

Improved endoxylanase production and colony morphology of *Aspergillus niger* DSM 26641 by γ -ray induced mutagenesis

Christoph Ottenheim^{a,b}, Katharina A. Werner^b, Wolfgang Zimmermann^{a*}, Jin
Chuan Wu^{b**}

^a Microbiology and Bioprocess Engineering, Institute of Biochemistry, Leipzig University,
04103 Leipzig, Germany;

^b Institute of Chemical and Engineering Sciences, Agency for Sciences, Technology and
Research (A*STAR), 1 Pesek Road, Jurong Island, Singapore 627833;

Christoph Johann Heinrich Ottenheim, christoph@ottenheim.net

Katharina Alicia Werner, k.a.werner@t-online.de

***Corresponding author:** Wolfgang Zimmermann;

E-mail: wolfgang.zimmermann@uni-leipzig.de; Tel: +49-3419736781; Fax: +49-3419736798

****Corresponding author:** Jin Chuan Wu;

E-mail: wu_jinchuan@ices.a-star.edu.sg; Tel: +65-67963803; Fax: +65-63166182

Abstract: *Aspergillus niger* DSM 26641 was exposed to ^{60}Co γ -radiation to enhance the β -1, 4-endoxylanase activity, restrict colony growth and improve robustness of pellets. The first promising mutant obtained after γ -radiation of the fungal spores at 50-2000 Gy showed a restricted colony growth and an 82% enhancement in β -1, 4-endoxylanase activity. The mutant was subjected to a second round of γ -radiation at 1400 Gy generating a mutant with double the β -1, 4-endoxylanase activity compared to the native strain. The selected final mutant, deposited as *Aspergillus niger* DSM 28712, showed a maximal saccharification activity of 26 U ml $^{-1}$ on xylan based broth, 48 U ml $^{-1}$ on lignocellulose hydrolysate and 375 U ml $^{-1}$ on lignocellulose hydrolysate supplemented with yeast extract and mineral salts.

Keywords: Filamentous Fungi, γ -Radiation Mutagenesis, Enzymes, Endoxylanase, Microbial Growth, Optimisation.

1. Introduction

Filamentous fungi are a good source for xylanotic enzymes owing to their naturally inhabiting as lignocellulose decomposers. Individual fungal xylanases have been successfully overexpressed in various organisms such as bacteria [1], filamentous fungi [2], yeasts [3] and transgenic plants [4]. For complete degradation of hemicellulose, however, at least two backbone-degrading enzymes (β -1,4-endoxylanase and β -xylosidase) and three accessory enzymes (arabinofuranosidase, α -glucuronidase and acetyl xylan esterase) are needed [5]. Therefore, production of enzymes from filamentous fungi as enzyme cocktails is more suitable for industrial applications in breaking down hemicellulose to fermentable sugars. Native filamentous fungi are usually not ideal for producing industrial enzymes due to the low enzyme activities or un-favored morphology. In most cases, the activity of the backbone-degrading enzyme, β -1, 4-endoxylanase, needs to be enhanced for more efficient hydrolysis of xylan, the major component of hemicellulose. In addition, filamentous fungi cannot be easily optimized by using well-developed agar plate supported screening methods due to their widespread and fast-growing morphology, so colony restriction needs to be developed for easier screening. Although a limited colony restriction can be achieved by adding chemicals like oxgal [6, 7], a permanent change to a more restricted morphology would be more desirable to achieve easier recycling and

1
2
3
4 handling of fungal pellets in submerged cultures, which is the preferred modus for
5
6
7 enzyme production in industry rather than the natural growth modus of filamentous
8
9
10 fungi. For instance, it has been shown that a hard and smooth pellet surface
11
12
13 favored the production of enzymes and organic acids like citric acid [8].
14

15
16 The complete degradation of hemicellulose needs the synergistic action of a
17
18 group of enzymes, so a random approach followed by suitable screening methods
19
20
21 is most promising for improving the enzyme cocktails produced by fungi. The
22
23
24 enzyme activity levels are not only influenced by their coding genes but also by
25
26
27 control elements. Some important control elements include repression by carbon
28
29
30 sources [9], expression of transcription factors [10] and expression of kinases and
31
32
33 phosphatases for sensing different carbon sources or controlling the cellular energy
34
35
36 states [11]. The use of γ -radiation for introducing random mutations has the
37
38
39 advantages of deep penetration and fast induction of various mutations such as
40
41
42 single and double strand breaks, creation of base-free sites and modification of
43
44
45 bases [12]. γ -radiation from ^{60}Co radiation sources is a low linear energy transfer
46
47
48 radiation (LET radiation) with a LET value of $1 \text{ keV } \mu\text{m}^{-1}$ and below and tends to
49
50
51 create clustered non-double strand breaks which can be repair-resistant or at-least
52
53
54 more difficult to repair than non-clustered DNA damages as shown by using
55
56
57 synthetic oligonucleotides [13]. In addition, it enhances non-DNA related cell
58
59
60
61
62
63
64
65

1
2
3
4 stress by creation of free radicals [14]. γ -radiation has been successfully used for 2-
5
6
7 fold enhancement of cellulase and galactosidase production in fungi [15,16].
8
9

