



# *Sclerotinia sclerotiorum*

J. , A. , F. , \* , \*

College of Plant Science and Technology, Huazhong Agricultural University, Wuhan 430070, China

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## ARTICLE INFO

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## ABSTRACT

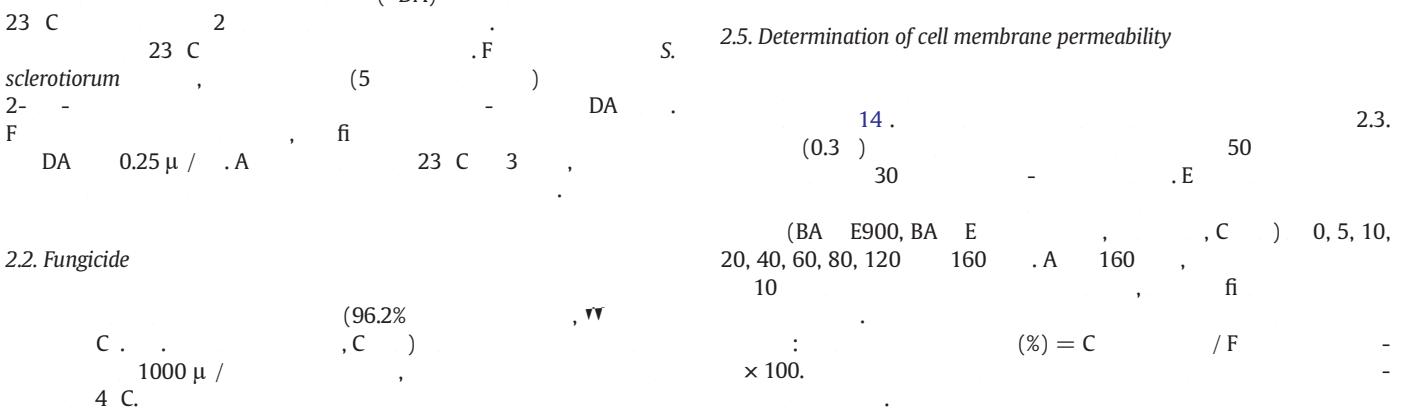
### *Sclerotinia sclerotiorum*

			2.3. Determination of glycerol content in mycelia							
			G	14,45	B	fl	,			
8.44	A	(V238A)	SsOs1	C	4	0.05	/	3.5		0.05 /
<i>S. sclerotiorum</i>	17.			fi		50			10	
				0.006, 0.008,		0.01	/		0, 0.0025, 0.003, 0.004, 0.005,	
				12					100	
				.A		630			fi	
<i>S. sclerotiorum</i>	24.		fl	fi					fi	
<i>Neurospora crassa</i>										
		34	A	DA						
			DCF	Bo-						
<i>trytis cinerea</i>	11,35.									
A										
17,24,37,39,40,48	.VV	S. sclerotiorum,	DCF	10,		20				
		14-17,	DCF			50				
<i>Sclerotiorum</i>			fi	S.		80	C	15	.A	
				630						
		(D)								
		(Sshk, SsYpd, Sssk1)	A							
SsHog)				(Sssk2, SsPbs,						

