

氮源受限条件下植物病原真菌氮调控基因表达特性

周晓罡^{1,2}, 姚春馨^{1,2}, 丁玉梅^{1,2}, 陶南^{1,2}, 孙茂林^{1,2}, 张绍松^{1,2}

1. 云南省农业科学院生物技术与种质资源研究所, 昆明650223 2. 云南省农业生物技术重点实验室, 昆明650223

ZHOU Xiao-Gang^{1,2}, YAO Chun-Xin^{1,2}, DING Yu-Mei^{1,2}, TAO Nan^{1,2}, SUN Mao-Lin^{1,2}, ZHANG Shao-Song^{1,2}

1. Biotechnology and Genetic Germplasm Institute, Yunnan Academy of Agricultural Sciences, Kunming 650223, China 2. Key Laboratory of Agricultural Biotechnology of Yunnan Province, Kunming 650223, China

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摘要 研究证实植物病害的发生往往是由于植物病原真菌分泌的效应子诱导引起的, 在此过程中, 调控效应基因表达能够了解病原菌的侵染过程。细胞的营养状况据推测对于效应基因的表达起着重要的作用。已有研究表明在氮胁迫条件下相同效应基因的诱导作用在植株体内和体外是一致的, 表明氮源缺乏的环境在植物体进化的早期就已经存在了。文章阐述了在氮受限条件下真菌致病系统中效应基因调控机制及其已经发现的氮调节基因特异性表达研究结果, 通过对比几个病原菌中氮调控基因的功能, 比较寄主植物体内和体外在氮限制条件下基因的诱导效应, 从而揭示出氮的有效性在寄主植物病害发展过程中起到重要作用。

关键词: 病原真菌 氮代谢 氮调控基因

Abstract: It has been confirmed that the occurrence of plant disease is caused by the effector molecules secreted by plant pathogens. The regulation effector gene expression is an important aspect in understanding of the infection process. The nutritional status of cells has been postulated to be a vital role for effector gene expression. Studies have indicated that the induction of the same effector genes during growth *in vitro* as those during growth *in planta* under nitrogen-starved conditions. This showed that the nitrogen poor environment existed in the early time of plant evolution. This paper describes the system in the pathogenesis of several fungal pathogens and nitrogen in the process of gene expression effects from the results of several species by comparing and contrasting the function of nitrogen regulatory genes, as well as by studying plants *in vivo* and *in vitro* gene under nitrogen limitation inductive effect in order to reveal the effectiveness of nitrogen in the development process of host plant disease is an important factor.

Keywords: [pathogenic fungi](#), [nitrogen metabolism](#), [nitrogen-regulated gene](#)

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通讯作者 孙茂林 Email: sun_maolin@tom.com

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- [1] van Baarlen P, van Belkum A, Thomma BPHJ. Disease induction by human microbial pathogens in plant-model systems: potential, problems and prospects. *Drug Discov Today*, 2007, 12(3-4): 167-173.
- [2] Chisholm ST, Coaker G, Day B, Staskawicz BJ. Host-microbe interactions: shaping the evolution of the plant immune response. *Cell*, 2006, 124(4): 803-814.
- [3] Dangl JL. Nibbling at the plant cell nucleus. *Science*, 2007, 315(5815): 1088- 1089.
- [4] Kamoun S. Groovy times: filamentous pathogen effectors revealed. *Curr Opin Plant Biol*, 2007, 10(4): 358-365.
- [5] Baldwin TK, Winnenburg R, Urban M, Rawlings C, Koehler J, Hammond-Kosack KE. The pathogen- host in-teractions database (PHI-base) provides insights into generic and novel themes of pathogenicity. *Mol Plant Microbe Interact*, 2006, 19(12): 1451-1462.

