

Multi enzyme systems involved in astin biosynthesis and their use in heterologous astin production



Project acronym: MESIAB
Project no: EIB.10.004



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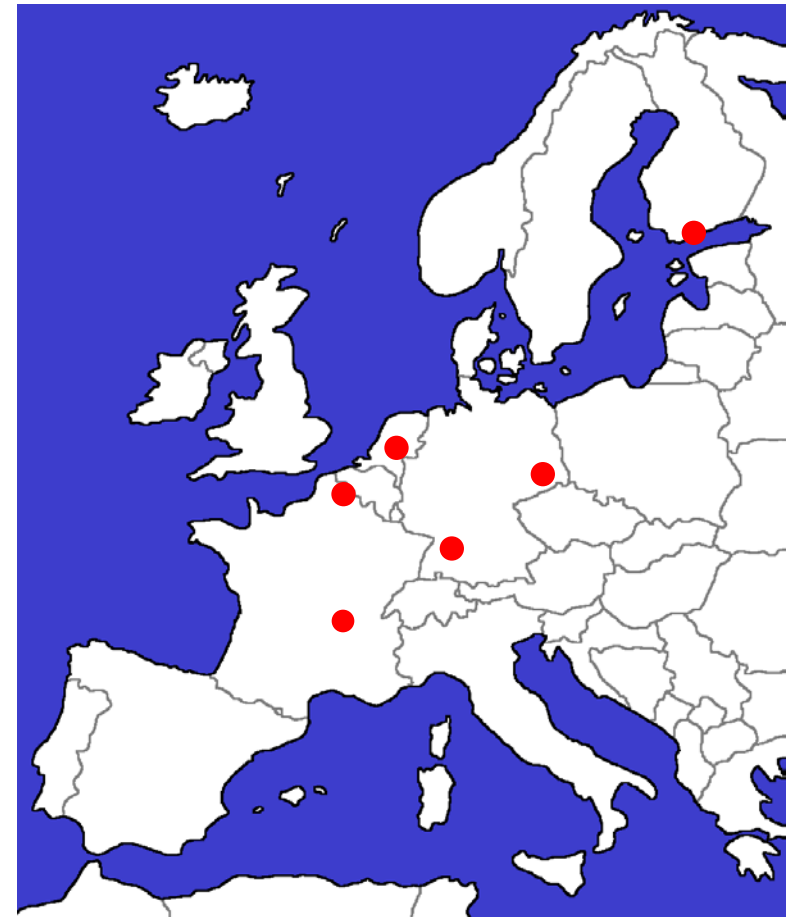
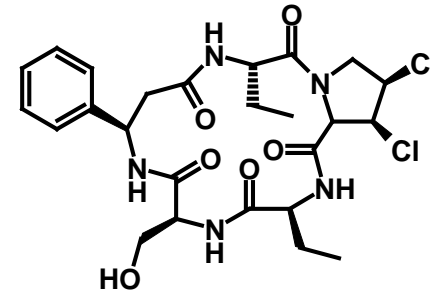


Fig. 1: <http://www.digitale-europakarte.de/europakarte.png>

Project aim

Enhancing the production of astins using molecular genetic tools and screening for new biological activities and novel applications of astins



Chemical structure of astin C

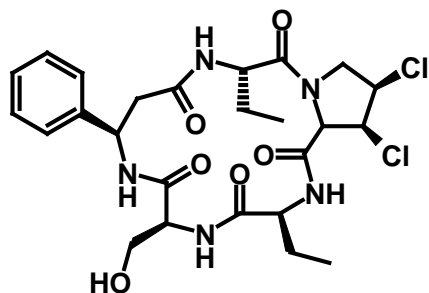
General project approach

Detection, cloning and heterologous expression of the individual genes involved in astin biosynthesis

Combining the genes into a cluster, expression of the cluster in *Streptomyces*, yeast and hairy root cultures

Screening for new biological activities and novel applications for astins using cell cultures and transcriptomic technologies

Detection of astins in *Aster tataricus*



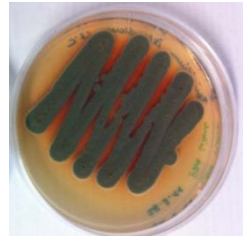
Chemical structure of astin C

Dried roots of *Aster tataricus*

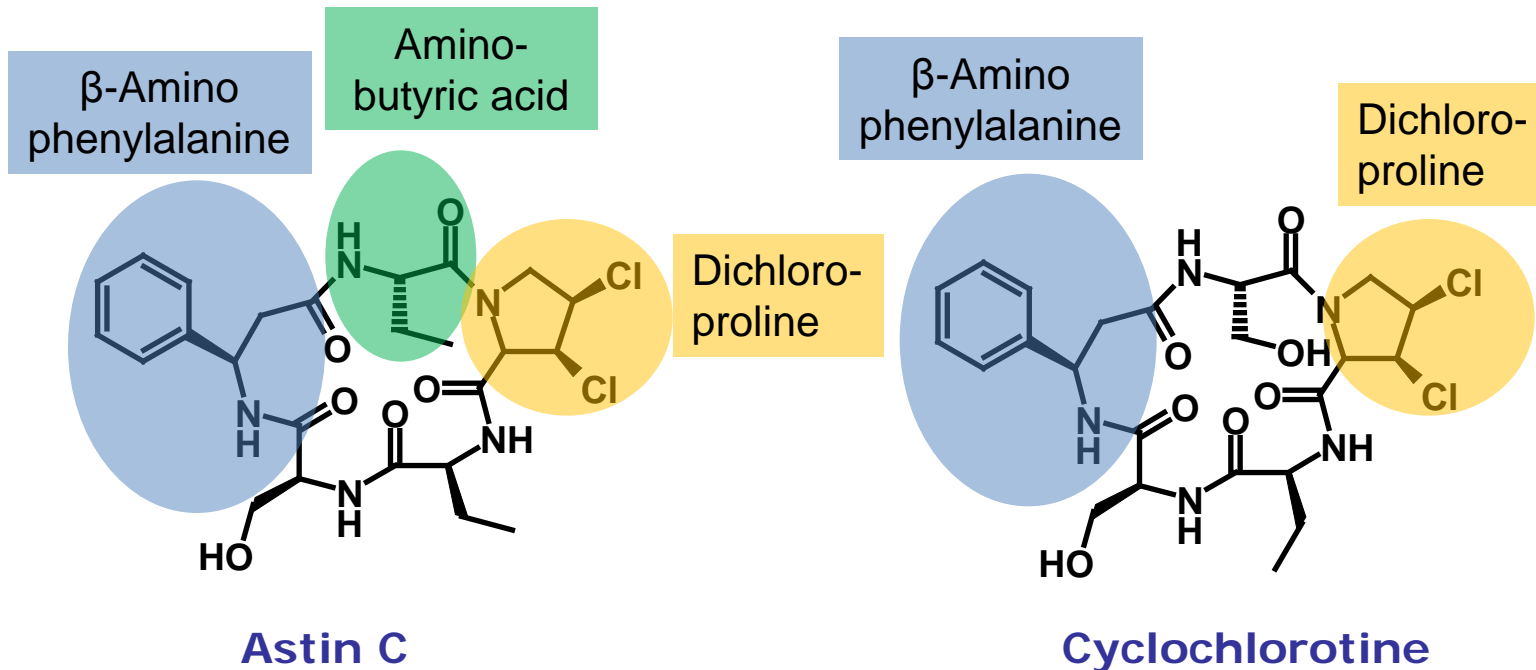
LC-MS detection methods were established for astins

