

Article

Comparative Characterization of G Protein α Subunits in *Aspergillus fumigatus*

Yong-Ho Choi ¹, Na-Young Lee ¹, Sung-Su Kim ², Hee-Soo Park ³ and Kwang-Soo Shin^{1,*}

¹ Department of Microbiology, Graduate School, Daejeon University, Daejeon 34520, Korea; youngho1107@gmail.com (Y.-H.C.), 1209leeny@gmail.com (N.-Y.L.)

² Department of Biomedical Laboratory Science, Daejeon University, Daejeon 34520, Korea; sungsu@dju.kr

³ School of Food Science and Biotechnology, Institute of Agricultural Science and Technology, Kyungpook National University, Daegu 41566, Korea; phsoo97@knu.ac.kr

* Correspondence: shinks@dju.kr

Received: 12 February 2020; Accepted: 03 April 2020; Published: date

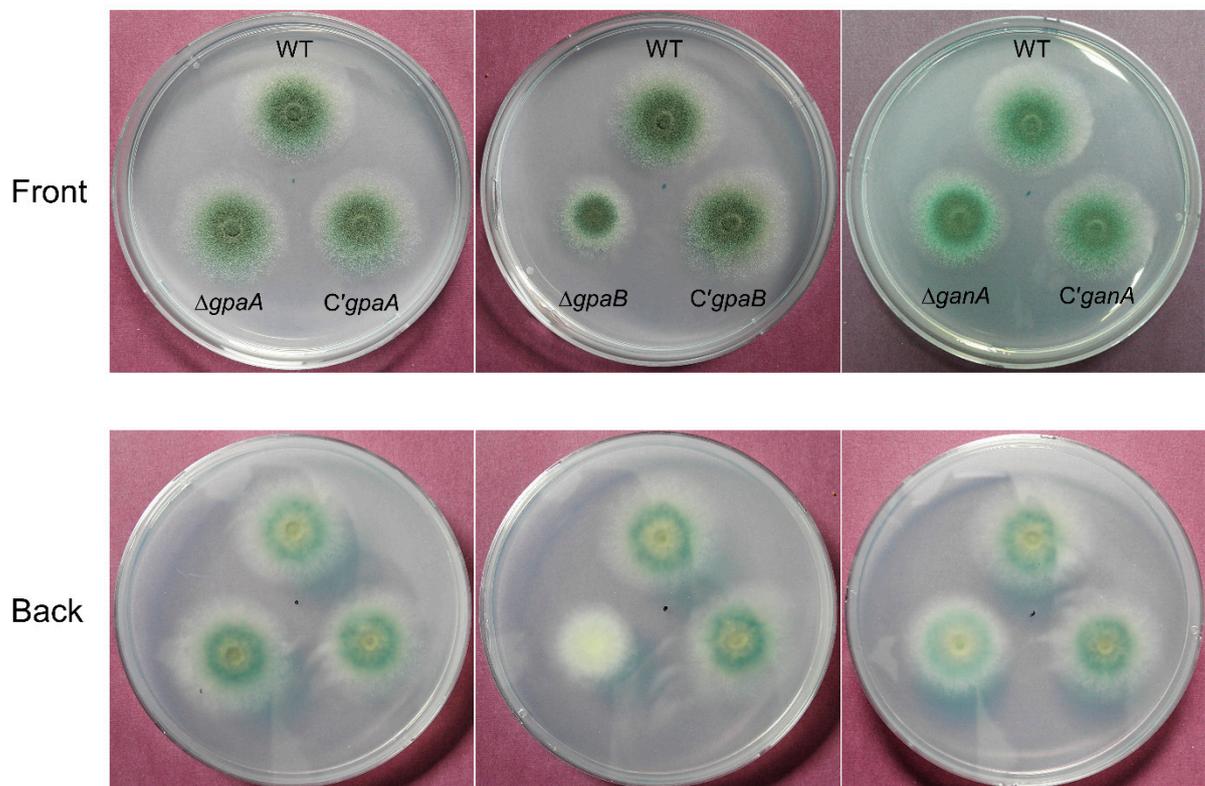


Figure S1. Colony morphologies of WT, mutant, and complemented strains. Each strain was inoculated on solid MMG with supplements and grown for 3 days.

1

Table S1. Oligonucleotides used in this study.