- [6] van der Does HC, Rep M. Virulence genes and the evolution of host specificity in plant-pathogenic fungi. *Mol Plant Microbe Interact*, 2007, 20(10): 1175-1182. 
- [7] Bolton MD, van Esse HP, Vossen JH, de Jonge R, Stule-meijer IJE, Stergiopoulos I, Stulemeijer IJE, van Den Berg GCM, Borrás-Hidalgo O, Dekker HL, De Koster CG, De Wit PJGM, Joosten MHAJ, Thomma BPHJ. The novel *Cladosporium fulvum* lysin motif effector Ecp6 is a virulence factor with orthologues in other fungal species. *Mol Microbiol*, 2008, 69(1): 119-136. 
- [8] Bolton MD, Thomma BPHJ, Nelson BD. *Sclerotinia sclerotiorum* (Lib.) de Bary: biology and molecular traits of a cosmopolitan pathogen. *Mol Plant Pathol*, 2006, 7(1): 1-16.
- [9] van Esse HP, Bolton MD, Stergiopoulos I, de Wit PJGM, Thomma BPHJ. The chitin-binding *Cladosporium fulvum* effector protein Avr4 is a virulence factor. *Mol Plant Microbe Interact*, 2007, 20(9): 1092-1101. 
- [10] van Esse HP, van't Klooster JW, Bolton MD, Yadeta K, van Baarlen P, Boeren S, Vervoort J, de Wita PJGM, Thomma BPHJ. The *Cladosporium fulvum* virulence protein Avr2 inhibits host proteases required for basal defense. *Plant Cell*, 2008, 20(7): 1948-1963. 
- [11] Rooney HC, van't Klooster JW, van der Hoorn RAL, Joosten MHAJ, Jones JDG, de Wit PJGM. *Cladosporium Avr2* inhibits tomato Rcr3 protease required for Cf-2-dependent disease resistance. *Science*, 2005, 308(5729): 1783-1786.
- [12] Bahn YS, Xue C, Indurn A, Rutherford JC, Heitman J, Cardenas ME. Sensing the environment: lessons from fungi. *Nat Rev Microbiol*, 2007, 5(1): 57-69. 
- [13] Talbot NJ, Ebbole DJ, Hamer JE. Identification and characterization of *MPG1*, a gene involved in pathogenicity from the rice blast fungus *Magnaporthe grisea*. *Plant Cell*, 1993, 5(11): 1575-1590.
- [14] van den Ackerveken GFJM, Dunn RM, Cozijnsen AJ, Vossen JP, van den Broek HW, de Wit PJGM. Nitrogen limitation induces expression of the avirulence gene *avr9* in the tomato pathogen *Cladosporium fulvum*. *Mol Gen Genet*, 1994, 243(3): 277-285. 
- [15] Benthin S, Schulze U, Nielsen J, Villadsen J. Growth energetics of *Lactococcus cremoris* FD1 during energy-, carbon-, and nitrogen-limitation in steady state and transient cultures. *Chem Eng Sci*, 1994, 49(5): 589-609.
- [16] Talbot NJ, McCafferty HRK, Ma M, Moore K, Hamer JE. Nitrogen starvation of the rice blast fungus *Magnaporthe grisea* may act as an environmental cue for disease symptom expression. *Physiol Mol Plant Pathol*, 1997, 50(3): 179-195. 
- [17] Donofrio NM, Oh Y, Lundy R, Pan H, Brown DE, Jeong JS, Coughlan S, Mitchell TK, Dean RA. Global gene expression during nitrogen starvation in the rice blast fungus, *Magnaporthe grisea*. *Fungal Genet Biol*, 2006, 43(9): 605-617. 
- [18] Wang Y, Wu J, Park ZY, Kim SG, Rakwal R, Agrawal GK, Kim ST, Kang KY. Comparative secretome investigation of *Magnaporthe oryzae* proteins responsive to nitrogen starvation. *J Proteome Res*, 2011, 10(7): 3136-3148. 