- We have detected astins from dried root samples of *Aster tataricus*
- These are mainly dichlorinated, but some do not have the same mass as published astins

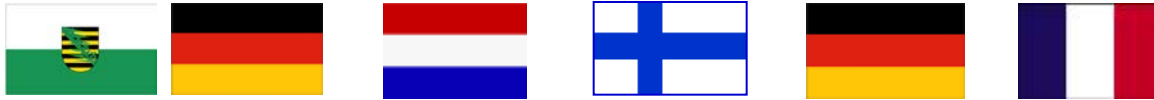
Cyclochlorotine from *Penicillium islandicum*



Penicillium islandicum



Detection of astin/ cyclochlorotine biosynthetic genes



- Both biosynthetic pathways can be assumed to require highly similar enzymes
- Working with *P. islandicum* should be faster and easier
- Thus, it was decided to sequence the *P. islandicum* genome and use the detected genes for searching for the corresponding genes in *A. tataricus*

P. islandicum sequence analysis



→ Candidate genes are present, but not clustered



Macrocyclisation of NRPS

Bacteria

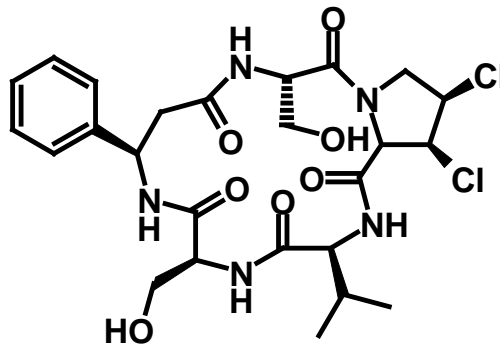
Du and Lou, 2009, Nat Prod Rep

Fungi

Gao et al., 2012, Nature ChemBiol

Te

CT



Cyclochlorotine

Specific signature

Present, ending the NRPS?

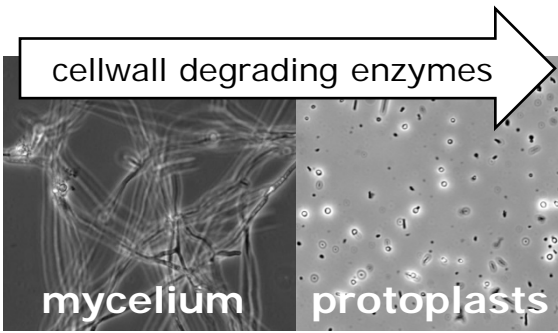
Literature data show that a terminal condensation (CT) domain is required for macrocyclisation of cyclic nonribosomal peptides in fungi

This information means that the NRPS involved in cyclochlorotine production will most likely contain a CT domain



1 Protoplasting successful

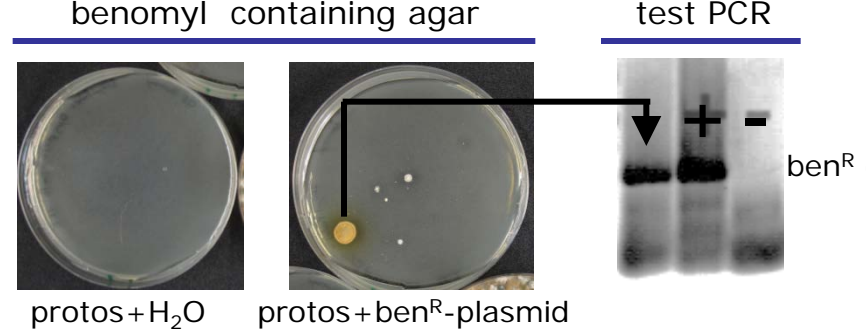
cellwall degrading enzymes



mycelium protoplasts

2 Protoplast-transformation successful

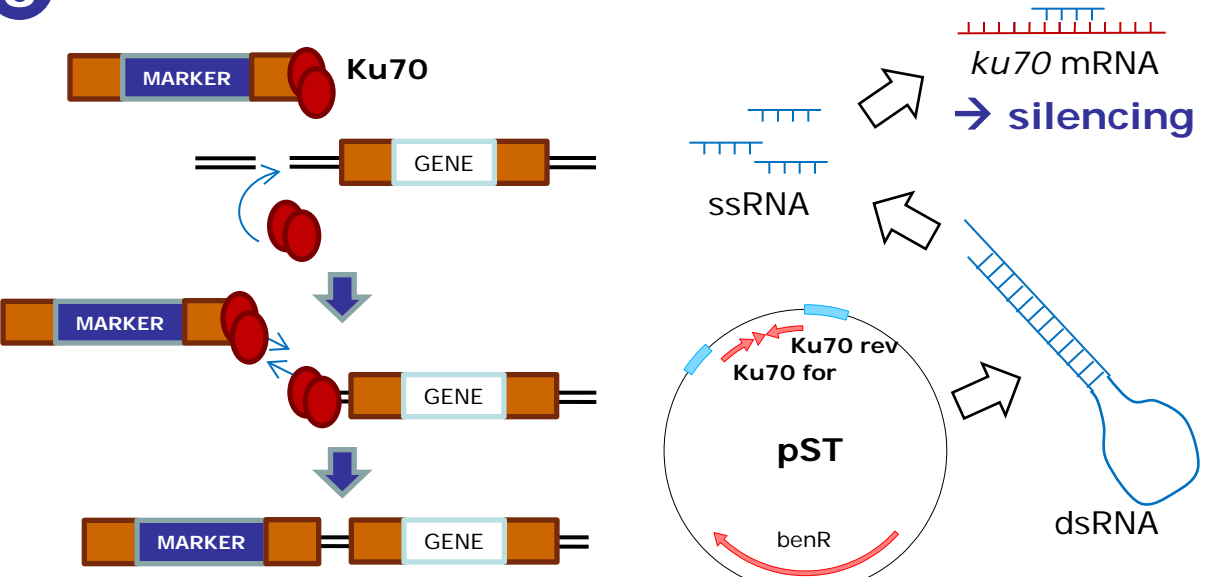
benomyl containing agar test PCR



protos + H₂O protos + ben^R-plasmid

ben^R gene

3 NHEJ deficient strain established



non-homologous end joining (NHEJ)

Ku70

MARKER

GENE

ssRNA

ku70 mRNA

→ silencing

dsRNA

Ku70 rev

Ku70 for

pST

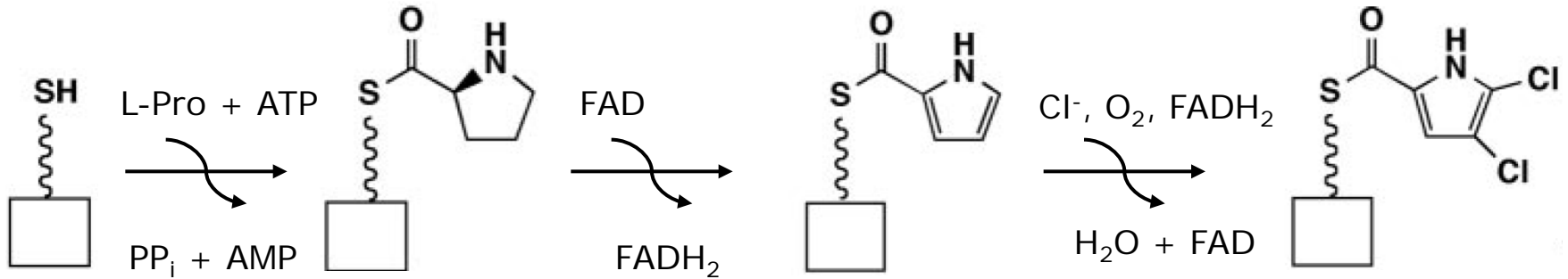
benR

Ku70 gene activity (qRT-PCR)