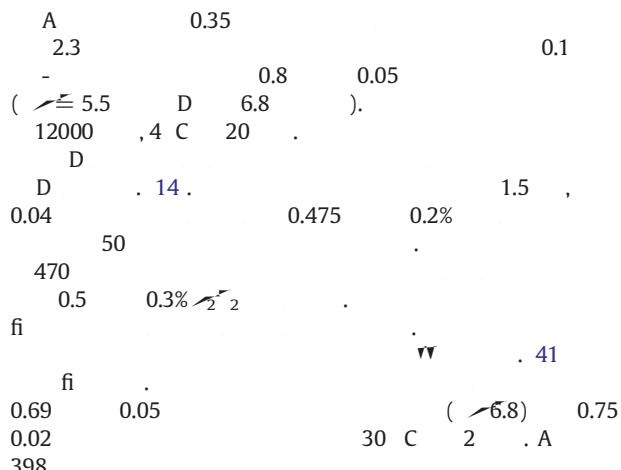
## 2. Materials and methods

### 2.1. Isolates of *S. sclerotiorum*

<i>sclerotiorum</i>			(G2, G3, G4, G5)	2011	48	A	S.			
			fi							



## 2.6. Determination of peroxidase and polyphenol oxidase activity



## 2.7. Cloning and sequencing of genes involved in two-component HK system and MAP kinase cascades

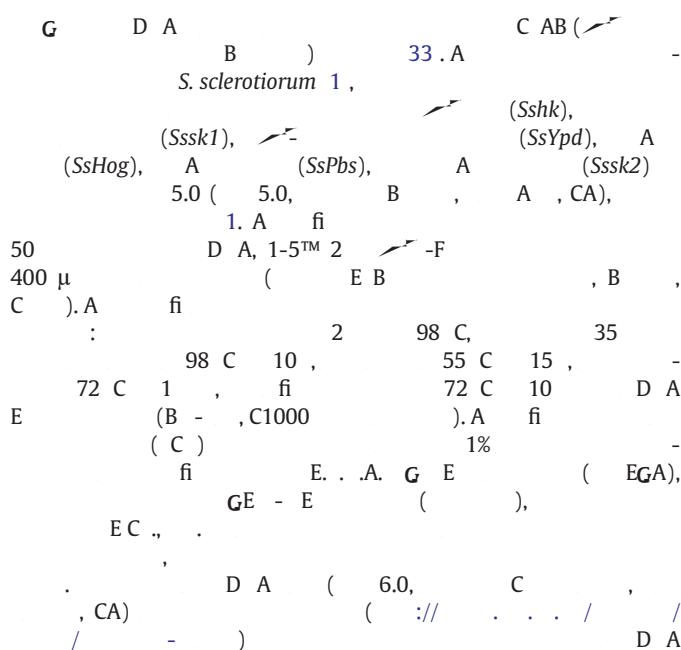
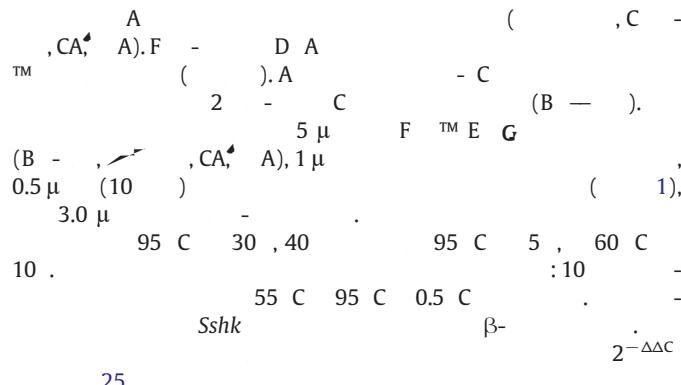


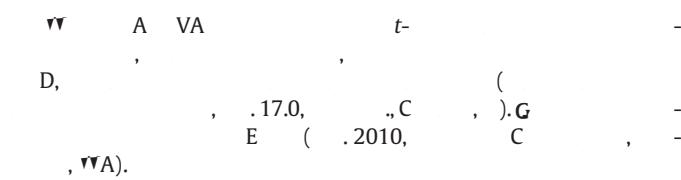
Table 1

(5'-3')		
F1	A GGGGGACAC AGCA AGC C	Sshk
1	AG ACAGG CC GCAAG GG	
-F	A GGC GAA CG AGAG	Sshog
	GACCA CCACCGA GGGCG	
-F	A GACAGA AA CC CAA AGA CG	Sspbs
-	CA CAAACCC GC CC C CA	
2-F	A GGAGCG CCACA A G	Sssk2
2-	AC GAGAGC CCAC A CGC G	
1-F	GACGA AGGCGA C CAAAACACGA	Sssk1
1-	A CCCACAACCACC CA A CG	
-F	A G CC C CCAC C ACC C	Ssypd
-	G GGCA AAAAC C CAAG	
β-F	GGA GC CC GACCAAG	β-
β-	AGCGGCCA CA G C AGG	
-F	AAGGG AGGGAAG AAGGCAA C	Sshk
-	G A AGGGCCAG GG AG G A G	

