



Dyadic Netherlands

Developing industrial enzymes and proteins

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Research Subsidiary of Dyadic Inc, USA

- Dyadic Nederland BV, Wageningen, The Netherlands
- www.dyadic.nl

Focus

- Discovery and development of enzymes for the bioenergy, food/feed, and paper and pulp industries.

Tools

- Fungal molecular biology
- Genomics
- Enzymology
- Fermentation technology



Team Dyadic Netherlands



Management: Wim van der Wilden

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Rienco van der Mooren

Marianne Cramer

Consultant: Jaap Visser

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Dyadic's Commercial Enzyme production platforms



Enzyme producing fungi currently employed by Dyadic:

Fungus	Current Market	Development
<i>Myceliophthora thermophila C1</i>	<i>Textile</i>	<i>Biofuels, Feed/food, P&P, specialty proteins</i>
<i>Trichoderma sp.</i>	<i>Feed, paper&pulp, textile</i>	<i>P&P, Feed, Biofuels</i>

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C1: developed into an efficient protein production system

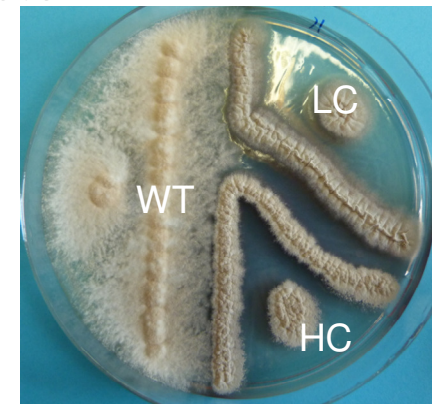


Myceliophthora thermophila C1 (formerly *Chrysosporium lucknowense C1*)

- Isolated from soil in eastern Russia as producer of neutral cellulases.
- Developed into a proprietary mature enzyme production system.

Main features:

- Low viscosity, fermentation to very high densities
 - High production levels (up to 100 g/L protein), easy scaling
 - Versatile genetic tools and hosts developed
 - Genome sequenced and annotated
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- Self affirmed GRAS status of a C1-cellulase product was acknowledged by the FDA (2009)



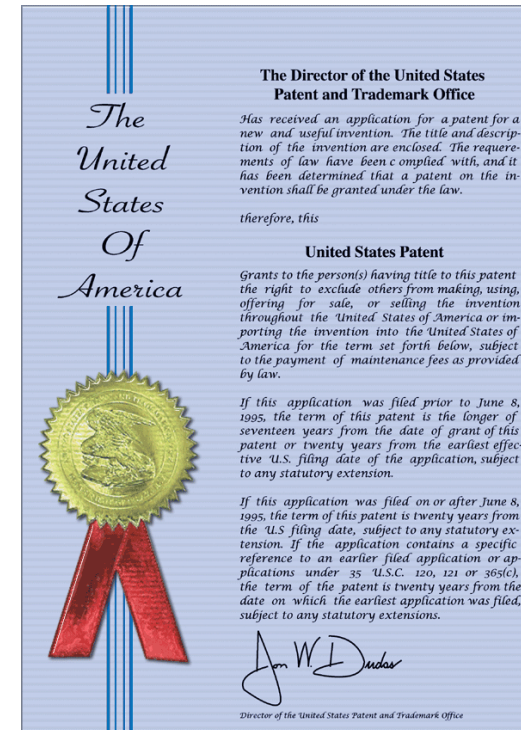
Intellectual Property



Dyadic fully owns the C1 fungus and technology:

- 10 US Patents granted
- 10 US Patent applications pending
- 74 Foreign patents issued
- 23 Foreign patents pending