10 Here we report the introduction of random mutations in *Aspergillus niger*
11
12 DSM 26641 by γ -radiation to enhance its β -1, 4-endoxylanase activity and restrict
13
14
15 its colony and pellet morphology for favorable industrial applications. This is the
16
17
18 first report on improving an *Aspergillus niger* strain in both its β -1, 4-endoxylanase
19
20
21 activity and cellular morphology.
22
23

24 25 **2. Materials and Methods**

26 27 **2.1 β -1, 4-endoxylanase assays**

28
29
30
31 The β -1, 4-endoxylanase activity was assayed by using either Remazol Brilliant
32
33
34 Blue dyed xylan (RBB-Xylan) or pure xylan as substrates.
35
36
37

38 For the activity assay using RBB-Xylan as the substrate, RBB-Xylan was
39
40
41 prepared following the developed protocol [17]. Spores or mycelia from selected
42
43
44 mutants were used to inoculate 10 ml broth containing 2 g l⁻¹ xylan, 0.1 g l⁻¹ yeast
45
46
47 extract, 0.1 g l⁻¹ peptone and 0.1 g l⁻¹ xylose in 15 ml tubes. The tubes were
48
49
50 incubated at 30°C and 200 rpm for 4 days. After successful growth, the tubes were
51
52
53 centrifuged for 10 min at 3,220×g at 4°C to separate the mycelia from the
54
55
56 supernatant. The culture supernatant was used to determine the β -1, 4-
57
58
59
60
61
62
63
64
65

1
2
3
4 endoxylanase activity following the protocol as described previously [18]. The
5
6
7 absorbance of the native strain *Aspergillus niger* DSM 26641 was set as 100%.
8
9

10
11 For the activity assay using xylan as the substrate, the produced sugars in the
12
13 supernatant were detected by the 3, 5-dinitrosalicylic acid (DNS) method [19].
14
15

16 17 **2.2 γ -Ray irradiation procedures**

18
19
20

21 *Aspergillus niger* DSM 26641, which was isolated from a soil sample in Singapore
22
23 and exhibited high β -1, 4-endoxylanase activity, was subjected to γ -radiation for
24
25 improving its performances. After mycelial growth for 3 days at 30°C on Potato
26
27 Dextrose Agar (PDA, potato extract 4 g l⁻¹, dextrose 20 g l⁻¹, agar 20 g l⁻¹), black
28
29 spores were formed by the fungus. The PDA plates were submerged in 10 ml
30
31 sterile ultrapure water each and the spores were brought into suspensions.
32
33
34
35
36
37

38
39 A γ -ray irradiation chamber 4000a containing a ⁶⁰Co radiation source inside
40
41 a shielded compartment was used for irradiation of the spores at a dose rate of 88
42
43 Gy h⁻¹. For estimation of γ -radiation sensitivity, the spore solution was exposed to
44
45 radiation doses of 50-2000 Gy. Subsequently, the irradiated spore samples were
46
47 spread on PDA plates at different dilutions to reach a countable amount of colony
48
49 forming units (CFU). Afterwards, morphologically different colonies were selected
50
51 from suitable plates and transferred to fresh PDA plates for further observation.
52
53
54
55
56
57
58
59
60
61
62
63
64
65

The obtained CFU values were used to estimate its D_{10} value. The D_{10} value represents the radiation dose at which the viable inoculum is reduced by 90%.

After screening the morphological different colonies for β -1,4-endoxylanase activity, spore solutions from three selected mutants each (different colony morphology; higher β -1,4-endoxylanase activity) were directly irradiated with a total dose of 1400 Gy (as estimated by the sensitivity experiments) and streaked out on PDA plates. Not only morphologically different but also similar colonies were isolated and screened for higher β -1, 4-endoxylanase activity.

2.3 Validation of mutants and final strain selection

For final validation of β -1,4-endoxylanase activity produced by the selected mutants, spores of the respective mutants and the original wildtype strain were used to inoculate 40 ml xylan broth in 50 ml shaking flasks in duplicate. The shaking flasks were incubated at 30°C and 200 rpm. In regular intervals, β -1, 4-endoxylanase activity was measured using RBB-Xylan as described above.

For final strain selection, spores of the selected mutant were spread on 10 PDA plates in a fitting dilution to obtain single colonies. Before spores were observed, colonies exhibiting restricted colony growth were picked and transferred into a well plate half filled with PDA. After observation of spores with the desired morphology, the wells were filled up with sterile water. The obtained spore

solutions were pooled and mixed. The mixed spore solution was streaked out in a fitting dilution on 10 new PDA plates and the same procedure was repeated. For investigating the changes in β -1, 4-endoxylanase activity during final strain selection, the spore solutions were stored and assayed using RBB-Xylan. In addition, the initial xylan broth, the hydrothermal palm oil empty fruit bunch fiber (EFB) hydrolysate, a lignocellulose derivate, and hydrothermal hydrolysate of EFB supplied with 0.05% yeast extract, mineral salts and 3% EFB were used for production of enzymes and the total saccharification activity of the enzyme solutions were assayed using the DNS method. The final mutant was deposited as *Aspergillus niger* DSM 28712. All other mutants were preserved at -80°C in a 20% glycerol stock. Details and pictures were recorded and stored in a publically available database (www.domse.org).