## 2.8. Quantitative RT-PCR

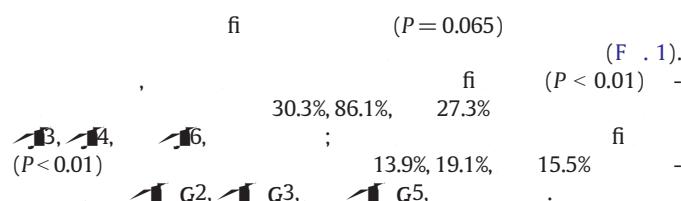


## 2.9. Data analysis



## 3. Results

### 3.1. Glycerol content



### 3.2. Cell membrane permeability and oxalic acid content

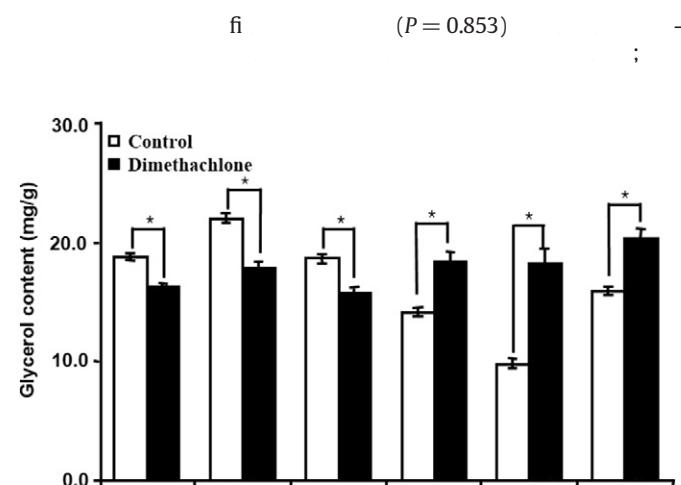
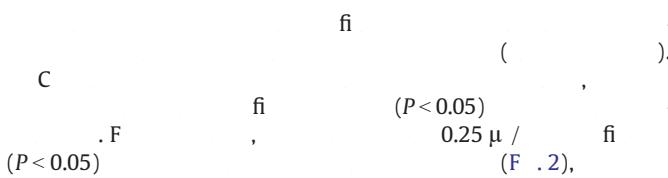
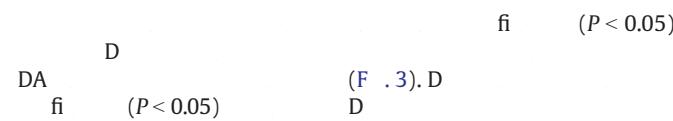