Name	Sequence (5'→3')	Purpose
oligo697	GCAATGTAAAGCTAACGTGCGTG	5' <i>AnpyrG</i> marker
oligo698	TGCCTTTAAGCTTCGGGTAGAG	3' <i>AnpyrG</i> marker
oligo710	CCCACCCCAAATCTACCTA	5' flanking region of <i>gpaA</i>
oligo1030	TTTGTAGGCTTTGGGCTGTTCAAACTTGGCGACAGTTGACGATTCAATT	5' <i>gpaA</i> with <i>AnpyrG</i> tail
oligo1031	CTGATCTACCCCTTGGAACGCAGCATGAGCCATTATACTTGACGGAGTGA	3' <i>gpaA</i> with <i>AnpyrG</i> tail
oligo713	ATCATCCGCATCAGAAGGGA	3' flanking region of <i>gpaA</i>
oligo738	CCACCTCCACTTTTTCTCC	5' nested of <i>gpaA</i>
oligo739	GGCGAGGCTGAATTCGAGAT	3' nested of <i>gpaA</i>
oligo716	GTTGGTGGCCGTATCGGGAC	5' flanking region of <i>gpaB</i>
oligo1032	TTTGTAGGCTTTGGGCTGTTCAAAAACAGCGAAAGGCCTAAAG	5' <i>gpaB</i> with <i>AnpyrG</i> tail
oligo1033	CTGATCTACCCCTTGGAACGCAGCATCATTCCAATCGTTCCATCA	3' <i>gpaB</i> with <i>AnpyrG</i> tail
oligo719	AAGAGCGAGGGGGAGAGTGA	3' flanking region of <i>gpaB</i>
oligo720	GCCCAAGTCACCATGGTCTG	5' nested of <i>gpaB</i>
oligo721	GGGCCACCGTTATGCTCTA	3' nested of <i>gpaB</i>
oligo360	CTCTGTTTCCCTCCACAAA	5' flanking region of <i>ganA</i>
oligo1034	TTTGTAGGCTTTGGGCTGTTCAAGGACAAAGCGTGGAGATAGG	5' <i>ganA</i> with <i>AnpyrG</i> tail
oligo1035	CTGATCTACCCCTTGGAACGCAGCAACCTGAGACGCCAAATTTTC	3' <i>ganA</i> with <i>AnpyrG</i> tail
oligo363	GACAACCCAGCTGATGACTGT	3' flanking region of <i>ganA</i>
oligo364	GTGCTCGAACTCCTTGCTCT	5' nested of <i>ganA</i>
oligo365	ACGGCTGAGCTTCAAGACAT	3' nested of <i>ganA</i>
oligo699	CCATGTGTGTCGAGTCCCTTC	5' <i>eflA</i> for qRT-PCR normalization
oligo700	GAACGTACAGCAACAGTCTGG	3' <i>eflA</i> for qRT-PCR normalization
oligo256	TTCCAAGCAGAGCTTGTCAC	5' <i>brlA</i> for qRT-PCR
oligo257	CCAGTTCTTTGCACTTGAA	3' <i>brlA</i> for qRT-PCR
oligo303	GCTACCACTCTGCATCCTCA	5' <i>abaA</i> for qRT-PCR
oligo304	TACGAGCTCCAGCATGATTC	3' <i>abaA</i> for qRT-PCR
oligo271	AAATCCATCACATCCACCCT	5' <i>gliZ</i> for qRT-PCR
oligo272	GGTTGTTTCATGGTCAGTTGC	3' <i>gliZ</i> for qRT-PCR
oligo1462	ACGTACCGCATGTTTGATGT	5' <i>gpaA</i> for qRT-PCR
oligo1463	CAGCTGATCGTACTCCGAGA	3' <i>gpaA</i> for qRT-PCR
oligo1464	CTCTGGAATGATCCATGCAC	5' <i>gpaB</i> for qRT-PCR
oligo1465	CAAAGAAATATGGCGCAGAA	3' <i>gpaB</i> for qRT-PCR
oligo1466	CCCGTTTAAGGACAACCTGGT	5' <i>ganA</i> for qRT-PCR
oligo1467	ACCAACGTCCATCATTCTGA	3' <i>ganA</i> for qRT-PCR

2



© 2020 by the authors. Submitted for possible open access publication under the terms and conditions of the Creative Commons Attribution (CC BY) license (<http://creativecommons.org/licenses/by/4.0/>).