3. Results and Discussion

3.1 Strain selection

Aspergillus niger is an attractive target for whole cell mutagenesis as it is generally recognized as safe and even used for producing food additives [20]. It is also one of the most important fungi used for industrial production of citric acid [21] and various enzymes. For hydrolytic enzymes, an improvement in production of β -1, 4-endoxylanase is not only favorable for more efficient hydrolysis of xylan but is also beneficial to cellulose decomposition by cellulases due to the removal of

1
2
3
4 xylan [22, 23]. The mutants created by γ -ray irradiation are not considered as
5
6
7 genetically modified organisms, so their release into the environment after usage
8
9
10 would be permitted, making the on-site enzyme production more cost-effective
11
12
13 [24].
14

15 16 17 **3.2 γ -Radiation sensitivity and mutant screening**

18
19 To achieve a high mutation probability and avoid extensive radiation
20
21 damages at the same time, two strategies were adapted. First, using spores instead
22
23
24 of mycelia for the γ -ray irradiation to achieve a higher level of stress resistance and
25
26
27 minimize the undesired side effects [25]. Second, a γ -radiation dosage of 1400 Gy
28
29
30 was applied, which equals 3-4 D_{10} , leading to a balance of damage and mutation
31
32
33 rates, similar to the established protocols [26]. The D_{10} value was estimated to be
34
35
36 400 Gy in the first round of γ -ray irradiation (Fig.1), similar to that estimated for
37
38
39 other strains of *Aspergillus niger* reported in literature [27, 28].
40
41

42
43 The first round of screening led to mutants that showed changes in spore
44
45
46 formation, spore color, spore quantity, mycelial growth rate, colonial growth rate
47
48
49 and formation of pigments (Fig.2). Totally, 40 mutants were isolated and tested for
50
51
52 their β -1, 4-endoxylanase activity, from mutant Pre_04 which completely lost its β -
53
54
55 1, 4-endoxylanase activity to the mutant Pre_M36 which increased its β -1, 4-
56
57
58 endoxylanase activity to 125% compared to the wild type strain. Totally 15
59
60
61 mutants showed an obviously higher β -1, 4-endoxylanase activity than the wild
62
63
64
65

1
2
3
4 type strain. In 7 out of 10 of the most active mutants, a loss or a very strong
5
6
7 reduction in spore formation was observed. The fungal sporulation system is
8
9 considered to be very sensitive, therefore easily susceptible to mutation damages
10
11 [29]. Sporulation is a very disturbing process for submersion production of
12
13 enzymes, but it is still useful for optimization, storage and easy multiplication and
14
15 thus should not be completely abolished. A delayed or weaker sporulation would
16
17 be more desirable. Therefore, in the second round of γ -ray irradiation, 3 mutants
18
19 that showed higher β -1, 4-endoxylanase activities beside morphological alterations
20
21 but still maintained the ability for spore formation were selected (Pre_M30: 82%
22
23 enhancement in β -1,4-endoxylanase activity, fewer spores and restricted colonial
24
25 growth; Pre_M15: 57% enhancement in β -1, 4-endoxylanase activity and slightly
26
27 slower mycelial growth; Pre_M40: 56% enhancement in β -1, 4-endoxylanase
28
29 activity and a change to a brownish spore pigmentation). The second round of γ -
30
31 ray irradiation of the spore solutions of the 3 selected mutants resulted in 101 new
32
33 mutants. All mutants showed a broad variation in β -1, 4-endoxylanase activity
34
35 similar to the mutants obtained in the first round.
36
37
38
39
40
41
42
43
44
45
46
47
48
49
50

51 **3.3 Validation of mutants and final strain selection**

52
53

54 For final validation, the best mutants from the second round of γ -ray
55
56 irradiation and their predecessors, the best two non-spore-forming mutants from
57
58
59
60
61
62
63
64
65

1
2
3
4 the first round of γ -ray irradiation and the wild type strain were tested for their β -1,
5
6
7 4-endoxylanase production for 89 h (Fig.3).
8
9

10 Pre_M15, Pre_40 and Pre_M30 showed roughly the same activity level at
11
12 110% (wild type 100%) when cultivated in 50 ml flasks. Pre_26 and Pre_36
13
14 showed a clearly higher activity (130%) than the wild type strain, but they were not
15
16 selected for further study due to the undesired loss in spore formation. Almost all
17
18 mutants from the second screening (M26, M20, M12, M09) round gave lower
19
20 activities, except the descendants of Pre_M30 (M42, M43) which showed an
21
22 activity level at 180%, almost double that of the wild type strain.
23
24
25
26
27
28
29
30

31 The obvious differences between observed activities in different screening
32
33 rounds and the final validation experiment might reflect the susceptibility of
34
35 filamentous fungi to the changes of environmental conditions such as the change of
36
37 culture volume and shape of the incubators [30].
38
39
40
41
42
43

