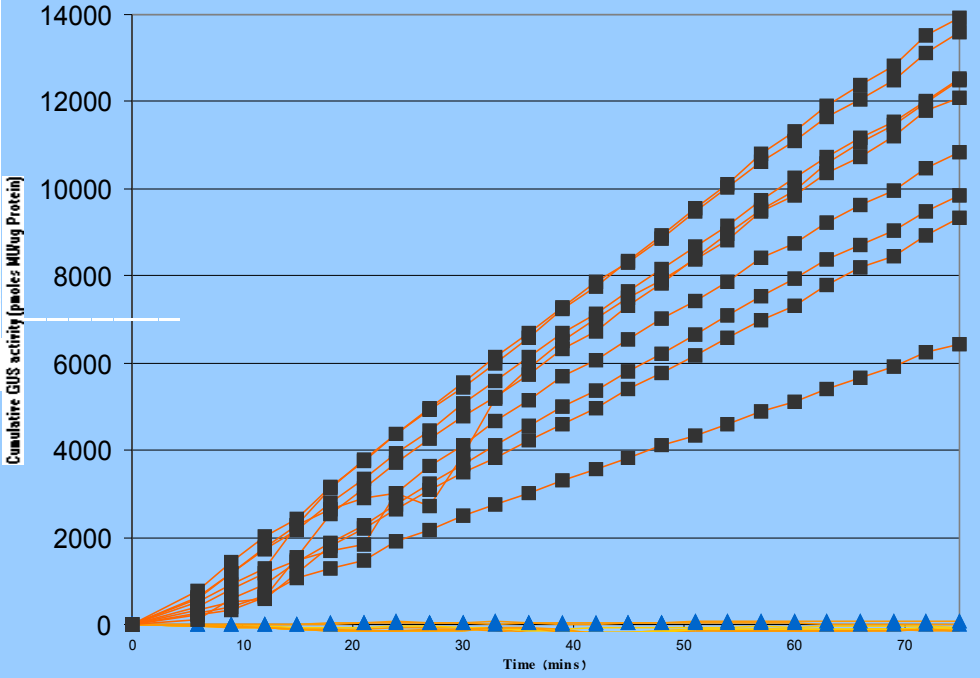


Figure 8 Seasonal GUS expression 2003/4

Error bars are standard errors of means from at least 9 independent replicates.

