

## Detection of Extracellular Enzyme Activity in *Penicillium* using Chromogenic Media

Ji Hwan Yoon<sup>1</sup>, Seung Beom Hong<sup>2</sup>, Seung Ju Ko<sup>2</sup> and Seong Hwan Kim<sup>1\*</sup>

<sup>1</sup>Department of Microbiology and Institute of Basic Sciences, Dankook University, Cheonan, Chungnam 330-714, Korea

<sup>2</sup>Korean Agricultural Culture Collection and National Institute of Agricultural Science and Technology, Rural Development Administration, Suwon, Kyungido 441-707, Korea

(Received September 23, 2007)

A total of 106 *Penicillium* species were tested to examine their ability of degrading cellobiose, pectin and xylan. The activity of  $\beta$ -glucosidase was generally strong in all the *Penicillium* species tested. *P. citrinum*, *P. charlesii*, *P. manginii* and *P. aurantiacum* showed the higher ability of producing  $\beta$ -glucosidase than other tested species. Pectinase activity was detected in 24 *Penicillium* species. *P. paracanescens*, *P. sizovae*, *P. sartoryi*, *P. chrysogenum*, and *P. claviforme* showed strong pectinase activity. In xylanase assay, 84 *Penicillium* species showed activity. Strong xylanase activity was detected from *P. megasporum*, *P. sartoryi*, *P. chrysogenum*, *P. glandicola*, *P. discolor*, and *P. coprophilum*. Overall, most of the *Penicillium* species tested showed strong  $\beta$ -glucosidase activity. The degree of pectinase and xylanase activity varied depending on *Penicillium* species.

**KEYWORDS:** Chromogenic media,  $\beta$ -Glucosidase, Pectinase, *Penicillium*, Xylanase

*Penicillium* is well-known fungus that served as original source of the first antibiotic Penicillin. The species belonging to *Penicillium* are generally called as green moulds due to their color of spores and mycelium. This fungus recognized generally by its dense brush-like spore-bearing structures is a taxonomically large and difficult genus encountered almost everywhere, and usually the most abundant genus of fungi in soils and one of major groups of indoor moulds (Domsch, 1980; Pitt *et al.*, 1979). The common occurrence of *Penicillium* species in food is a particular problem because some species produce toxins and may render food inedible or even dangerous (de Hoog *et al.*, 2000; Pitt *et al.*, 2000). On the other hand, some species of *Penicillium* are beneficial to humans. Cheeses are ripened with species of *Penicillium* and are quite safe to eat.

*Penicillium* also has been known as producers of useful enzymes that degrade proteins, xylan, starch, lipids etc. from agricultural wastes (Techapun *et al.*, 2003). However, in spite of its prevalence in nature, not many species of *Penicillium* have been known for their ability of producing extracellular enzymes. Recently, Krogh *et al.* (2004) tried to screen some *Penicillium* fungi for cellulase and xylanase production. To better use of this fungus in industry such as food, textile, and bioremediation, further search for the source of enzymes needs to be proceeded. In this context this study was aimed at evaluation of vast numbers of *Penicillium* species using chromogenic medium (Castro *et al.*, 1995) that allowed us to examine

the degree of  $\beta$ -glucosidase, pectinase and xylanase the activity.

Totally, 106 *Penicillium* species were provided by Korean Agricultural Culture Collection (KACC, Suwon, Korea) and used for the present study. Fungal cultures were maintained on potato dextrose agar (Difco, USA). For the test of extracellular enzyme activities, the cultures were grown on PDA and transferred to chromogenic media containing each carbon source as enzymatic substrates (Ten *et al.*, 2004). Chromogenic media contained 0.1% yeast nitrogen base (Difco, USA), and 0.5% D-cellobiose (Sigma, USA), polygalacturonic acid (MP Bio-medical, France), xylan from oat spelts (Sigma, USA) as enzymatic carbon source, 0.5% Congo Red dye (Sigma, USA) for chromogenic reaction (Teather and Wood, 1982), and 1.5% agar powder (Yoon *et al.*, 2007). After 5 days of culturing at 25°C, evaluation of enzyme activity was conducted by measuring clear zone (plaque) peripherally formed around the fungal colony resulting from reaction between chromogenic substrates and the enzyme produced by the fungus. The degree of enzyme activity was expressed as strong (S), moderate (M), and weak or no activity (N) by measuring the size of clear zone. When the clear zone size was over 0.5 cm, the activity was treated as strong. Moderate activity was recorded when clear zone size was 0.1~0.4 cm. When clear zone size was less than 0.1 cm, the activity was treated as no activity. There was little difference between the sizes of clear zone measured by a ruler and shown on photo due to the resolution of photo image. The values measured by a ruler were used for comparative determination of clear zone