- [19] Pieterse CMJ, Verbakel HM, Spaans JH, Davidse LC, Govers F. Increased expression of the calmodulin gene of the late blight fungus *Phytophthora infestans* during pathogenesis on potato. *Mol Plant Microbe Interact*, 1993, 6(2): 164-172. 
- [20] Pérez-García A, Snoeijs SS, Joosten MHAJ, Goosen T, de Wit PJGM. Expression of the avirulence gene *Avr9* of the fungal tomato pathogen *Cladosporium fulvum* is regulated by the global nitrogen response factor NRF1. *Mol Plant Microbe Interact*, 2001, 14(3): 316-325. 
- [21] Soundararajan S, Jedd G, Li XL, Ramos-Pamplona M, Chua NH, Naqvi NI. Woronin body function in *Magnaporthe grisea* is essential for efficient pathogenesis and for survival during nitrogen starvation stress. *Plant Cell*, 2004, 16(6): 1564-1574. 
- [22] Donofrio NM, Oh Y, Lundy R, Pan H, Brown DE, Jeong JS, Coughlan S, Mitchell TK, Dean RA. Global gene expression during nitrogen starvation in the rice blast fungus, *Magnaporthe grisea*. *Fungal Genet Biol*, 2006, 43(9): 605-617. 
- [23] Brown SH, Yarden O, Gollop N, Chen S, Zveibil A, Belausov E, Freeman S. Differential protein expression in *Colletotrichum acutatum*: changes associated with reactive oxygen species and nitrogen starvation implicated in pathogenicity on strawberry. *Mol Plant Pathol*, 2008, 9(2): 171-190. 
- [24] Stephenson SA, Hatfield J, Rusu AG, Maclean DJ, Mansners JM. *CfDN3*: an essential pathogenicity gene of *Colletotrichum gloeosporioides* necessary to avert a hypersensitive-like response in the host *Stylosanthes guianensis*. *Mol Plant Microbe Interact*, 2000, 13(9): 929-941. 
- [25] 周晓罡, 苏源, 李成云, 丁玉梅, 张绍松, 孙茂林, 李进斌. 氮胁迫条件下稻瘟病菌分泌蛋白致病性分析. *植物病理学报*, 2009, 39(5): 491-500.
- [26] Marzluf GA. Genetic regulation of nitrogen metabolism in the fungi. *Microbiol Mol Biol Rev*, 1997, 61(1): 17-32.
- [27] Caddick MX, Arst HN, Taylor LH, Johnson RI, Brownlee AG. Cloning of the regulatory gene *areA* mediating nitrogen metabolite repression in *Aspergillus nidulans*. *EMBO J*, 1986, 5(5): 1087-1090.
- [28] Stewart V, Vollmer SJ. Molecular cloning of *nlt-2*, a regulatory gene required for nitrogen metabolite repression in *Neurospora crassa*. *Gene*, 1986, 46(2-3): 291-295.
- [29] Morozoc IY, Martinez MG, Jones MG, Caddick MX. A defined sequence within the 3' UTR of the *areA* transcript is sufficient to mediate nitrogen metabolite signalling via accelerated deadenylation. *Mol Microbiol*, 2000, 37(5): 1248-1257. 
- [30] Andrianopoulos A, Kourambas S, Sharp JA, Davis MA, Hynes MJ. Characterization of the *Aspergillus nidulans nmrA* gene involved in nitrogen metabolite repression. *J Bacteriol*, 1998, 180(7): 1973-1977.
- [31] Fu YH, Young JL, Marzluf GA. Molecular cloning and characterization of a negative-acting nitrogen regulatory gene of *Neurospora crassa*.