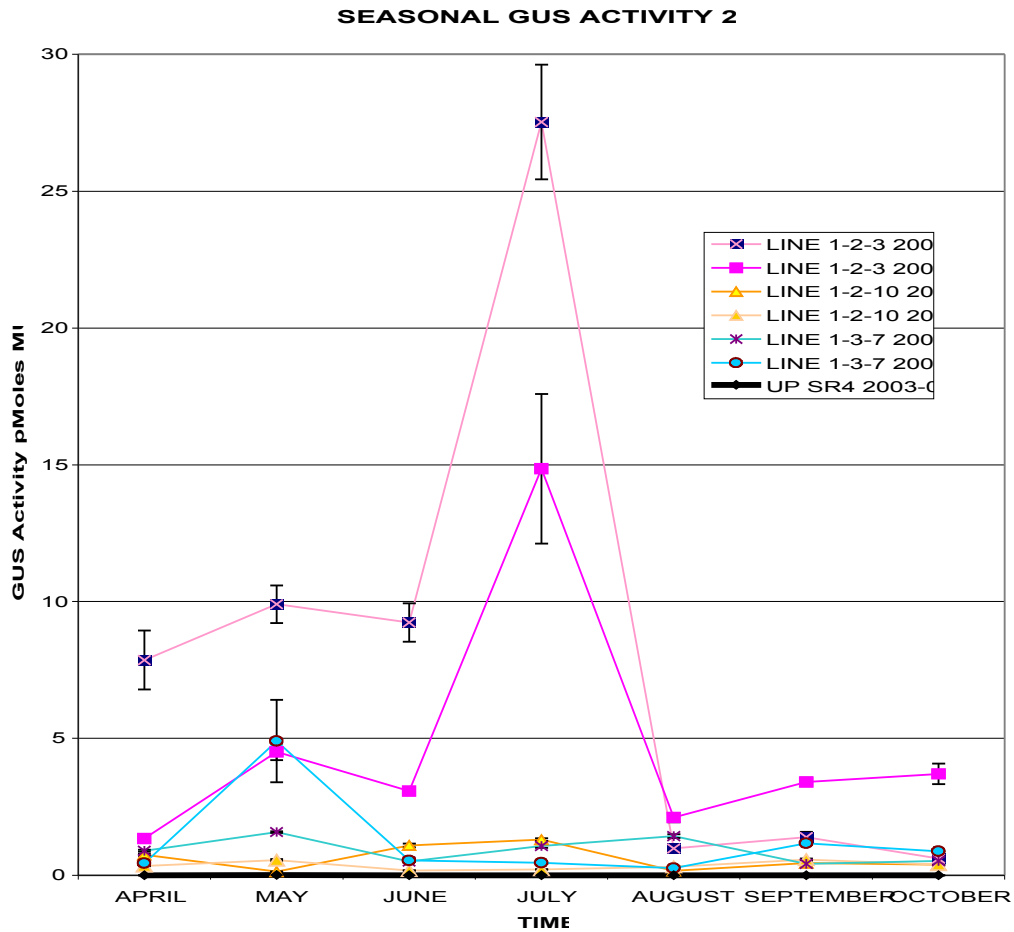
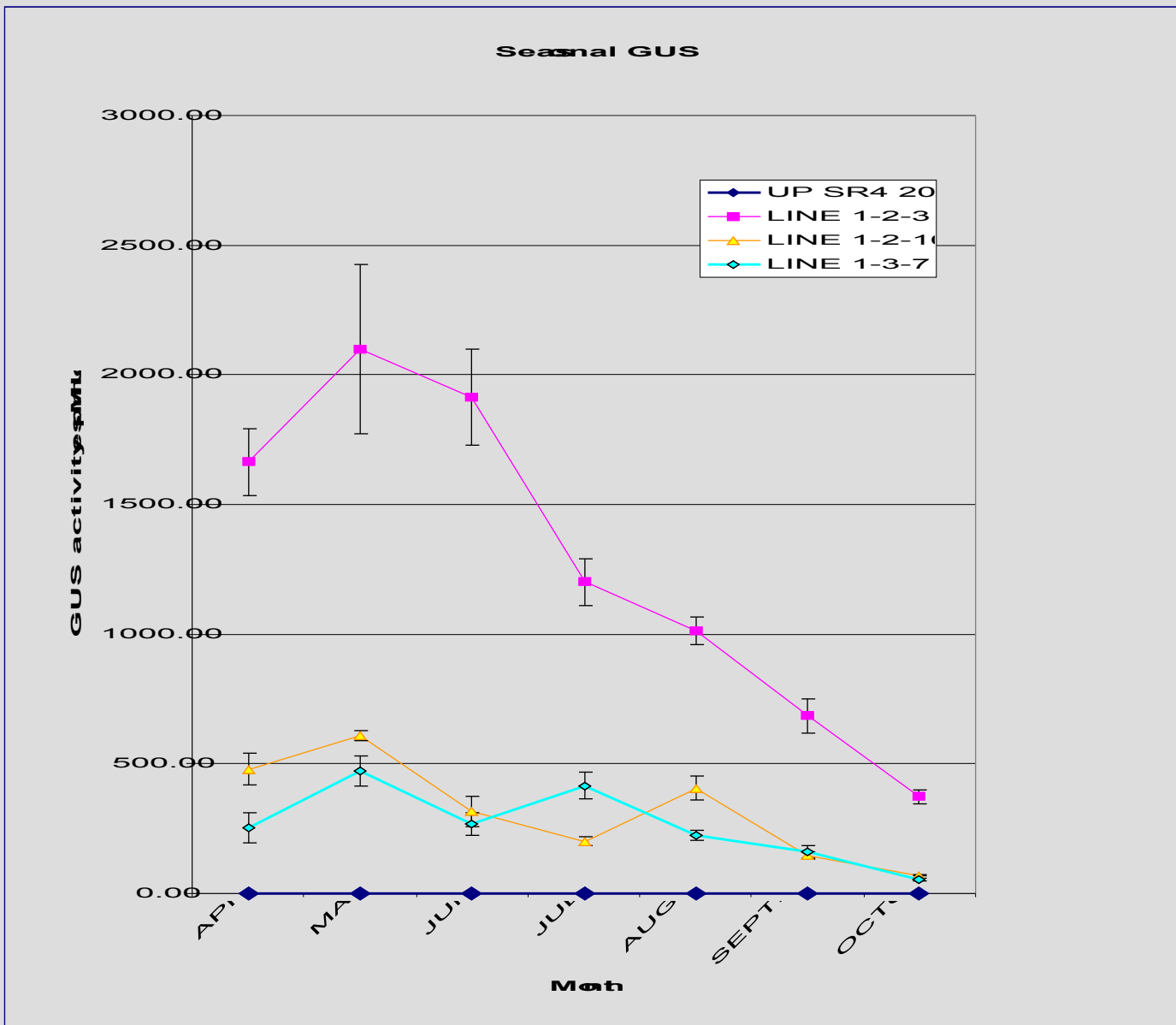