Strain	Ku70 gene activity (qRT-PCR)
WT	1.0
clone 1	~0.18
clone 2	~0.15
clone 3	~0.12
clone 4	~0.48

-90%

Prolyl dehydrogenase (PRODH)



peptidyl carrier

protein

proline

adenyltransferase

acyl-CoA

dehydrogenase

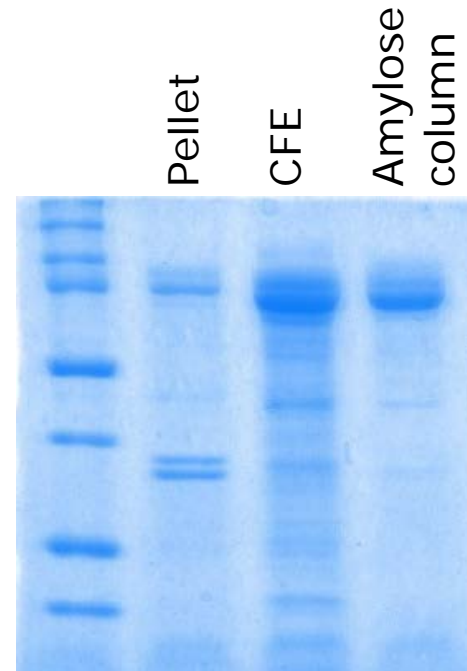
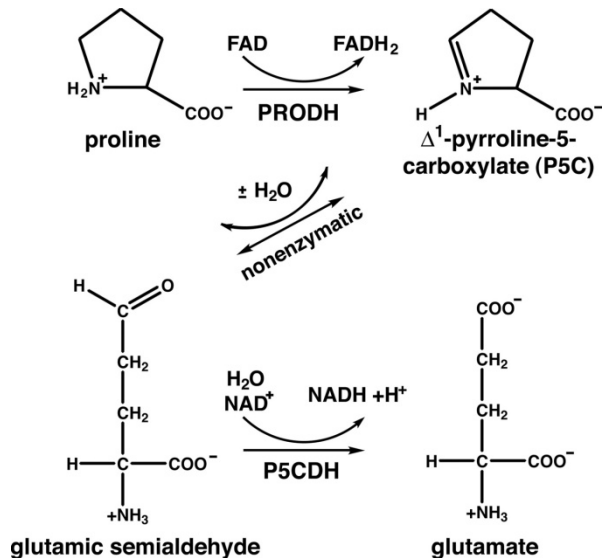
FADH₂-dependent

halogenase

Prolyl dehydrogenase (PRODH)

Thermophilic proline dehydrogenase

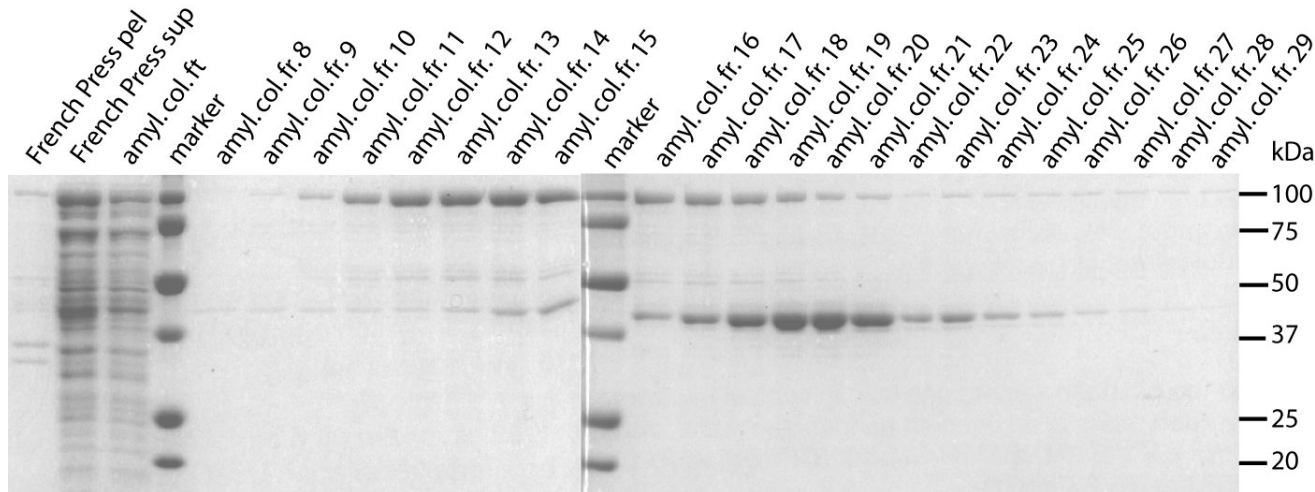
- PRODH from *Thermus thermophilus*
- High overproduction of MBP-tagged flavoenzyme
- Active as fusion and after removal of MBP tag
- Enzyme fully stable at 80 °C



Prolyl dehydrogenase (PRODH) Fungal proline dehydrogenase



- 3 Putative PRODH genes in *Penicillium islandicum*
- One PRODH produced in soluble form
- One-step purification of MBP-tagged flavoenzyme
- Active as fusion and after removal of MBP tag