\*Corresponding author <E-mail: piceae@naver.com>

**Table 1.** Detection of extracellular enzyme activities in *Penicillium* spp. on chromogenic medium

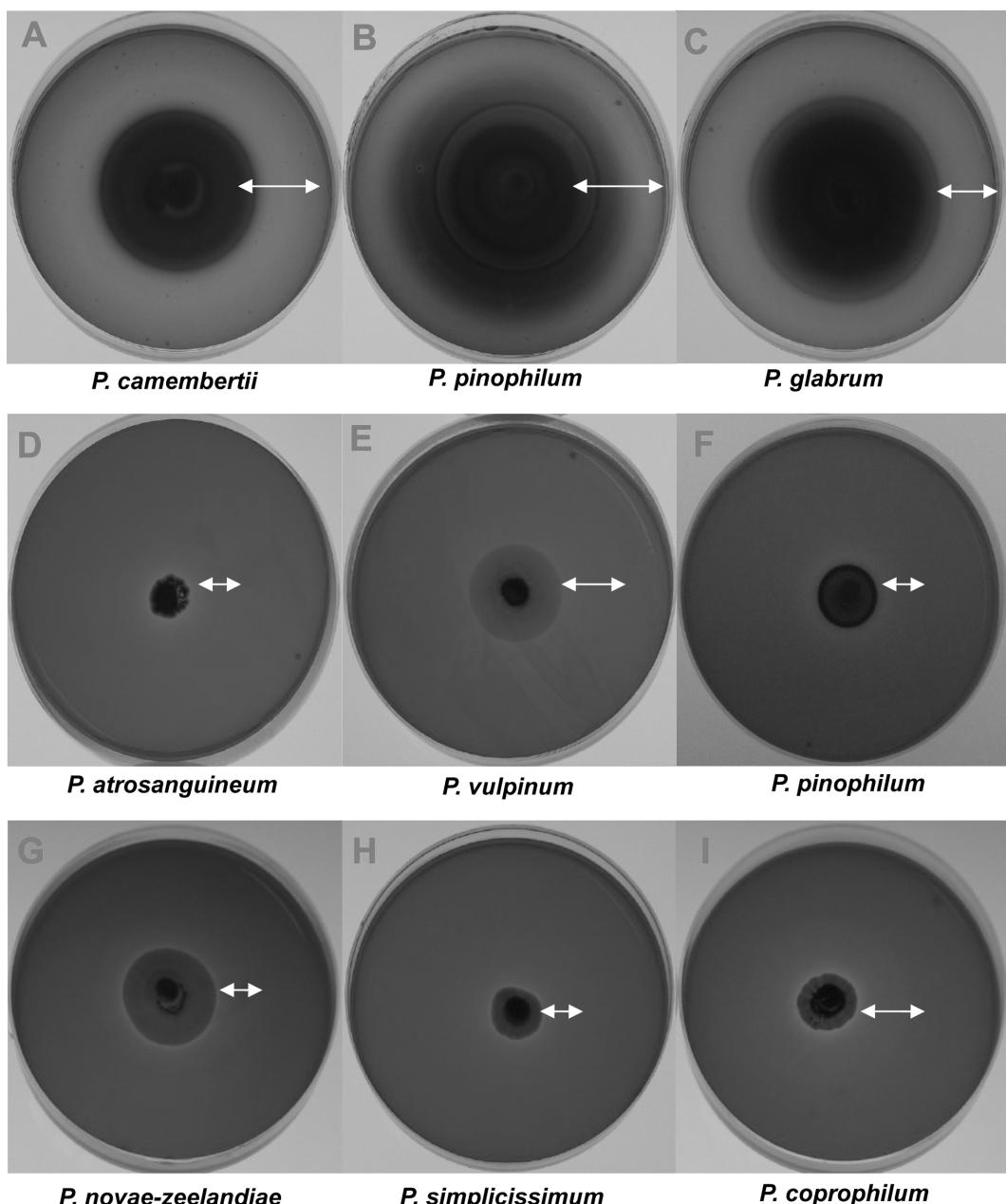
Species/KACC no.	$\beta$ -glucosidase	Pectinase	Xylanase	Species/KACC no.	$\beta$ -Glucosidase	Pectinase	Xylanase
<i>P. aculeatum</i> /40586	S	N	M	<i>P. lanosum</i> /41914	S	N	N
<i>P. adametzii</i> /40587	S	N	N	<i>P. lividum</i> /40609	S	N	N
<i>P. aethiopicum</i> /41335	S	N	M	<i>P. luteum</i> /40610	S	N	S
<i>P. album</i> /40588	S	N	N	<i>P. madritii</i> /41481	S	N	S
<i>P. allahabadense</i> /40632	S	S	S	<i>P. manginii</i> /41482	S	N	M
<i>P. allii</i> /41337	S	M	M	<i>P. megasporum</i> /40637	S	N	S
<i>P. amagasakiense</i> /40633	S	S	S	<i>P. melanoconidium</i> /41350	S	N	N
<i>P. atramentosum</i> /40589	S	S	S	<i>P. melinii</i> /41497	S	N	N
<i>P. atrosanguineum</i> /40590	S	S	S	<i>P. miczynskii</i> /40612	S	N	M
<i>P. aurantiacum</i> /40634	S	N	N	<i>P. moldavicum</i> /41505	S	M	M
<i>P. aurantiogriseum</i> /40591	S	N	N	<i>P. mononematosum</i> /41916	S	N	M
<i>P. avellaneum</i> /40598	S	N	N	<i>P. nalgiovense</i> /41351	S	M	N
<i>P. baradicum</i> /41580	S	N	M	<i>P. neoechinulatum</i> /41917	S	N	M
<i>P. brasiliandum</i> /41527	S	N	M	<i>P. novae-zeelandiae</i> /41506	S	M	M
<i>P. brevicompactum</i> /41339	S	N	M	<i>P. ochraceum</i> /40613	S	N	M
<i>P. camemberti</i> /41890	S	N	S	<i>P. ochrochloron</i> /41510	S	N	M
<i>P. canescens</i> /40662	S	N	S	<i>P. oxalicum</i> /41013	S	N	N
<i>P. caseicolum</i> /40635	S	N	N	<i>P. palitans</i> /41919	S	N	M
<i>P. charlesii</i> /41340	S	N	M	<i>P. pallidum</i> /40638	S	N	M
<i>P. chermesinum</i> /41449	S	N	N	<i>P. paneum</i> /41920	S	N	M
<i>P. chrysogenum</i> /41891	S	N	M	<i>P. paracanescens</i> /41933	S	S	N
<i>P. citrinum</i> /40601	S	N	M	<i>P. paxilli</i> /41534	S	S	M
<i>P. claviforme</i> /40381	S	N	M	<i>P. pinophilum</i> /40615	S	N	S
<i>P. commune</i> /41341	S	N	N	<i>P. polonicum</i> /41353	M	N	N