44 Mutant M42 was selected for the final strain selection as it showed almost
45
46 double β -1, 4-endoxylanase activity of the wild type strain with restricted colony
47
48 growth. The occurrence of the widespread colony morphology could be found
49
50 initially in up to 8% of the colonies after streaking a suitably diluted solution on
51
52 PDA plates. After selecting colonies with restricted growth followed by pooling
53
54 and re-plating, the occurrence of widespread colony growth dropped from 8%
55
56
57
58
59
60
61
62
63
64
65

(initial mutant) to 3.8% (first round), 2.7% (second round), 0.6% (third round) and below 0.04% (final round) of all counted colonies. Not only the desired colony morphology was observed but the β -1, 4-endoxylanase activity enhanced significantly in each round (Fig.4). The repeated selection and pooling of spores from restricted colonies stabilized the obtained mutant strains. Almost no occurrence of the wild type growth modus was observed. Additionally, after each selection round, an obvious increase in β -1, 4-endoxylanase activity was observed, suggesting a possible relation between the observed restricted colony growth and the enhanced β -1, 4-endoxylanase activity. The relation between morphology and enzyme activity was reported earlier but is still poorly understood [31]. Therefore, one of the possible explanations lies within the ability of the mutant to form smaller pellets, lowering the amount of undersupplied or even dead mycelia, increasing the pellet density and subsequently the enzyme production. However, possibilities like the changes of enzyme activities or metabolic pathways are other possible explanations.

When assayed for the saccharification activity using the DNS method, the wild type and mutant strain exhibited similar activities (27 U ml^{-1} and 26 U ml^{-1}) on xylan. This stands in opposite to the different β -1, 4-endoxylanase activities which were observed earlier using the RBB-Xylan assay. The main reason might be the fundamental differences of the two assays. The RBB-Xylan assay measures

solely the liberated dyed xylan oligomers and thus only β -1, 4-endoxylanase activity, while the DNS assay measures the liberation of reducing carbohydrate units directly. Therefore other enzyme activities such as β -xylosidase activity might be higher in the supernatant of the wild type strain leading to similar overall reducing carbohydrate units. The wild type strain showed a higher activity than the mutant (59 U ml⁻¹ vs 48 U ml⁻¹) on hydrothermal hydrolysate of EFB. For the EFB hydrolysate supplied with additional carbon source, nitrogen source and salts, the mutant produced a more active supernatant than the wild type strain (375 U ml⁻¹ vs 337 U ml⁻¹). The saccharification activity of 375 U ml⁻¹ of the mutant is higher than the reported 210 U ml⁻¹ of a wild type strain [32] but lower than that of a genome shuffled strain (427.5 U ml⁻¹) [33].

3.4 Morphological comparison between wild type and mutant strains

In comparison with the wild type strain *Aspergillus niger* DSM 26641, the finally stabilized mutant M42, deposited as *Aspergillus niger* DSM 28712, showed all desired morphological attributes. When growing on PDA plates (Fig.5a-b), colonies of the mutant were well separated from each other at higher colony density. The spore formation of the mutant was delayed for 1 day and a yellow pigment was formed in excess after 4 days. Under the microscope (Fig.5c-d), the wild type strain revealed a more widespread colony growth than the mutant. The mutant grew on PDA in a dense and more uplifted way with no obvious growth

1
2
3
4 after 4 days. When growing in submerged cultures (Fig.5e-f), the mutant formed
5
6
7 small, restricted, robust and regular pellets, while the wild type strain exhibited the
8
9
10 opposite growth pattern. The pellets of the mutant were easier to handle for
11
12
13 recycling purposes.
14

15 16 **3.5 Conclusions**

17
18
19 *Aspergillus niger* DSM 26641, a native β -1, 4-endoxylanase-producing
20
21
22 filamentous fungus, was successfully improved by several rounds of γ -radiation
23
24
25 and subsequent strain stabilization. The finally selected mutant, *Aspergillus niger*
26
27
28 DSM 28712, showed double β -1,4-endoxylanase activity compared to the wild
29
30
31 type strain and a restricted widespread filamentous growth on agar plates and in
32
33
34 submerged cultures. This makes it easier to be picked up from agar plates for
35
36
37 screening for further improvement by γ -radiation and other mutagenesis methods
38
39
40 and more suitable for submerged fermentation for industrial applications.
41
42
43
44
45
46
47
48
49
50
51
52
53
54
55
56
57
58
59
60
61
62
63
64
65

Acknowledgements

This work was supported by the Science and Engineering Research Council (SERC) of the Agency for Science, Technology and Research (A*STAR) under the Value-Added Chemicals from Lignocellulose (VACL) Program (SERC grant no. 0921590133) and Biomass to Chemicals Program (SERC grant no 1124004027) through the A*STAR Research Attachment Program (ARAP) Scholarship. We thank Dr. Meredith E. K. Calvert for providing high resolution colony pictures using the TLL bioimaging & biocomputing facility.