Large freedom to operate



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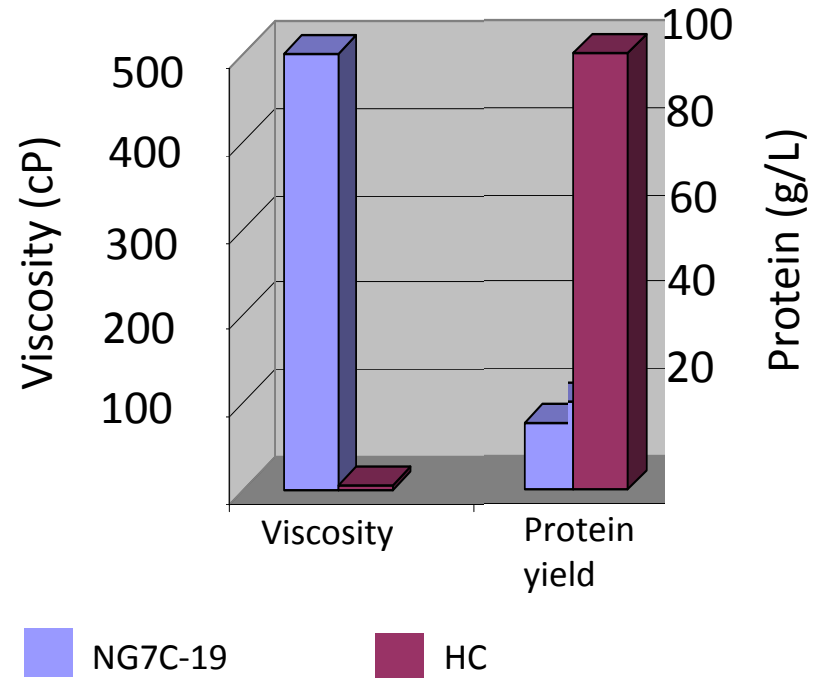
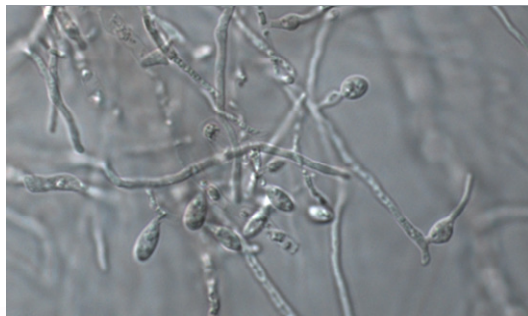
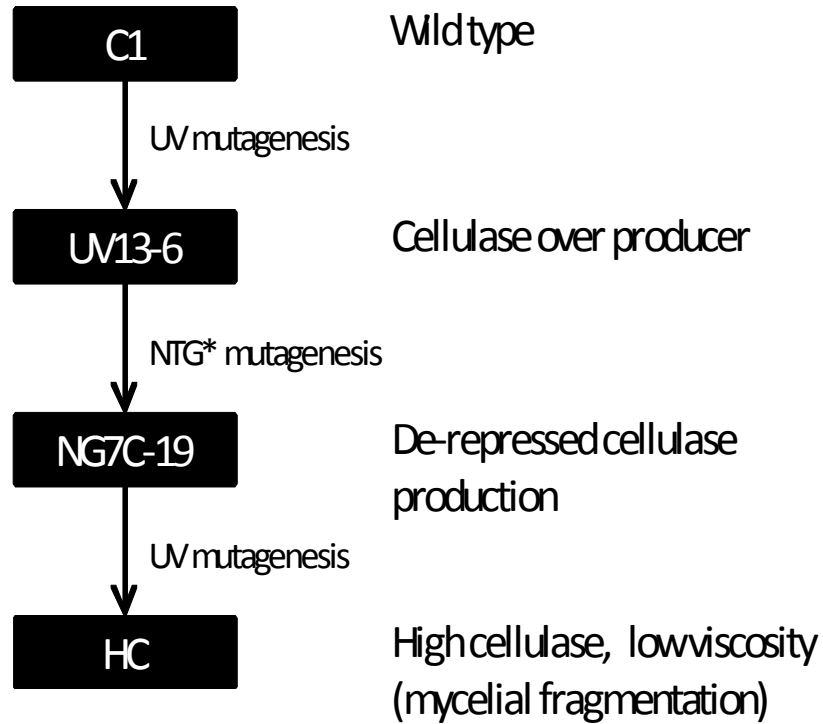
Steps towards a mature enzyme production system



1. Development of protein hyper-producing strains
2. Development of versatile genetic tools and host strains
3. Exploration of the enzymatic potential by genomics
4. Construction of tailored strains for desired enzymes and enzyme mixtures



Development of protein hyper-producing strains



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Development of versatile genetic tools and strains