First fungal prolyl dehydrogenase



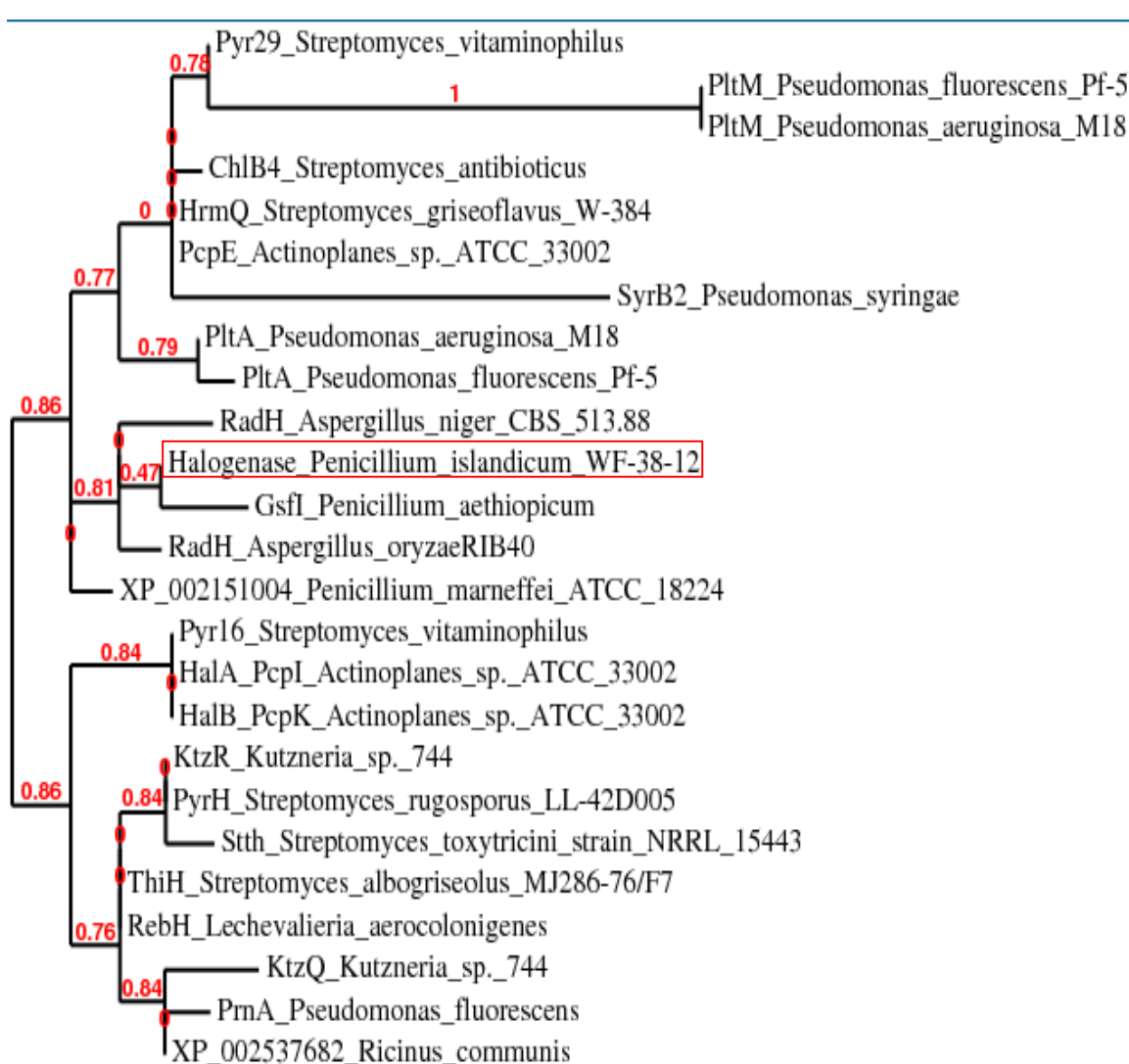
Flavin-dependent halogenases



pyrrole
halogenases

putative
halogenases
from fungi

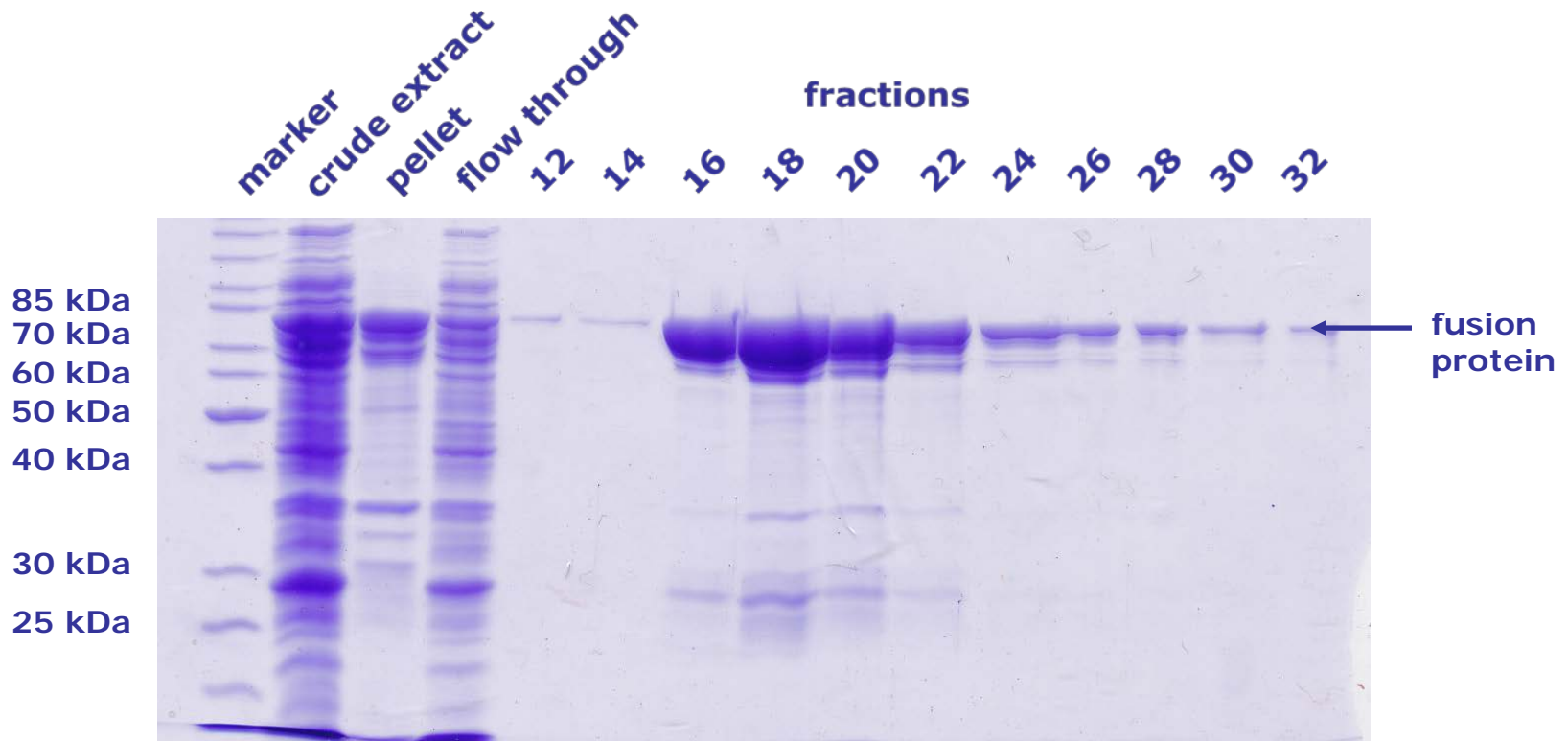
tryptophan
halogenases



Halogenase from *P. islandicum*

Purification of the halogenase

50 μ M IPTG, over night at 20 °C
GST fusion protein: 83.6 kDa



Halogenase can be purified using a maltose-binding protein tag.