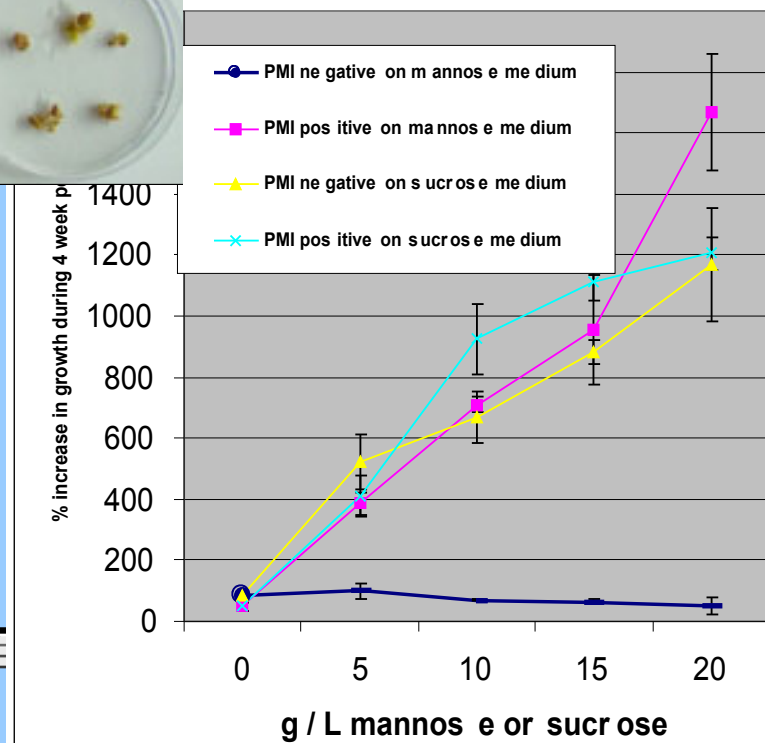
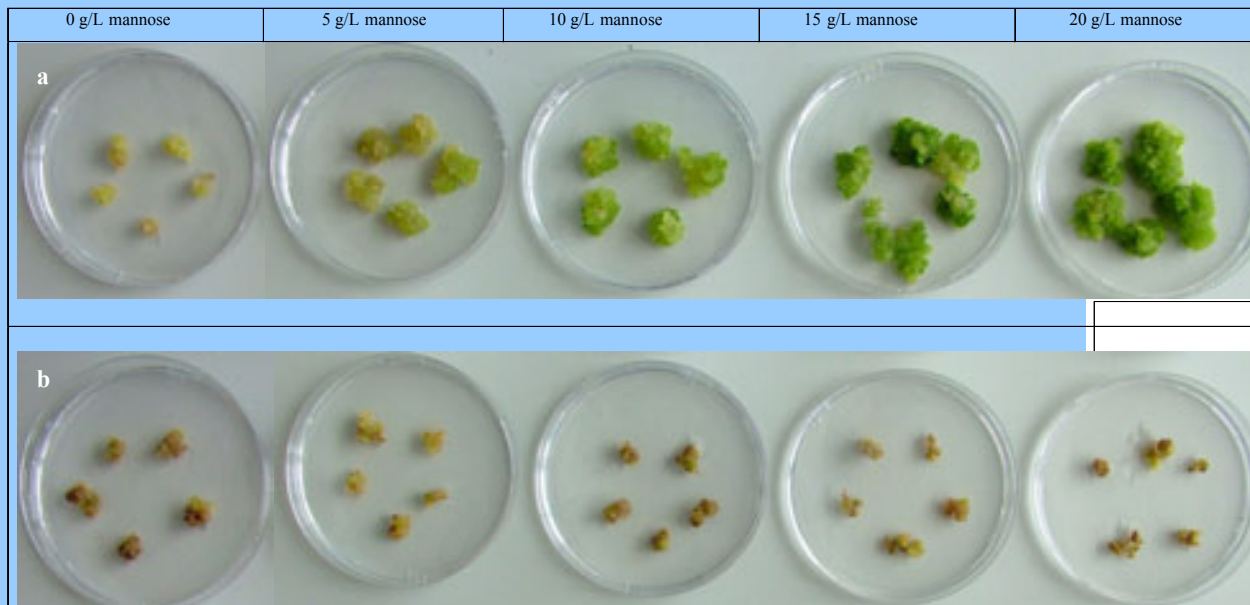
Figure 9 Seasonal Depression 2005