- [32] Lowry JA, Atchley WR. Molecular evolution of the GATA family of transcription factors: conservation within the DNA-binding domain. *J Mol Evol*, 2000, 50(2): 103-115.
- [33] van Kan JA, van den Ackerveken GF, de Wit PJ. Cloning and characterization of cDNA of avirulence gene *avr9* of the fungal pathogen *Cladosporium fulvum*, causal agent of tomato leaf mold. *Mol Plant Microbe Interact*, 1991, 4(1): 52-59. 
- [34] Thomma BPHJ, Bolton MD, Clergeot PH, de Wit PJGM. Nitrogen controls in *planta* expression of *Cladosporium fulvum Avr9* but no other effector genes. *Mol Plant Pathol*, 2006, 7(2): 125-130. 
- [35] Snoeijers SS, Vossen P, Goosen T, van den Broek HW, de Wit PJGM. Transcription of the avirulence gene *Avr9* of the fungal tomato pathogen *Cladosporium fulvum* is regulated by a GATA-type transcription factor in *Aspergillus nidulans*. *Mol Gen Genet*, 1999, 261(4-5): 653-659. 
- [36] Snoeijers SS, Pérez-García A, Goosen T, de Wit PJGM. Promoter analysis of the avirulence gene *Avr9* of the fungal tomato pathogen *Cladosporium fulvum* is the model filamentous fungus *Aspergillus nidulans*. *Curr Genet*, 2003, 43(2): 96-102.
- [37] Froeliger EH, Carpenter BE. *Nut1*, a major nitrogen regulatory gene in *Magnaporthe grisea*, is dispensable for pathogenicity. *Mol Gen Genet*, 1996, 251(6): 647-656.
- [38] Lau G, Hamer JE. Regulatory genes controlling *Mpg1* expression and pathogenicity in the rice blast fungus *Magnaporthe grisea*. *Plant Cell*, 1996, 8(5): 771-781.
- [39] Tudzynski B, Homann V, Feng B, Marzluf G. Isolation, characterization and disruption of the *areA* nitrogen regulatory gene of *Gibberella fujikuroi*. *Mol Gen Genet*, 1999, 261 (1): 106-114. 
- [40] Mihlan M, Homann V, Liu TW, Tudzynski B. AREA directly mediates nitrogen regulation of gibberellin biosynthesis in *Gibberella fujikuroi*, but its activity is not affected by NMR. *Mol Microbiol*, 2003, 47(4): 975-991. 
- [41] Pellier AL, Lauge R, Veneault-Fourrey C, Langin T. *CLNR1*, the AREA/NIT2-like global nitro-gen regulator of the plant fungal pathogen *Colletotrichum lindemuthianum* is required for the infection cycle. *Mol Microbiol*, 2003, 48(3): 639-655. 
- [42] Divon HV, Ziv C, Davydov O, Yarden O, Fluhr R. The global nitrogen regulator, *FNR1*, regulates fungal nutrition-genes and fitness during *Fusarium oxysporum* pathogenesis. *Mol Plant Pathol*, 2006, 7(6): 485-497.
- [43] Divon HH, Rothan-Denoyes B, Davydov O, Pietro AD, Fluhr R. Nitrogen responsive genes are differentially regulated *in planta* during *Fusarium oxysporum* f. sp. *lycopersici* infection. *Mol Plant Pathol*, 2005, 6(4): 459-470.
- [44] Kim H, Woloshuk CP. Role of AREA, a regulator of ni-trogen metabolism, during colonization of maize kernels and fumonisin biosynthesis in *Fusarium verticillioides*. *Fungal Genet Biol*, 2008, 45(6): 947-953. 
- [45] Burger G, Strauss J, Scazzocchio C, Lang BF. *nirA*, the pathway-specific regulatory gene of ni-trate assimilation in *Aspergillus nidulans*, encodes a putative GAL4-type zinc finger protein and contains four introns in highly con-served regions. *Mol Cell Biol*, 1991, 11(11): 5746-5755.
- [46] Horowitz S, Freeman S, Zveibil A, Yarden O. A defect in *nir1*, a *nirA*-like transcription factor, con-fers morphological abnormalities and loss of pathogenicity in *Colletotrichum acutatum*. *Mol Plant Pathol*, 2006, 7(5): 341-354. 
- [47] Soanes DM, Kershaw MJ, Cooley RN, Talbot NJ. Regulation of the *MPG1* hydrophobin gene in the rice blast fungus *Magnaporthe grisea*. *Mol Plant Microbe Interact*, 2002, 15(12): 1253-1267. 
- [48] Solomon PS, Tan KC, Oliver RP. The nutrient supply of pathogenic fungi: a fertile field for study. *Mol Plant Pathol*, 2003, 4(3): 203-210. 
- [49] Solomon PS, Oliver RP. Evidence that gamma-aminobutyric acid is a major nitrogen source during *Cladosporium fulvum* infection of tomato. *Planta*, 2002, 214(3): 414-420.
- [50] Keon J, Antoniw J, Carzaniga R, Deller S, Ward JL, Baker JM, Beale MH, Hammond-Kosack K, Rudd JJ. Transcrip-tional adaptation of *Mycosphaerella graminicola* to programmed cell death (PCD) of its susceptible wheat host. *Mol Plant Microbe Interact*, 2007, 20(2): 178-193. 
- [51] Tavernier V, Cadiou S, Pageau K, Lauge' R, Reisdorf-Cren M, Langin T, Masclaux-Daubresse C. The plant ni-trogen mobilization promoted by *Colletotrichum lin-demuthianum* in Phaseolus leaves depends on fungus pathogenicity. *J Exp Bot*, 2007, 58(12): 3351-3360. 
- [52] Solomon PS, Oliver RP. The nitrogen content of the to-mato leaf apoplast increases during infection by *Cladosporium fulvum*. *Planta*, 2001, 213(2): 241-249.
- [53] Cove DJ. Chlorate toxicity in *Aspergillus nidulans*. Studies of mutants altered innitrate assimilation. *Mol Gen Genet*, 1976, 146(2): 147-159.
- [54] Howard K, Foster SG, Cooley RN, Caten CE. Disruption, replacement, and cosuppression of nitrate assimilation genes in *Stagonospora nodorum*. *FungalGenet Biol*, 1999, 26(2): 152- 162.
- [55] Coleman M, Henricot B, Arnau J, Oliver RP. Starva-tion-induced genes of the tomato pathogen *Cladosporium fulvum* are also induced during growth *in planta*. *Mol Plant Microbe Interact*, 1997, 10(9): 1106-1109. 
- [56] Pieterse CM, Derkzen AM, Folders J, Govers F. Expression of the *Phytophthora infestans ipiB* and *ipiO* genes *in planta* and *in vitro*. *Mol Gen Genet*, 1994, 244(3): 269-277.
- [57] Foster AJ, Jenkinson JM, Talbot NJ. Trehalose synthesis and metabolism are required at different stages of plant infection by *Magnaporthe*