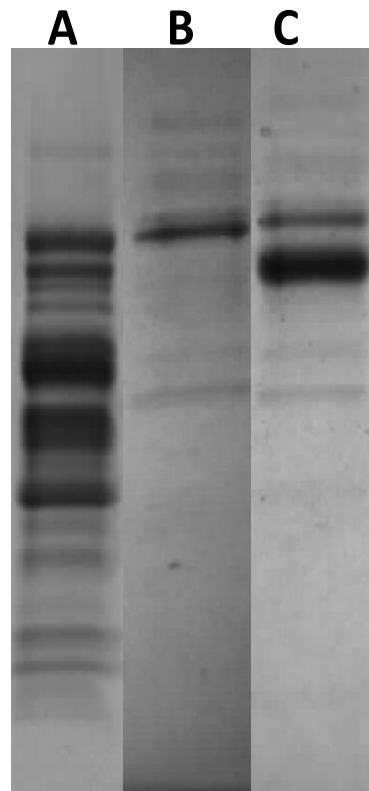
Genetic tools developed

- Transformation system: high efficiency, stable integration (stable progeny).
- Several genetic markers available: auxotrophic and dominant
- Allows for multiple rounds of transformation
- Gene expression: variety of expression signals.
- Constitutive, inducible at various strengths
- Protein production: efficient secretion signals.
- Targeted gene disruption: efficiency up to 90% locus specific integration.
- Based on self cloning: No foreign DNA needed



Development of versatile genetic tools and strains

- Construction of “clean” background strains (LC-strains)
- For production of relatively pure single enzyme



← Individual target enzyme

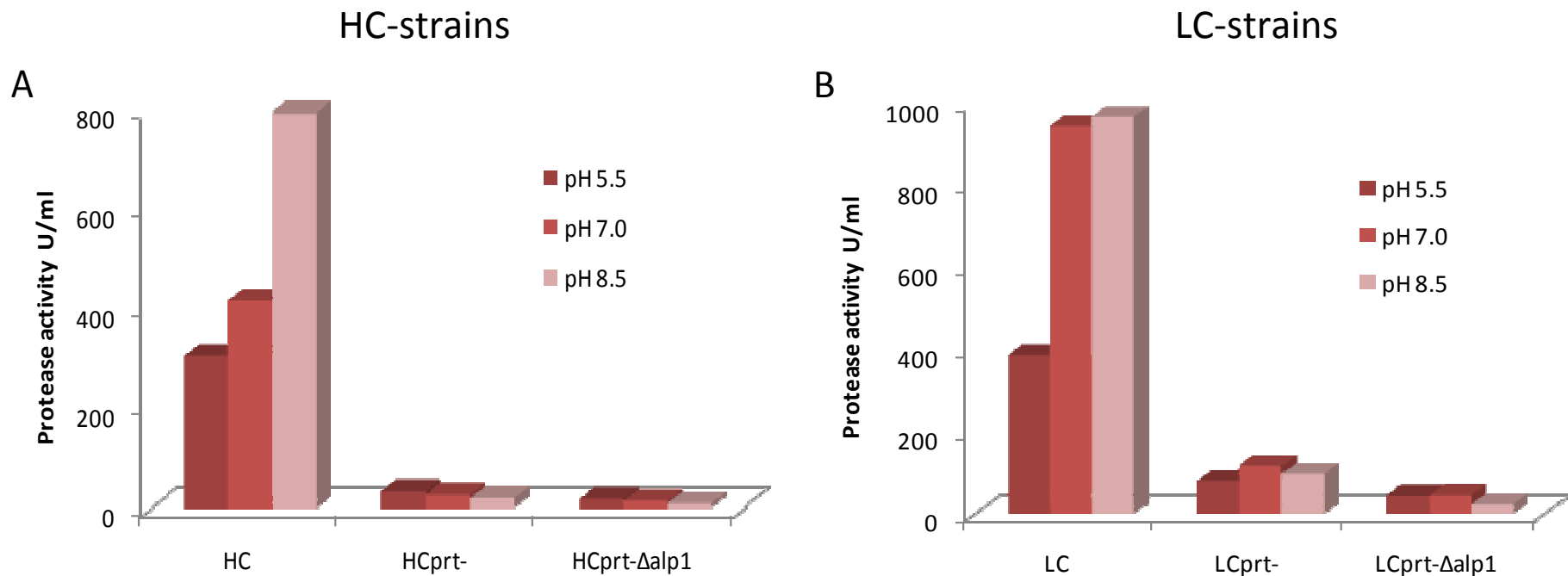
- (A) Baseline C1-strain:
High cellulolytic activities
Diverse enzyme mixture
Up to 100 g/L total protein
- (B) Low cellulase background strain (LC):
Almost no cellulolytic activities
Very few endogenous secreted
Suited for enzyme characterization
- (C) ~ Up to 80% target protein