Other GM Elm Species

- ❖ *U. americana*, *U. glabra*, *U. parvifolia* shown to be suitable for *Agrobacterium* systems
- ❖ GUS histochemistry
- ❖ PCR products
- ❖ Shoot formation

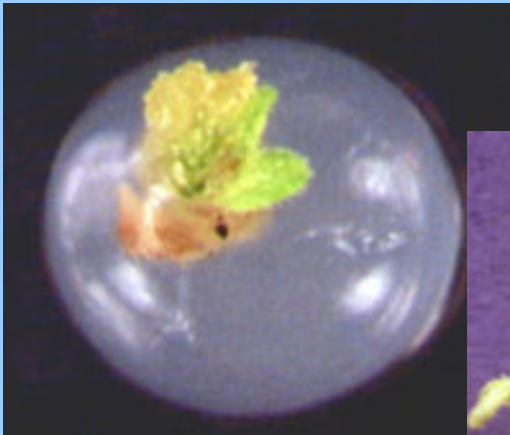
Figure 22 Growth of pmi positive transformant [(a)-upper] compared to pmi negative [(b)-lower] on phytagel solidified DKW media without sucrose but containing various levels of mannose.



Elm Cryopreservation

❖ Recovery from Freezing

❖ Shoot Regeneration



Elm Modifications

- ❖ Modify Internal Architecture: Ri-plasmid GM regenerants
- ❖ 3 Anti-Fungal Genes
- ❖ *Urtica dioica* Agglutinin UDA
- ❖ Dahlia Protein Dahlin D
- ❖ Radish Anti-Fungal Protein R
- ❖ GM Regenerants all three constructs
- ❖ Exploring alternative Anti-Fungals & Markers