- [58] Wilson RA, Jenkinson JM, Gibson RP, Littlechild JA, Wang ZY, Talbot NJ. Tps1 regulates the pentose phos-phate pathway, nitrogen metabolism and fungal virulence. *EMBO J*, 2007, 26(15): 3673-3685. crossref
- [59] Pullan ST, Chandra G, Bibb MJ, Merrick M. Genome-wide analysis of the role of GlnR in *Streptomyces venezuelae* provides new insights into global nitrogen regulation in actinomycetes. *BMC Genomics*, 2011, 12(175): 1471-1485.
- [60] Chang JH, Urbach JM, Law TF, Arnold LW, Hu A, Gom-bar S, Grant SR, Ausubel FM, Dangl J L. A high-throughput, near-saturating screen for type III effector genes from *Pseudomonas syringae*. *Proc Natl Acad Sci USA*, 2005, 102(7): 2549-2554. crossref
- [61] Guttman DS, Vinatzer BA, Sarkar SF, Ranall MV, Kettler G, Greenberg JT. A functional screen for the type III (Hrp) secretome of the plant pathogen *Pseudomonas syringae*. *Science*, 2002, 295(5560): 1722-1726.
- [62] Whisson SC, Boevink PC, Moleleki L, Avrova AO, Morales JG, Gilroy EM, Armstrong MR, Grouffaud S, West PV, Chapman S, Hein I, Toth IK, Pritchard L, Birch PRJ. A translocation signal for delivery of oomycete effector proteins into host plant cells. *Nature*, 2007, 450 (7166): 115-118.
- [63] Jiang RHY, Tripathy S, Govers F, Tyler BM. RXLR effector reservoir in two *Phytophthora* species is dominated by a single rapidly evolving super family with more than 700 members. *Proc Natl Acad Sci USA*, 2008, 105(12): 4874-4879. crossref
- [64] Weiland JJ. Production of protease isozymes by *Aphanomyces cochlioides* and *Aphanomyces euteiches*. *Physiol Mol Plant Pathol*, 2004, 65 (5): 225-233. crossref
- [65] Rico A, Preston GM. *Pseudomonas syringae* pv. tomato DC3000 uses constitutive and apoplast-induced nutrient assimilation pathways to catabolize nutrients that are abundant in the tomato apoplast. *Mol Plant Microbe Interact*, 2008, 21(2): 269-282.
- [66] Lee SJ, Kelley BS, Damasceno CM, St John B, Kim BS, Kim BD, Rose JKC. A functional screen to characterize the secretomes of eukaryotic pathogens and their hosts *in planta*. *Mol Plant Microbe Interact*, 2006, 19(12): 1368-1377. crossref

[1] 万英;王旭丽;黄云;邱德文;蒋玲活.细极链格孢菌 $HIS3$ 和 $LEU2$ 基因的cDNA序列的分离和鉴定[J].遗传, 2008,35(4): 251-256