References

- [1] C. Teng, H. Jia, Q. Yan, P. Zhou, Z. Jiang, High-level expression of extracellular secretion of a β -xylosidase gene from *Paecilomyces thermophila* in *Escherichia coli*, *Bioresource Technol.* 102 (2011) 1822-1830.
- [2] G. E. Anasontzis, S. Zerva, P. M. Stathopoulou, K. Haralampidis, G. Diallinas, A. D. Karagouni, D. G. Hatzinikolaou, Homologous overexpression of xylanase in *Fusarium oxysporum* increases ethanol productivity during consolidated bioprocessing (CBP) of lignocellulosics, *J. Biotechnol.* 152 (2011) 16-23.
- [3] B. Tian, Y. Xu, W. Cai, Q. Huang, Y. Gao, X. Li, J. Huang, Molecular Cloning and Overexpression of an Endo- β -1,4-xylanase Gene from

- 1
2
3
4 *Aspergillus niger* in Industrial *Saccharomyces cerevisiae* YS2 Strain, Appl.
5
6
7 Biochem. Biotech 170 (2013) 320-328.
8
9
- 10 [4] H.-J. Bae, H.-J. Kim, Y. S. Kim, Production of a recombinant xylanase in
11
12 plants and its potential for pulp biobleaching applications, Bioresource
13
14 Technol. 99 (2008) 3513-3519.
15
16
17
18 [5] V. Juturu, J.C. Wu, Microbial xylanases: Engineering, production and
19
20 industrial applications, Biotechnol. Adv. 30 (2012) 1219-1227.
21
22
23
24 [6] M. I. Rajoka, A. Yasmeen, Improved productivity of beta-fructofuranosidase
25
26 by a derepressed mutant of *Aspergillus niger* from conventional and non-
27
28 conventional substrates, World J. Microb. Biot. 21 (2005) 471-478.
29
30
31
32
33 [7] P.H. Tsao, Selection media for isolation of pathogenic fungi, Annu. Rev.
34
35 Phytopathol. 8 (1970) 157-186.
36
37
38
39 [8] M. Papagianni, M. Mattey, Morphological development of *Aspergillus niger*
40
41 in submerged citric acid fermentation as a function of the spore inoculum
42
43 level, Application of neural network and cluster analysis for characterization
44
45 of mycelial morphology, Microb. Cell Fact. 5:3 (2006).
46
47
48
49
50 [9] G. J. G. Ruijter, J. Visser, Carbon repression in aspergilla, FEMS Microbiol.
51
52 Lett. 151 (2006) 1574-6968.
53
54
55
56 [10] R. Rauscher, E. Würleitner, C. Wacenovsky, N. Aro, A. R. Stricker,
57
58 S. Zeilinger, C. P. Kubicek, M. Penttilä, R. L. Mach, Transcriptional
59
60
61
62
63
64
65

1
2
3
4 Regulation of xyn1, Encoding Xylanase I, in *Hypocrea jecorina*, Eukaryot.
5
6 Cell. 5 (2006) 447-56.
7
8

9
10 [11] N. A. Brown, P. F. de Gouvea, N. G. Krohn, M. Savoldi, G. H. Goldman,
11
12 Functional characterisation of the non-essential protein kinases and
13
14 phosphatases regulating *Aspergillus nidulans* hydrolytic enzyme production,
15
16 Biotechnol. Biofuels. 6 (2013) article 91.
17
18

19
20 [12] W. D. Henner, S. M. Grunberg, W. A. Haseltine, Sites and structure of gamma
21
22 radiation-induced DNA strand breaks, J. Biol. Chem. 257 (1982) 11750-
23
24 11754.
25
26
27

28
29 [13] M. Hada, A.G. Georgakilas, Formation of Clustered DNA Damage after
30
31 High-LET Irradiation: A review, J. Radiat. Res. 49 (2008) 203-210.
32
33

34
35 [14] N. P. McNamara, H. I. J. Black, N. A. Beresford, N. R. Parekh, Effects of
36
37 acute gamma irradiation on chemical, physical and biological properties of
38
39 soils, Appl. Soil Ecol. 24 (2003) 117–132.
40
41
42

43
44 [15] V. H. Vu, T. A. Pham, K. Kim, Improvement of fungal cellulase production
45
46 by mutation and optimization of solid state fermentation, Mycobiology 39
47
48 (2011) 20-25.
49
50

51
52 [16] M. S. Awan, N. Tabbasam, N. Ayub, M. E. Babar, Mehboob-ur-Rahman, S.
53
54 M. Rana, M. I. Rajoka, Gamma radiation induced mutagenesis in *Aspergillus*
55
56
57

- 1
2
3
4 *niger* to enhance its microbial fermentation activity for industrial enzyme
5
6
7 production, Mol. Biol. Rep. 38 (2011) 1367-1374.
8
9
- [17] P. Biely, D. Mislovicova, R. Toman, Remazol brilliant blue-xylan: A soluble
10
11 chromogenic substrate for xylanases, Meth. Enzymol. 160 (1988) 536-541.
12
13
14
- [18] C. Ottenheim, C. Verdejo, W. Zimmermann, J. C. Wu, Hemicellulase
15
16 production by *Aspergillus niger* DSM 26641 in hydrothermal palm oil empty
17
18 fruit bunch hydrolysate and transcriptome analysis, J. Biosci. Bioeng. (2014),
19
20
21 <http://dx.doi.org/10.1016/j.jbiosc.2014.05.014> (in press).
22
23
24
25
26
- [19] G. L. Miller, Use of dinitrosalicylic acid reagent for determination of reducing
27
28 sugar, Anal. Chem. 31 (1959) 426–428.
29
30
31
- [20] S. K. Shankar, V. H. Mulimani, α -Galactosidase production by *Aspergillus*
32
33 *oryzae* in solid-state fermentation, Bioresource Technol. 98 (2007) 958-961.
34
35
36
37
- [21] M. Papagianni, Advances in citric acid fermentation by *Aspergillus niger*:
38
39 Biochemical aspects, membrane transport and modelling, Biotechnol. Adv. 25
40
41
42 (2007) 244-263.
43
44
45
46
- [22] M. A. Kabel, H. van den Borne, J.-P. Vincken, A. G. J. Voragen, H. A.
47
48 Schols, Structural differences of xylans affect their interaction with cellulose,
49
50 Carbohydr. Polym. 69 (2007) 94-105.
51
52
53
54
55
56
57
58
59
60
61
62
63
64
65