AFP-Elm Regenerants

- ❖ *U. procera* plants obtained after AFP gene transfer









Fungal Challenge via Feeding Groove

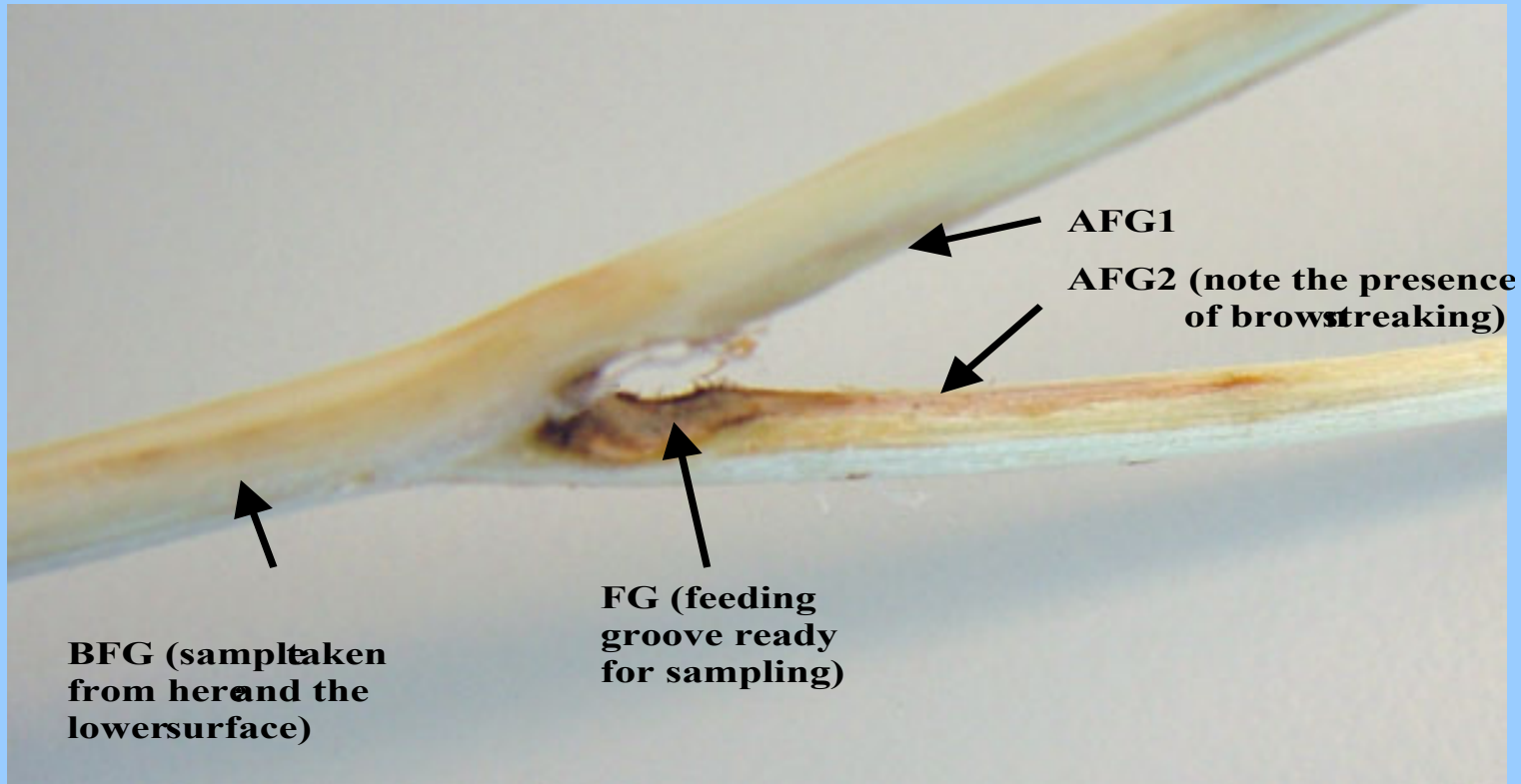


2.3.4 Raw data for feeding groove trials 2005-2006

Table 6 Key to Tables 7a-7g & 8a-8e

Plant Line	<i>Ulmus procera</i> SR4 line Genotype
Groove No.	Name given to identify feeding groove (N.B. The experiment was performed double blind and this identity was only un-encoded once the results had been assessed)
Diam. mm below groove	Twig Diameter (in mm) below Feeding Groove
Beetle present Yes or No	Was the beetle still present at the end of the two day period when the paper was removed?
Growth of inoculum from paper	Fungal growth from filter paper post infection on Malt Extract Agar
FG1&2	Sample taken from Feeding Groove
AFG1&2	Sample taken from above Feeding Groove
BFG	Sample taken from below Feeding Groove
Xylem Streaking	Brown Streaking of xylem evident on cutting and removal of bark layer (Graded I for small amount of streaking to III for a lot of streaking)
?	<i>Ophiostoma</i> growth
Both streaking and fungal growth together	? I light streaking ? II medium streaking ? III dark streaking

Figure 6 Brown streaking and fungal resolution sampling points shown on a stem piece following removal of bark prior to sampling