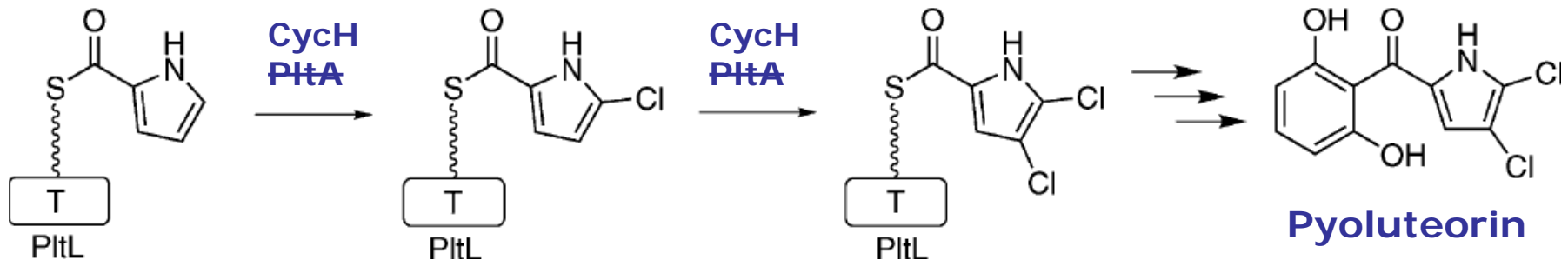
Demonstration of halogenase activity

Activity assay using a chemically synthesised substrate mimic has not been successful, yet.



Complementation of pyoluteorin biosynthesis in *Pseudomonas fluorescens*.

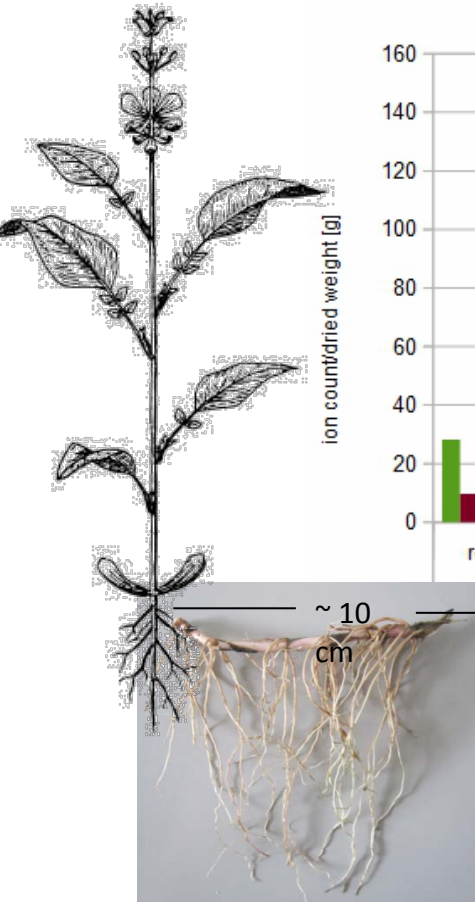
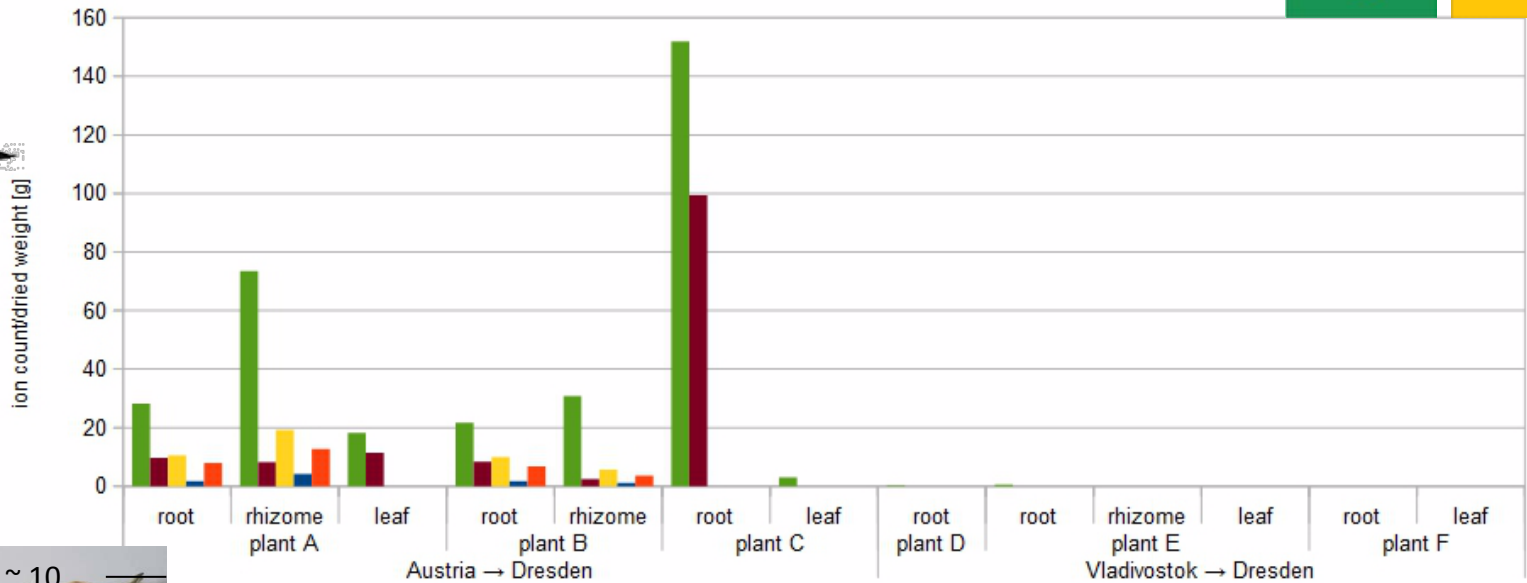
The pyrrole halogenase gene *pltA* is inactivated and will be substituted by the halogenase gene *cycH* from the cyclochlorotine producer *P. islandicum* to show halogenating activity *in vivo* by complementing pyoluteorin biosynthesis.



Localisation of astins in *A. tataricus*

Astin composition and concentration

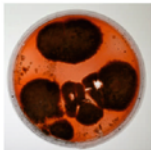
■ astin A/B ■ astin C ■ astin F ■ astin H/E ■ astin G



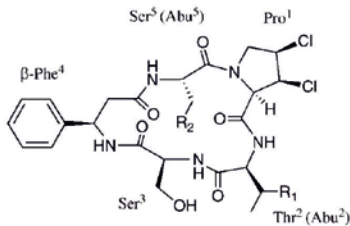
→ Presence of astins depends on plant origin

Astins, plant or fungal metabolites?