- 1
2
3
4 [23] Q. Qing, C. E. Wyman, Supplementation with xylanase and β -xylosidase to
5
6
7 reduce xylo-oligomer and xylan inhibition of enzymatic hydrolysis of
8
9
10 cellulose and pretreated corn stover, *Biotechnol. Biofuels* 4:18 (2011).
11
12
13 [24] L. Viikari, J. Vehmaanpera, A. Koivula, Lignocellulosic ethanol: From
14
15
16 science to industry, *Biomass Bioenergy* 46 (2012) 13-24.
17
18
19 [25] T. T. Wyatt, H. A. Wösten, J. Dijksterhuis, Fungal spores for dispersion in
20
21
22 space and time, *Adv. Appl. Microbiol.* 85 (2013) 43-91.
23
24
25 [26] Y. Toyoshima, A. Takahashi, H. Tanaka, J. Watanabe, Y. Mogi, T. Yamazaki,
26
27
28 R. Hamada, K. Iwashita, K. Satoh, I. Narumi, Lethal and mutagenic effects of
29
30
31 ion beams and γ -rays in *Aspergillus oryzae*, *Mutat. Res.-Fund. Mol. M.* 740
32
33
34 (2012) 43-49.
35
36
37 [27] Y. G. Saleh, M. S. Mayo, D. G. Ahearn, Resistance of some common fungi to
38
39
40 gamma irradiation, *Appl. Environ. Microbiol.* 54 (1988) 2134-2135.
41
42
43 [28] G. Blank, D. Corrigan, Comparison of resistance of fungal spores to gamma
44
45
46 and electron beam radiation, *Int. J. Food Microbiol.* 26 (1995) 269-77.
47
48
49 [29] H.-S. Park, J.-H. Yu, Genetic control of asexual sporulation in filamentous
50
51
52 fungi, *Curr. Opin. Microbiol.* 15 (2012) 669-677.
53
54
55 [30] A. M. Gibson, A. D. Hocking, Advances in the predictive modelling of fungal
56
57
58 growth in food, *Trends Food Sci. Tech.* 8 (1997) 353-358.
59
60
61
62
63
64
65

- 1
2
3
4 [31] S. de Nicolás-Santiago, C. Regalado-González, B. García-Almendárez, F. J.
5
6
7 Fernández, A. Téllez-Jurado, S. Huerta-Ochoa, Physiological, morphological,
8
9 and mannanase production studies on *Aspergillus niger* uam-gs1 mutants,
10
11 Electron. J. Biotechn. 9 (2006) 50-60.
12
13
14
15 [32] A. R. Shah, D. Madamwar, Xylanase production by a newly isolated
16
17 *Aspergillus foetidus* strain and its characterization, Process Biochem. 40
18
19 (2005) 1763 – 1771.
20
21
22
23 [33] A. M. A. El-Bondkly, Molecular Identification Using ITS Sequences and
24
25 Genome Shuffling to Improve 2-Deoxyglucose Tolerance and Xylanase
26
27 Activity of Marine-Derived Fungus, *Aspergillus sp.* NRCF5, Appl. Biochem.
28
29 Biotech. 167 (2012) 2160-2173.
30
31
32
33
34
35
36
37
38
39
40
41
42
43
44
45
46
47
48
49
50
51
52
53
54
55
56
57
58
59
60
61
62
63
64
65

Figure legends

Fig. 1. Relation between γ -radiation dose and survival rate

Fig. 2. Comparison of selected colony morphologies after the first round of γ -radiation. The wild type strain *Aspergillus niger* DSM 26641 is marked red

Fig.3. Time courses of β -1, 4-endoxylanase activity of selected mutants. The activity of the wildtype strain at 65 h was set as 100% (Error bars = standard deviation values)

Fig.4. Change of β -1, 4-endoxylanase activity of mutant Pre_M30 during 3 generations of final strain selection of mutant M42. The initial activity was set as 100% (Error bars = standard deviation values)

Fig.5. Comparison of colony morphologies between the wild type *Aspergillus niger* DSM 26641 and the final mutant *Aspergillus niger* DSM 28712. a and b: colonies on agar plates; c and d: colonies on agar plates in magnification; e and f: pellets in submerged cultures