Development of versatile genetic tools and strains



Host strains with low protease activity developed

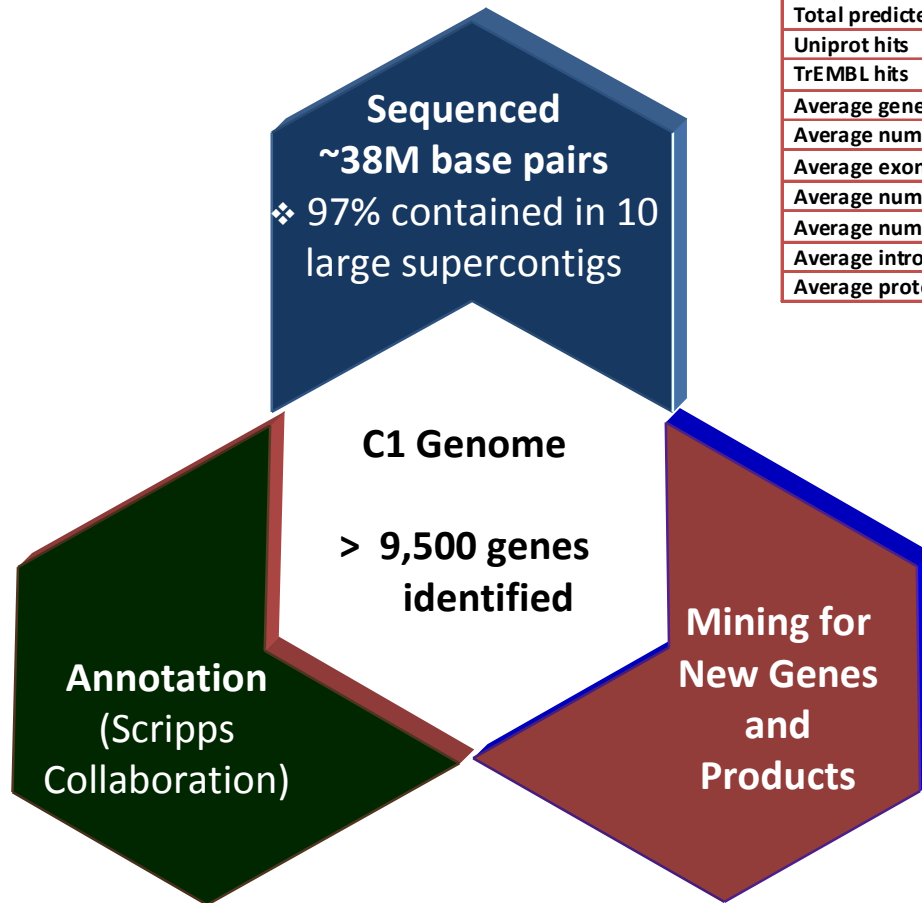


Particularly important for production of heterologous enzymes

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Exploration of the C1-enzymatic potential by genomics



Feature	Genemark	Geneid	Glimmer
Total predicted genes	9807	16012	8545
Uniprot hits	3151	4234	3831
TrEMBL hits	5007	3813	2856
Average gene length ¹ (bp)	2223	2271	2313
Average number of exons per gene ¹	3.47	3.66	3.70
Average exon length ¹ (bp)	549	499	503
Average number of introns per gene ¹	2.79	3.10	2.91
Average number of introns per gene (incl. 0) ¹	2.47	2.66	2.70
Average intron length ¹ (bp)	130	169	169
Average protein length ¹ (aa)	636	609	621

A large number of genes putatively encoding **industrially important** enzymes discovered:

- ~250 Carbohydrate-active Enzymes (CAZy)
- ~150 proteases
- ~700 oxido-reductases
- ~75 lipases / esterases.

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Exploration of the enzymatic potential by genomics



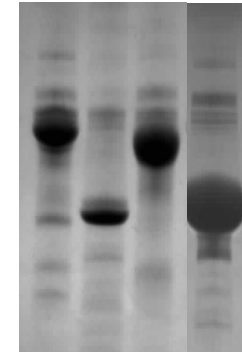
Opportunities

1. Identification of new targets for commercialization.

- Cellulases, xylanases, amylases, proteases, lipases, phytases
- Via enzyme library construction and analysis

2. C1-strain improvement

- Identification of key genes involved in expression, secretion, post-translational modification, other aspects of physiology.