P.islandicum (10 strains) ●



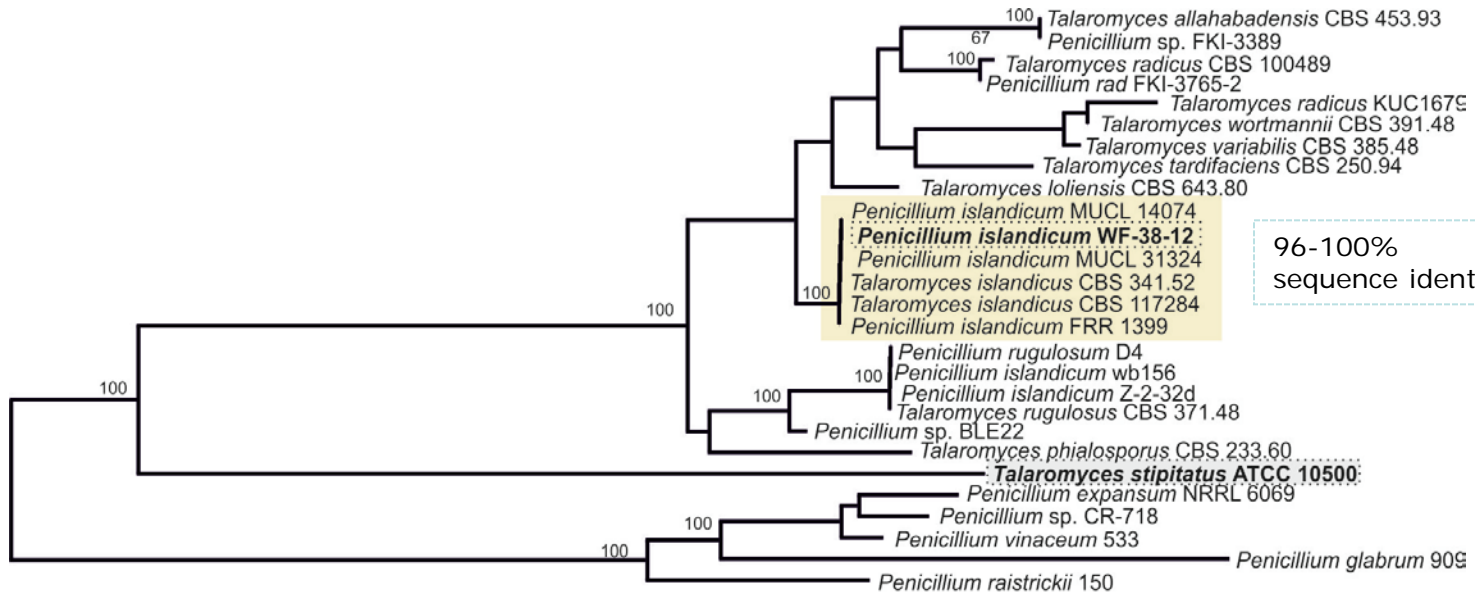
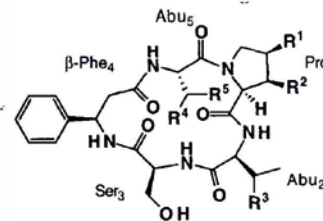
cyclochlorotines



A.tataricus (2 plants) ●

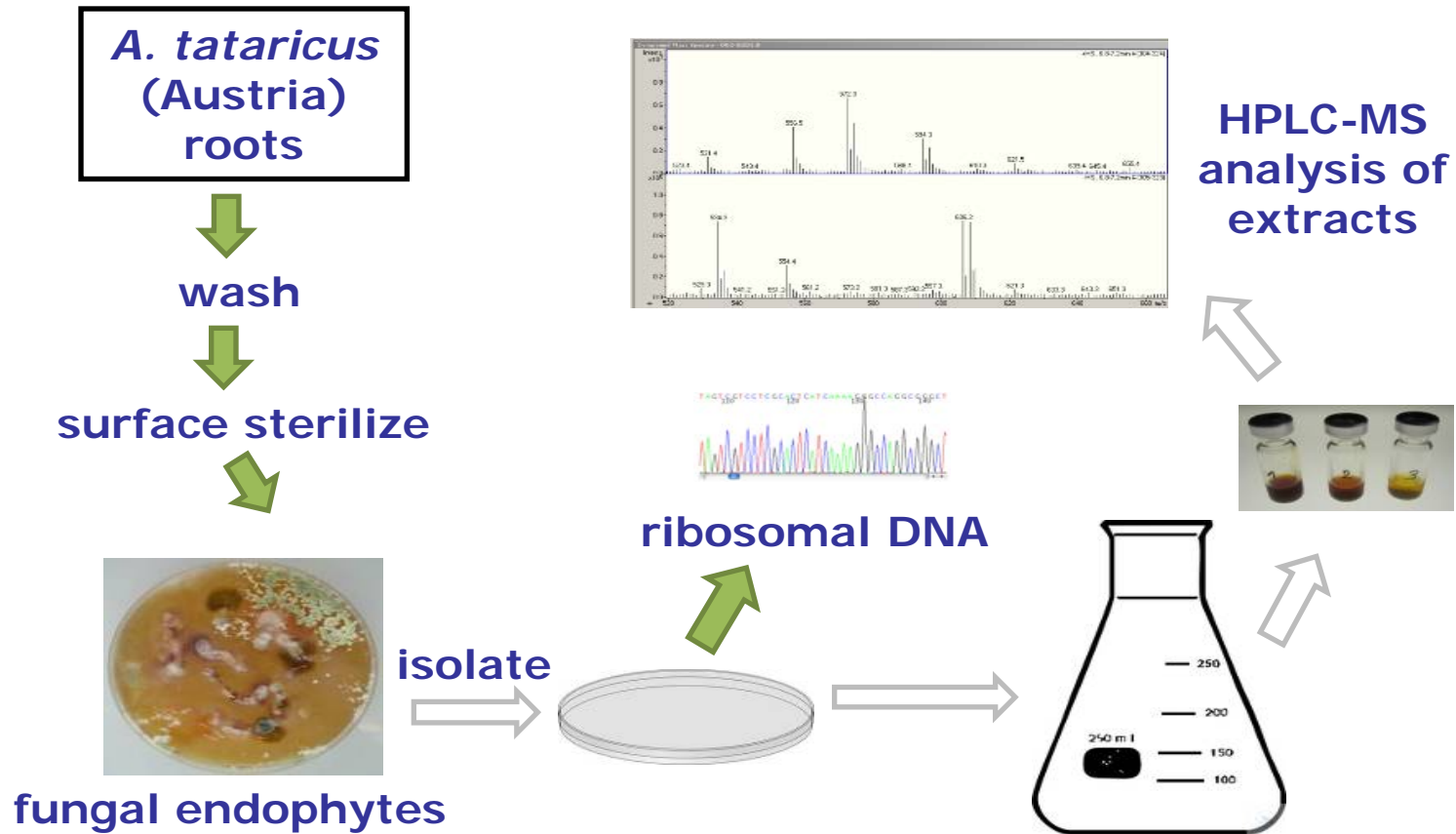


astins



Cyclochlorotines were identified from strains of *P. islandicum* isolated from soil or spoiled food; none of them produces astins.

Astins, plant or fungal metabolites?



Astins could not be detected in any of the fungi isolated from *A. tataricus* roots.

Generation of sterile plants

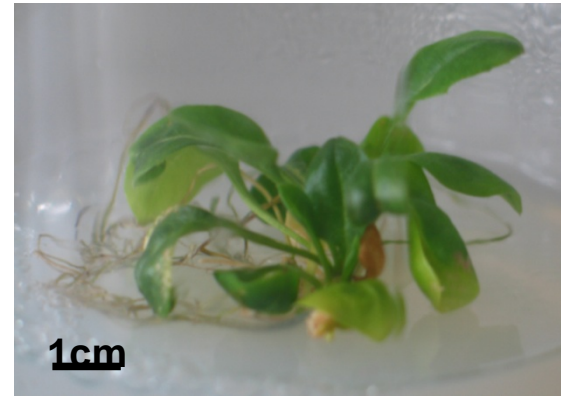


Is there astin production in sterile plants?

Cultivation of sterile seeds on MS medium without any hormones

Cultivation of sterilised inflorescence tissue on MS medium (naphthyl acetic acid or benzyl amino purine) to induce regeneration of small, sterile plants

These plants turned out to be not really sterile!