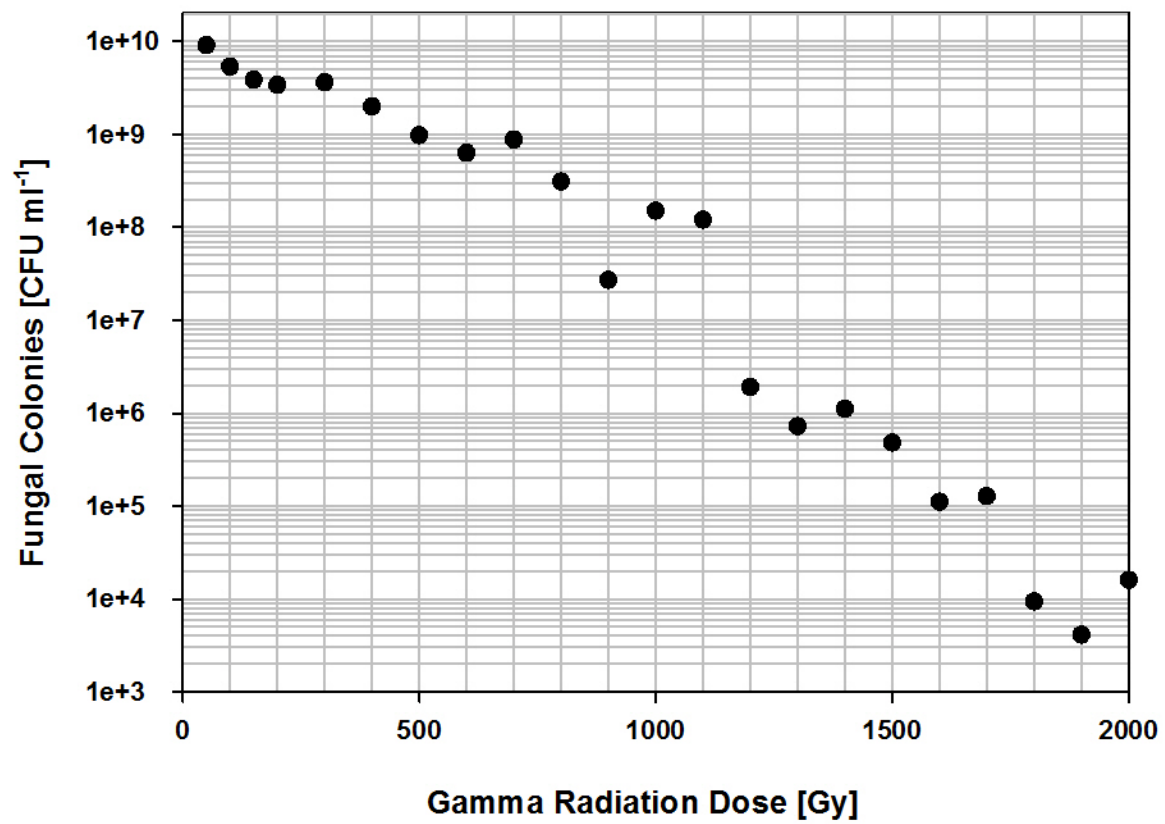


Fig.1



Fig.2

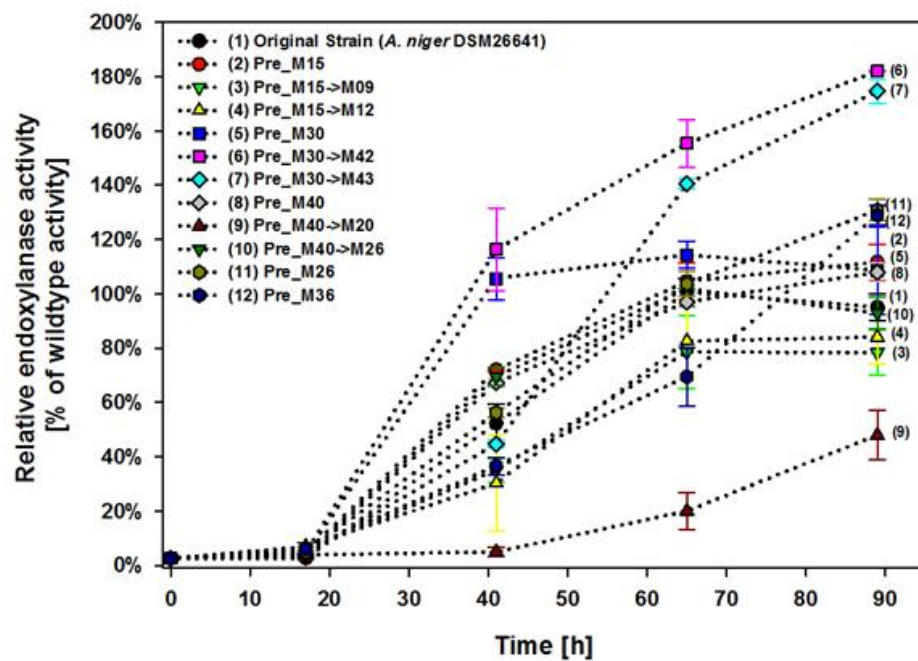