Generating leads for new/improved products

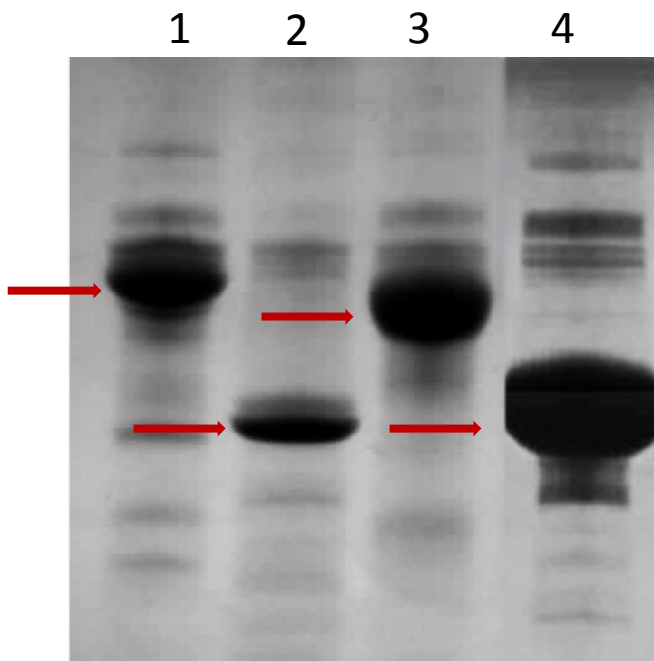
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Construction of tailored C1-strains for desired enzymes

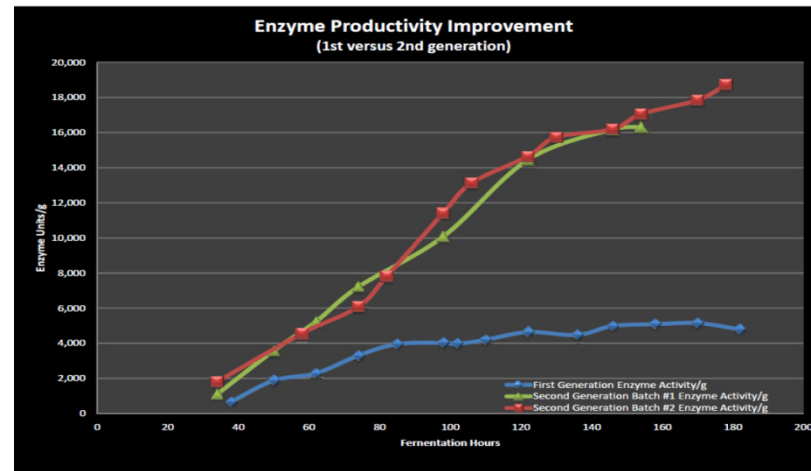


A library of single industrial enzymes produced at high levels is in place



SDS-Page analysis of end of fermentation broth of single enzyme producing LC-strains

- LC-strains routinely produce target single enzymes at the g/L scale.
- Up to 30 g/L of secreted protein has been obtained.
- 5- fold improvement within one year (commercial scale production).



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Construction of tailored C1-strains for desired enzymes



Features C1-enzyme library

- Broad spectrum of xylan, cellulose, pectin and other plant fiber degrading enzymes expressed and characterized.
- Enzymes show a broad temperature and pH profile.
- Enzymes have synergistic activity and can in combinations be used for the efficient degradation of plant-derived fibers.
- Applications in biofuels/biochemicals, paper&pulp, feed, food.

Several new products developed

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Construction of tailored C1-strains for desired enzymes



Functional **heterologous** enzymes/proteins produced at the g/L scale:

Source	Enzyme/protein
Fungal	Xylanases, amylase, cellulase, endo-polygalacturonase, oxidase, phytase
Bacterial	Xylosidase, Cellulase
Bacterial-directed evolution	Animal feed enzyme
Human	Immunoglobulin IgG1

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Construction of tailored C1-strains for desired enzymes



Dedicated enzyme mixtures for lignocellulosic fuels and chemicals

- Produced by a **single C1-strain** in a single fermentation
- **Fast liquefaction** of highly viscous lignocellulosic biomass
- **Fast saccharification**: Ethanol process completed in 72h.
- **Broad temperature range**: Active between 32 °C and 65 °C.
- **Broad pH-range**: High activity between pH 4.5 and 8.
- **Active on a variety of biomass substrates**: Corn Stover, Wheat Straw, Wheat bran, Sugar Cane Bagasse, Switch Grass, Sorghum, Wood and Paper Waste

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Summary C1-technology platform



- C1 is a proprietary **mature enzyme production system**.
- **High extracellular protein/enzyme** production levels obtained.
- **Easy scaling**, proven at the 150M³ fermentation scale.
- Proven for **heterologous enzyme and protein** production.
- Versatile genetic tools and hosts developed: **highly programmable**.
- Genome sequenced and annotated: **enzymatic potential elucidated**
- **Self affirmed GRAS** status of a C1-product was acknowledged by the FDA (2009)
- **Commercial products** developed, e.g. in the bioenergy and paper&pulp space.