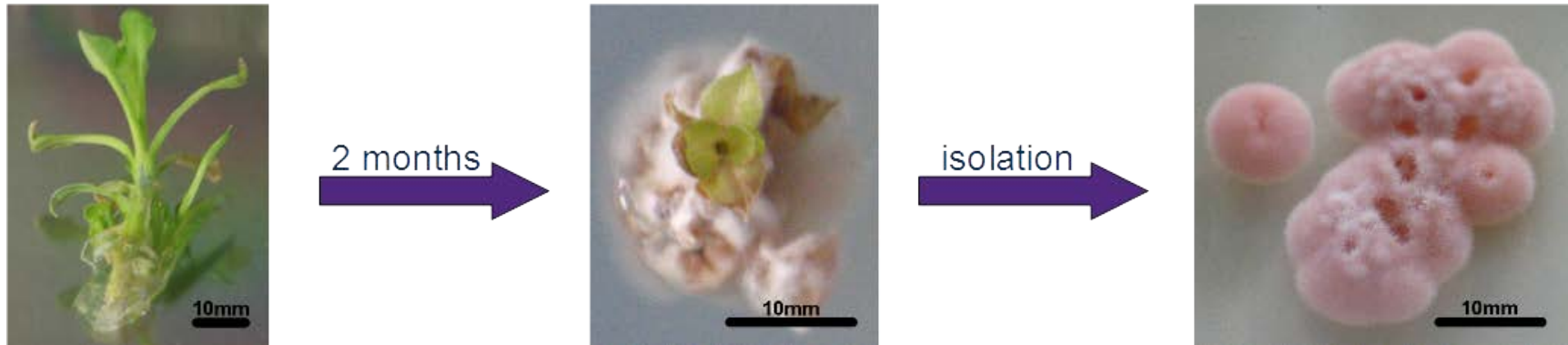


Sterile plant from sterilised seeds

Germinating plant from inflorescence axis on MS medium + 1µg/ml BAP



Growth of "sterile" plants



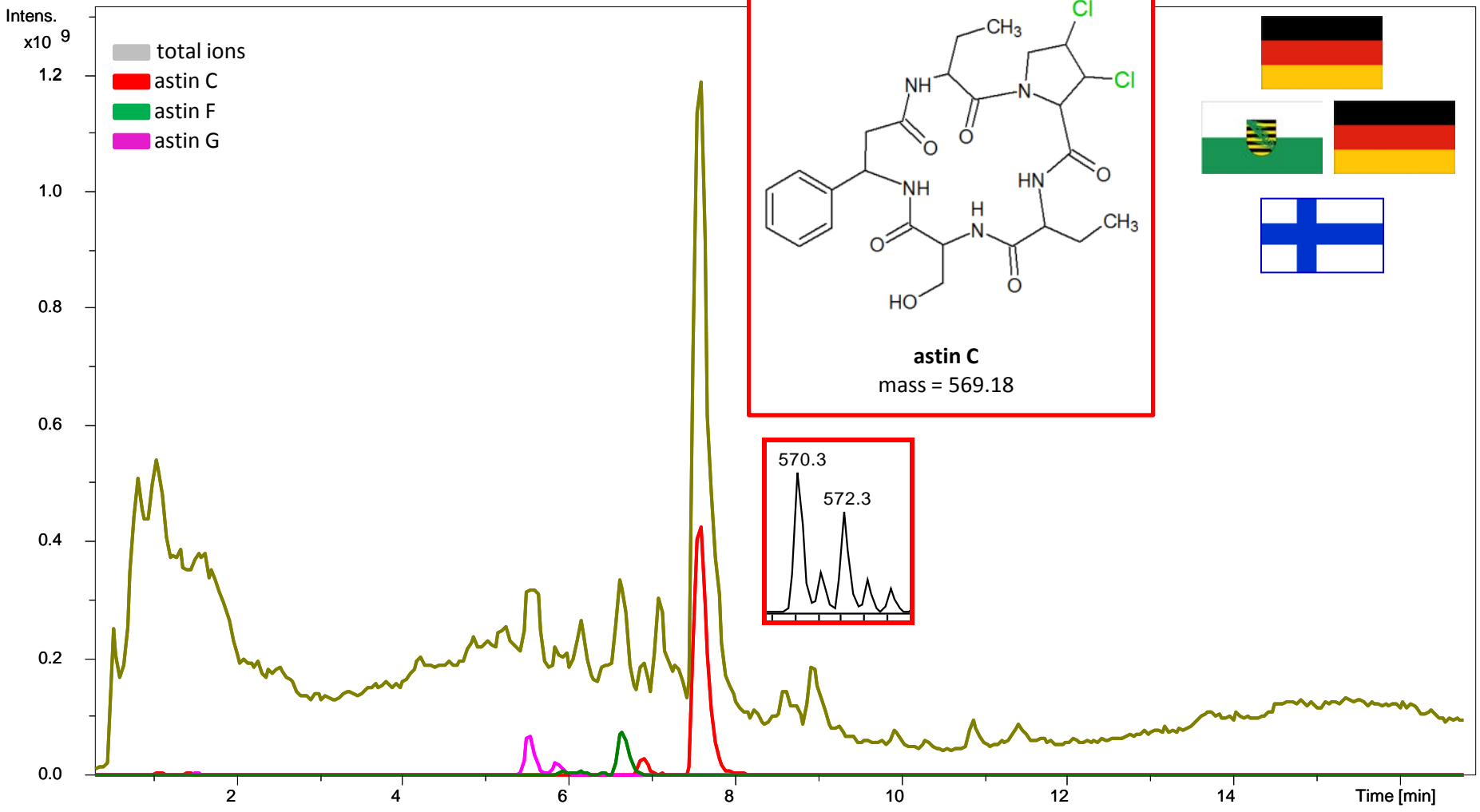
Generation of sterile *in vitro* cultures of *Aster tataricus* (Austrian) via hormone treatment (auxin and cytokinin for inducing shoot and roots)

After a few months, growth of a pink fungus could be observed. This fungus could not be removed by treatment with antimycotica like amphotericin.