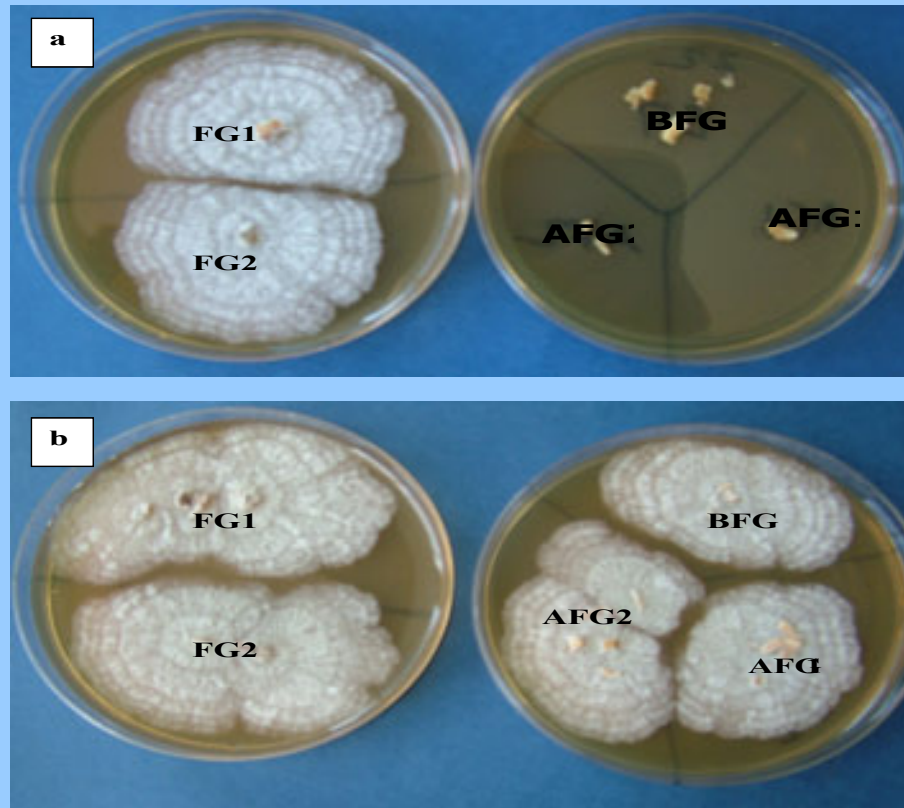


Figure 7 MEA plates containing hips from a) an antifungal gene containing line that exhibits resistance to the spread of the fungus compared with b) an untransformed, sensitive *P. coccinea*

Construct / Aspect	Fungus Isolated Above or Below FGroove	Brown Streaking Above or Below FGroove	Both Fungus Isolated & Brown Streaking	NO Fungus Isolated & NO Brown Streaking	Number Feeding Grooves
SR4	88.9%	44.4%	38.9%	19.4%	36
DP8	68.8%	27.1%	22.9%	29.2%	48
Dam6	63.9%	31.2%	23.0%	34.4%	61
U	52.6%	21.1%	15.8%	42.1%	19
R2425	45.5%	31.8%	13.6%	40.9%	22
C1(Ri)	60%	45%	35%	35%	20

Appendix 5

Genetic elements shown to be functionally effective

Cauliflower Mosaic Virus 35S promoter
Cestrum Mosaic Virus promoter
Nopaline synthase terminator
Phosphinothricin acetyl transferase/Bar
Neomycin phosphotransferase II
Ubiquitin 3 promoter
Green fluorescent protein (gfp)
Betaglucuronidase (gusA)
GusA intron
Rol A,B,C,D promoters
Rol AD
Nopaline synthase promoter
Nopaline synthase
Agropine synthase promoter
Agropine synthase
Agropine synthase terminator
Mannopine synthase
Tryptophan 2-monooxygenase
Indole 3-acetamide hydrolase
Isopentenyl transferase
Atriplex canescens untranslated region (UTR)
Cauliflower Mosaic Virus 35S promoter

Genotypes Available

- *Ulmus procera* SR4
- **SR4 + gusA, gfp, nptII, pmi**
- **SR4 + UDA**
- **SR4 + D**
- **SR4 + R**
- **SR4 C1**
- *Ulmus glabra*, *parvifolia*, *americana*
- *Sorbus aucuparia*
- *Quercus robur*
- *Salix* spp. Bowles' Hybrid, Jorr, Tora
- *Salix* spp. Dwarf Hybrids (x5)
- *Alnus glutinosa*
- *Populus* spp. hybrid

Elm Habitat Restoration



Acknowledgements

- ❖ Richard Irvine, Laura Marshall, Lorraine Gow, Amit Garg, Bob Crow, Angie McHugh
- ❖ Forestry Commission
Clive Brasier, Sue Kirk
- ❖ University of Abertay Dundee
- ❖ BioDundee
- ❖ Institute of Forest Biotechnology
- ❖ <http://www.forestbiotech.org>