<i>P. concentricum</i> /41895	S	N	M	<i>P. purpurogenum</i> /40161	S	N	S
<i>P. coprophilum</i> /41896	S	S	S	<i>P. raciborskii</i> /41536	M	N	M
<i>P. coralligerum</i> /41427	S	N	M	<i>P. raistrickii</i> /41539	S	N	S
<i>P. corylophilum</i> /40688	S	N	M	<i>P. resticulosum</i> /41921	S	S	S
<i>P. cremeogriseum</i> /41439	S	N	M	<i>P. rolfssii</i> /41545	S	N	M
<i>P. cyclopium</i> /41006	S	N	M	<i>P. roqueforti</i> /40617	S	S	S
<i>P. daleae</i> /41408	S	N	M	<i>P. rugulosum</i> /41355	S	N	S
<i>P. decumbens</i> /40604	S	N	S	<i>P. sartoryi</i> /41416	M	N	M
<i>P. digitatum</i> /40822	S	N	S	<i>P. sclerotigenum</i> /41546	S	S	M
<i>P. dipodomysis</i> /41899	S	N	M	<i>P. sclerotiorum</i> /40618	S	S	M
<i>P. discolor</i> /41344	S	N	S	<i>P. simplicissimum</i> /40619	S	N	M
<i>P. echinulantum</i> /40605	S	N	M	<i>P. sizovae</i> /41565	S	N	M
<i>P. expansum</i> /40606	S	N	M	<i>P. skrjabinii</i> /41567	S	N	M
<i>P. fennelliae</i> /41903	S	N	M	<i>P. solitum</i> /40818	S	S	S
<i>P. flavigenum</i> /41904	S	N	N	<i>P. soppii</i> /41568	S	N	M
<i>P. freii</i> /41345	S	N	M	<i>P. spinulosum</i> /40620	S	S	S
<i>P. funiculosum</i> /41346	S	N	S	<i>P. steckii</i> /41434	S	N	S
<i>P. glabrum</i> /40607	S	N	M	<i>P. tricolor</i> /41356	S	M	M
<i>P. glandicola</i> /41643	S	N	M	<i>P. turbatum</i> /40623	S	N	M
<i>P. griseofulvum</i> /41347	S	N	M	<i>P. ulaiense</i> /41357	S	N	M
<i>P. griseopurpureum</i> /40248	S	N	S	<i>P. variabile</i> /41358	S	M	M
<i>P. herquei</i> /41455	S	N	S	<i>P. velutinum</i> /40624	S	N	S
<i>P. hirsutum</i> /41348	S	N	S	<i>P. verrucosum</i> /41927	S	N	M
<i>P. hordei</i> /41349	S	N	N	<i>P. verruculosum</i> /41625	S	N	M
<i>P. inflatum</i> /41461	S	N	N	<i>P. vinaceum</i> /40626	S	S	S
<i>P. italicum</i> /40826	S	S	S	<i>P. viridicatum</i> /41359	S	S	S
<i>P. janczewskii</i> /41410	S	N	S	<i>P. vulpinum</i> /40627	S	S	S
<i>P. jensenii</i> /40636	S	N	N	<i>P. waksmanii</i> /40628	S	M	M
<i>P. kabunicum</i> /41479	S	S	S	<i>P. westlingii</i> /41597	S	M	N