Isolation of a novel fungus

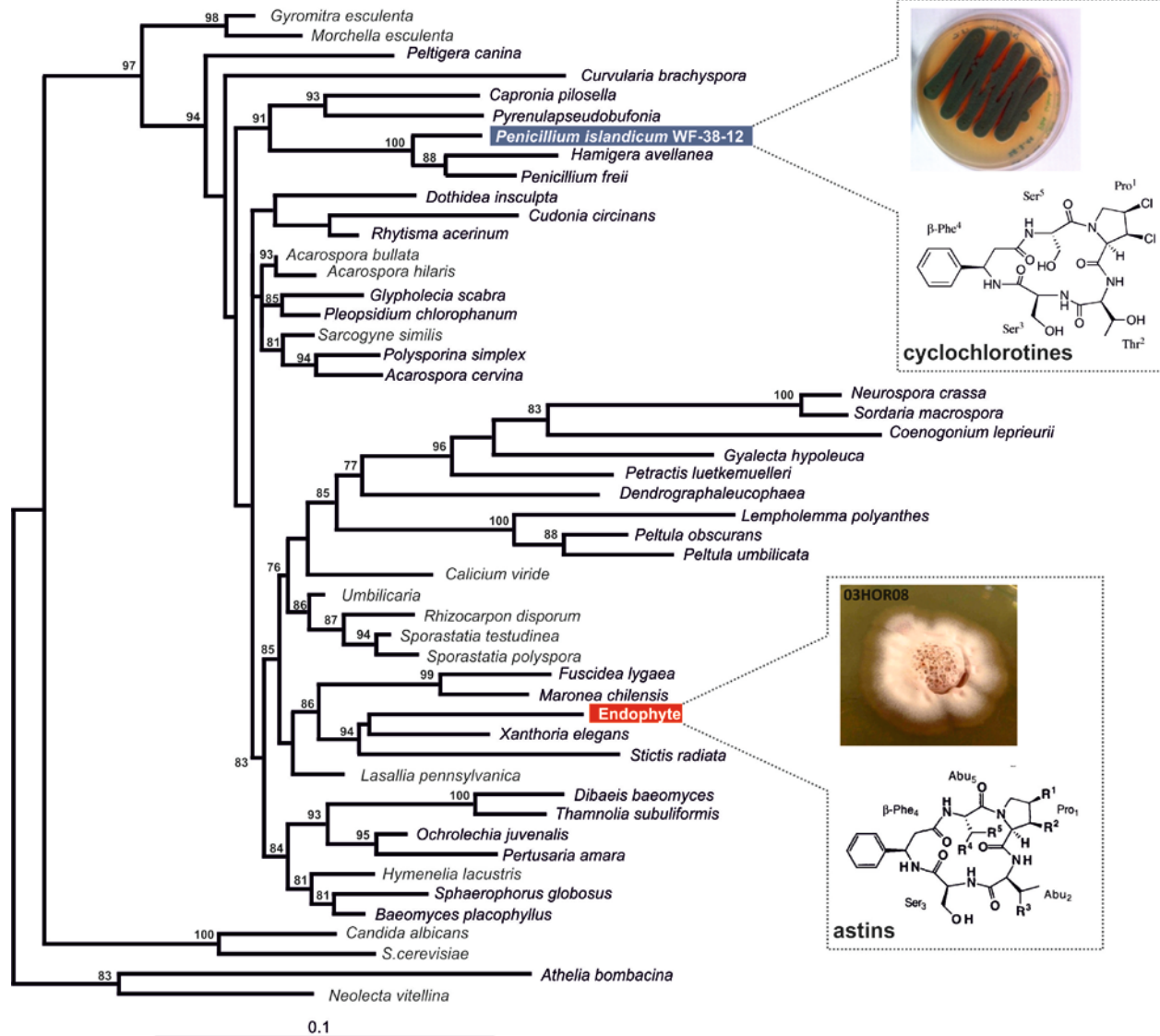


Production of astins by the novel fungus



Astin C is the main compound. Astins produced by the new fungus are identical to those isolated from dried *A. tataricus* roots.

An endophyte which produces astins



Astins were identified in a fungal endophyte isolated from *A. tataricus* which is not closely related to *P. islandicum*.

Genome sequencing of endophyte

**Astin producing
endophyte**

vs.

Penicillium islandicum



genome parameters

length: 28 MB
GC content: 53 %
coverage: 57 fold
scaffolds: 395

length: 34 MB
GC content: 45,22 %
coverage: 41 fold
scaffolds: 330

predicted secondary metabolite gene clusters (by antiSMASH)

PKS: 13
NRPS: 5
(1/5 is 5-modular)
NRPS-PKS: 1
terpene: 5
other: 9
sum 33

PKS: 19
NRPS: 17
(2/17 are 5-modular)
NRPS-PKS: 8
terpene: 4
lantipeptide 1
other: 11
sum 60

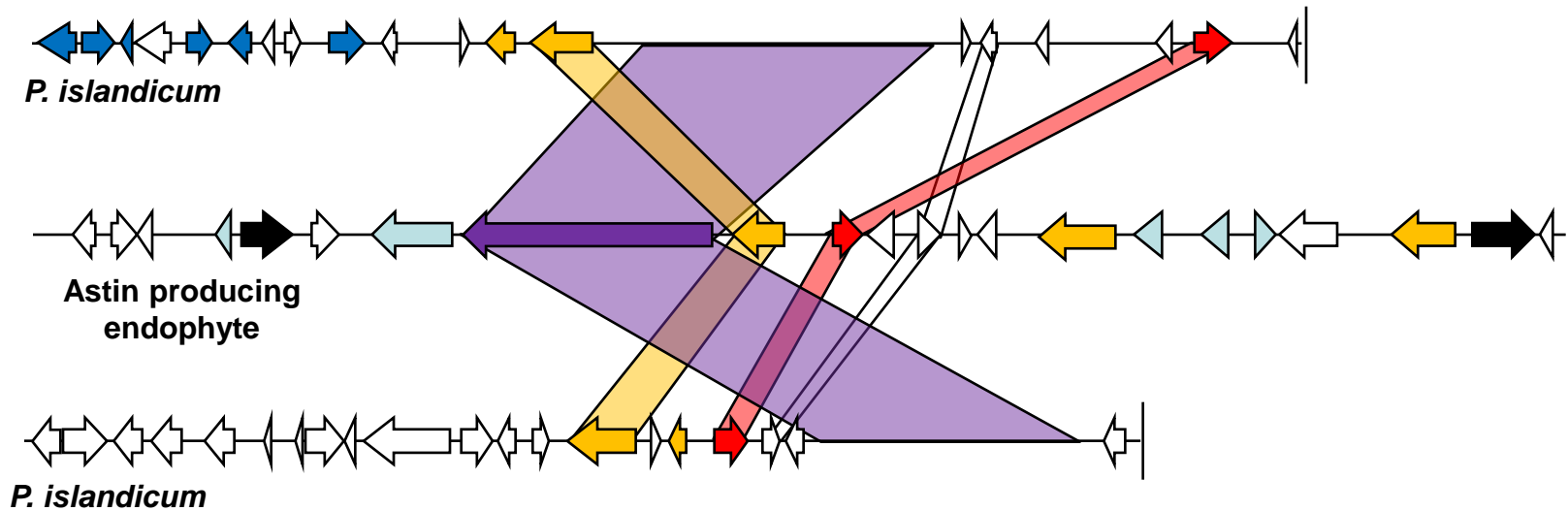
Partial astin gene cluster



Comparison of gene clusters



Genes with high similarity

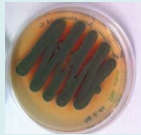


There is a gene cluster present in the new fungus with high similarity to two gene clusters in the cyclochlorotine producer *P. islandicum*

Summary



Modified proposal



Cyclochlorotines
P. islandicum

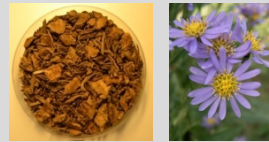
Genome sequencing

Identification of NRPS
gene cluster

Cloning of
individual genes

Enzymatic activity
and gene inactivation

Proposal



Astins
A. tataricus

Identification of individual
biosynthetic genes

Expression and
enzymatic activity

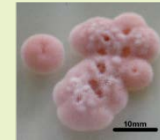
Construction of gene cluster

Heterologous expression

Production of astins

Bioactivities

What we achieved



Astins

Isolation of a novel
fungus producing
astins

Draft genome
sequence

Astins can now be
produced by
fermentation