Fig.3

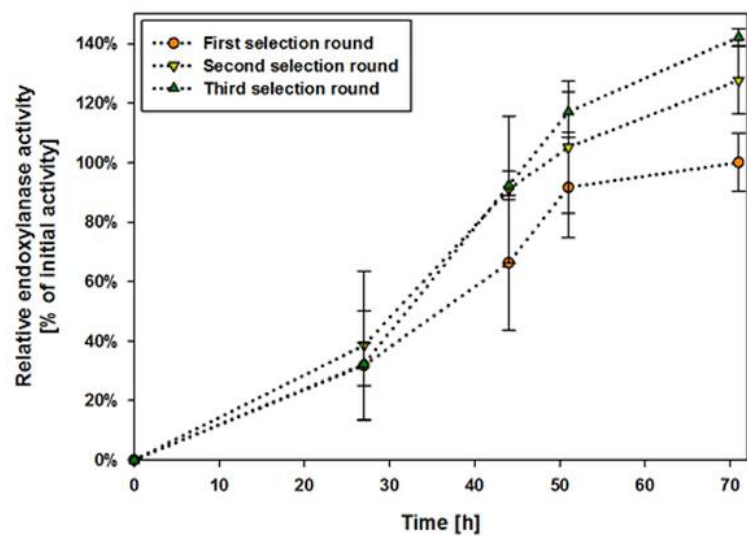


Fig.4

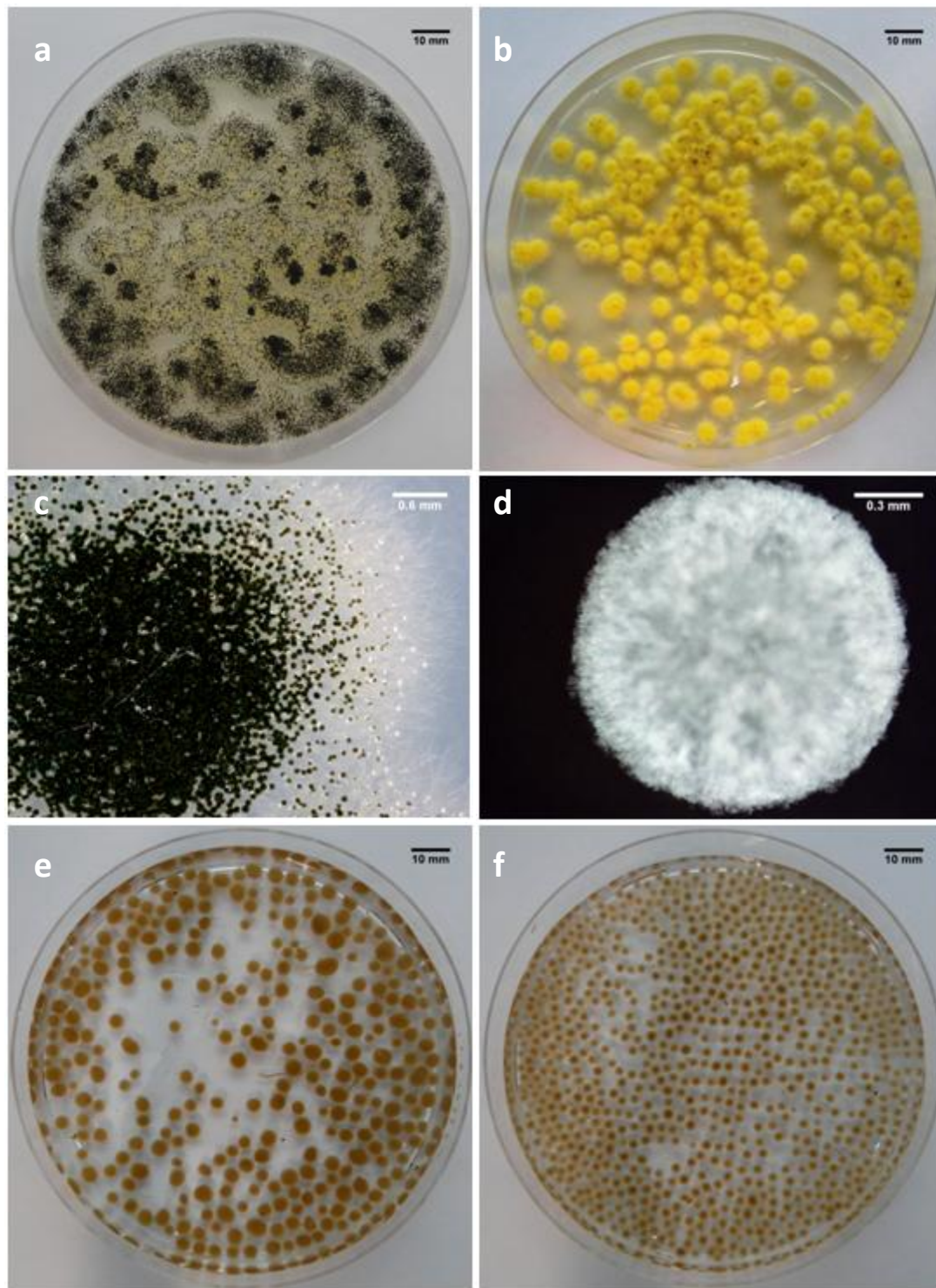


Fig.5