S = strong enzyme activity, M = moderate enzyme activity, and N = weak or no enzyme activity.

size. All the measurement values were obtained from 5 replicate plates of each species.

Table 1 shows the results of the detection of three extracellular enzymes in 106 species of *Penicillium*. D-cellulose was degraded more easily than other substrates because all the species in Table 1 showed activity. One hundred and three of 106 species showed strong  $\beta$ -glucosidase activity. Thus, the activity of  $\beta$ -glucosidase is generally strong in most of the *Penicillium* species tested (Table 1, Fig. 1). Examples of  $\beta$ -glucosidase activity formed on chromogenic media are shown in Figs. 1A-C. The breaking activity of pectin was observed from 24 of

106 species. Among these 24 pectinase positive species, 18 species showed strong activity. Especially, *P. coprophilum*, *P. sartoryi*, *P. sizovae*, *P. pinophilum* and *P. paracrenulatum* showed stronger pectinase activity than others. This indicates not many *Penicillium* species can degrade pectin compounds strongly. Examples of pectinase activity on chromogenic media are given in Figs. 1D-F. Compared to the number of *Penicillium* species that could degrade pectin, more number of *Penicillium* species showed the ability of xylan degradation. Twenty three species showed strong xylanase activity and fifty one species showed moderate activity. In xylanase assay, 84 *Peni-*



**Fig. 1.** Examples of observation of different enzyme activities produced by diverse *Penicillium* species on chromogenic media. A, B, C :  $\beta$ -glucosidase. D, E, F : pectinase. G, H, I : xylanase. Arrows indicate clear zone.