Fig. 1. G  
sclerotiorum. \*\*\*(E)  
Sclerotinia  
( $\alpha = 0.05$ ). V



### 3.3. Peroxidase and polyphenol oxidase activity



### 3.4. Newly identified point mutations in the Sshk and Sspbs genes

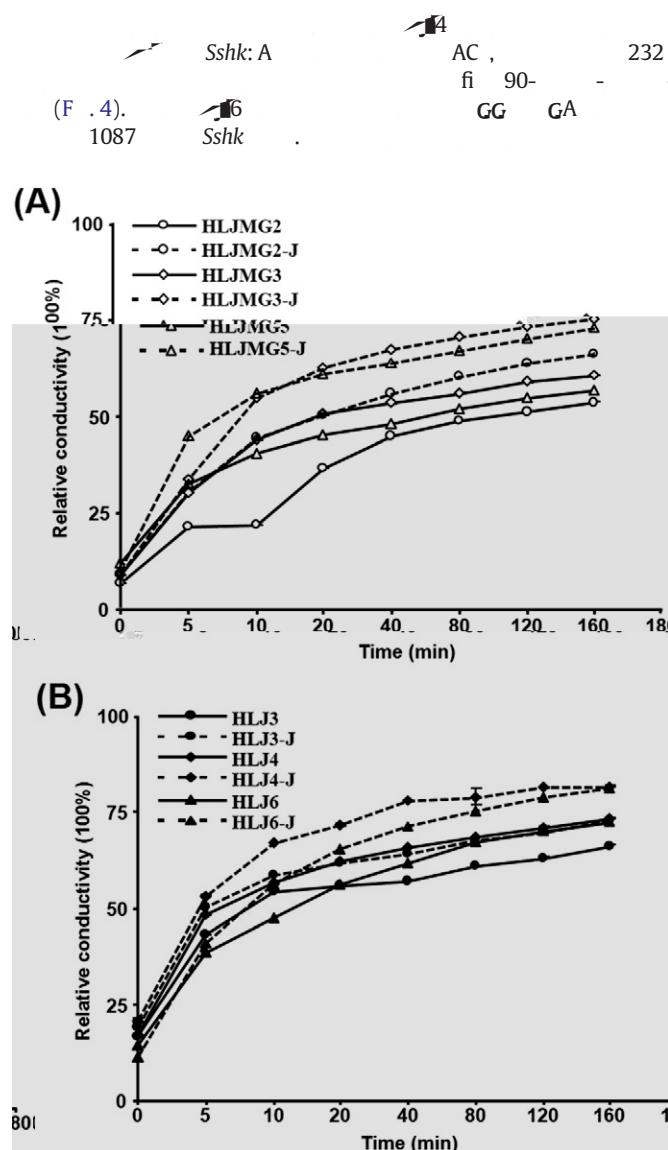


Fig. 2.

*Sclerotinia sclerotiorum* (A)

(B)

(E)

t-

E

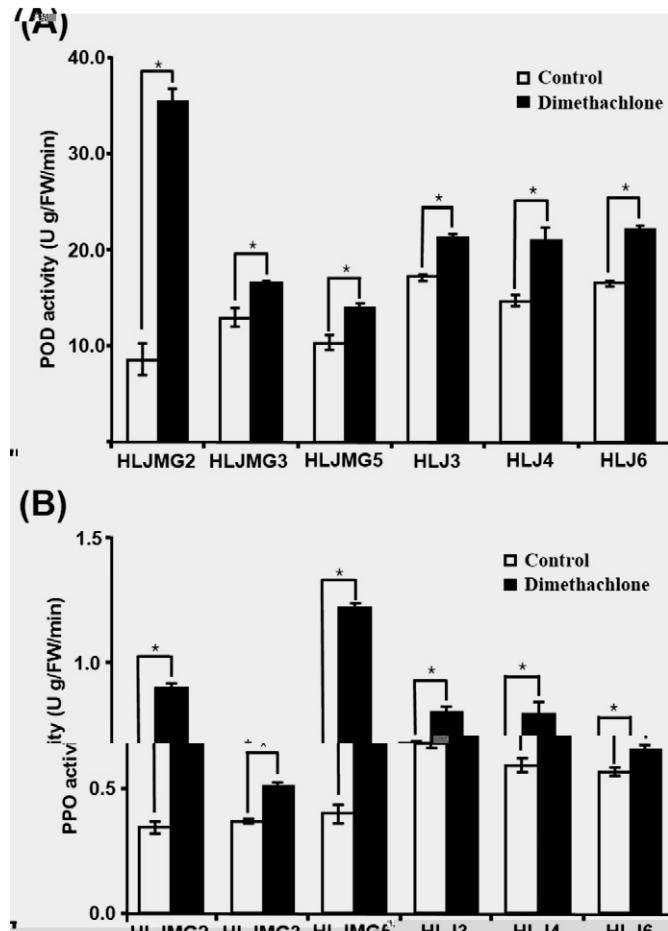


Fig. 3. (D) (A)