*cillium* species showed the activity. In these xylanase positive species, 33 species showed strong activity and 51 species displayed moderate activity. Strongest xylanase activity was detected from *P. coprophilum*, *P. simplicissimum*, *P. sartoryi* and, *P. discolor*. Figs. 1G-H show the examples of xylanase activity checked on chromogenic media. It seems that *Penicillium* is good sources of finding noble fungal xylanase.

Among the 106 species tested, *P. allahabadense*, *P. atrosanguineum*, *P. coprophilum*, *P. italicum*, *P. kabunicum*, *P. resticulosum*, *P. roqueforti*, *P. solitum*, *P. spinulosum*, *P. viridicatum*, *P. vulpinum*, and *P. waksmanii* showed strong activities of all the three extracellular enzymes tested (Table 1). Since there has been not much information in literature on the activity of  $\beta$ -glucosidase, pectinase, and xylanase in these 12 species, the results of Table 1 are quite useful for the understanding of their biochemical properties and nutrient physiology.

In conclusion, our work generated basic information on the vast numbers of *Penicillium* species. The genus contained many species that can be used for production of extracellular  $\beta$ -glucosidase, pectinase, and xylanase. Since all the species tested in this study is available from KACC, our results will be useful to the researchers who are searching for the tested enzyme sources. The genome sequence project on the human pathogen *Penicillium marneffei* (de Hoog *et al.*, 2000) has been performed and it is currently in assembly stage. Thus, its genome information will be available soon at genome project database in GenBank database (<http://www.ncbi.nlm.nih.gov/>). The genomic information of *P. marneffei* surely provides useful background information and opportunity for the search of genes encoding extracellular enzymes from *Penicillium* spp. used in this study.

## Acknowledgment

This study was supported by a grant (project no : 20060401034815) of BioGreen21 program from the Rural Development of Administration in Korea.

## References

- Castro, G. R., Baigori, M. D. and Sineriz, F. 1995. A plate technique for screening of inulin degrading microorganisms. *J. Microbiol. Methods* **22**: 51-56.
- de Hoog, G. S., Guarro, J., Gene, J. and Figueras, M. J. 2000. *Atlas of Clinical Fungi*, 2nd ed, vol. 1. Centraalbureau voor Schimmelcultures, Utrecht, The Netherlands.
- Domsch, K. H., Gams, W. and Anderson, T.-H. 1980. *Compendium of soil fungi*. Academic Press, London 2 vol.
- Krogh, K. B. R., Morkeberg, A., Friscad, J. C. and Olsson, L. 2004. Screening genus *Penicillium* for producers of cellulolytic and xylanolytic enzymes. *Appl. Biochem. Biotechnol.* **113**: 389-401.
- Pitt, J. I. 1979. The genus *Penicillium* and its teleomorphic states *Eupenicillium* and *Talaromyces*. Academic press, London.
- Pitt, J. I., Basilico, J. C., Abarca, M. L. and Lopez, C. 2000. Mycotoxins and toxicogenic fungi. *Med Mycol.* **38**: 41-46.
- Teather, R. M. and Wood, P. J. 1982. Use of congo red-polysaccharide interactions in enumeration and characterization of cellulolytic bacteria from bovine rumen. *Appl. Environ. Microbiol.* **43**: 777-780.
- Techapun, C., Poosaran, N., Waanabe, M. and Sasaki, K. 2003. Thermostable and alkaline-tolerant microbial cellulase-free xylanases produced from agricultural wastes and the properties required for use in pulp bleaching bioprocesses: a review. *Process Biochem.* **38**: 1327-1340.
- Ten, L. N., Im, W. T., Kim, M. K., Kang, M. S. and Lee, S. T. 2004. Development of a plate technique for screening of polysaccharide-degrading microorganisms by using a mixture of insoluble chromogenic substrates. *J. Microbiol. Methods* **56**: 375-382.
- Yoon, J. H., Park, J. E., Suh, D. Y., Hong, S. B., Ko, S. J. and Kim, S. H. 2007. Comparison of dyes for easy detection of extracellular cellulase in fungi. *Mycobiology* **35**: 21-24.