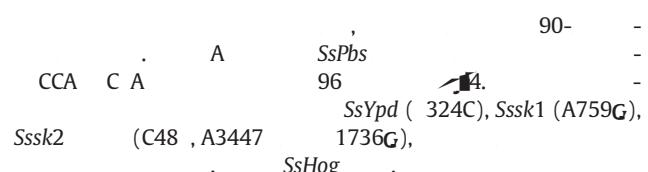
fi (a = 0.05). V

( ) (B)

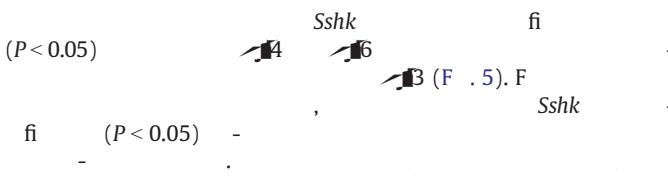
*Sclerotinia sclerotiorum*. \*\*\*

E

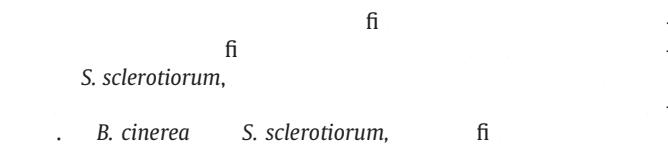
t-

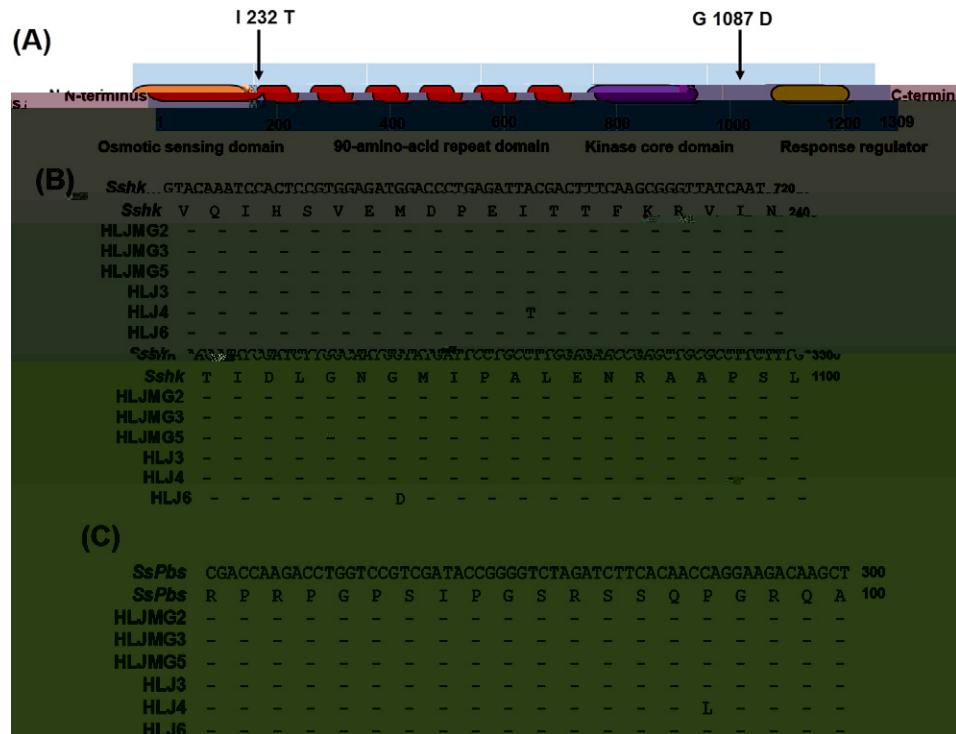


### 3.5. Transcription levels of the Sshk gene

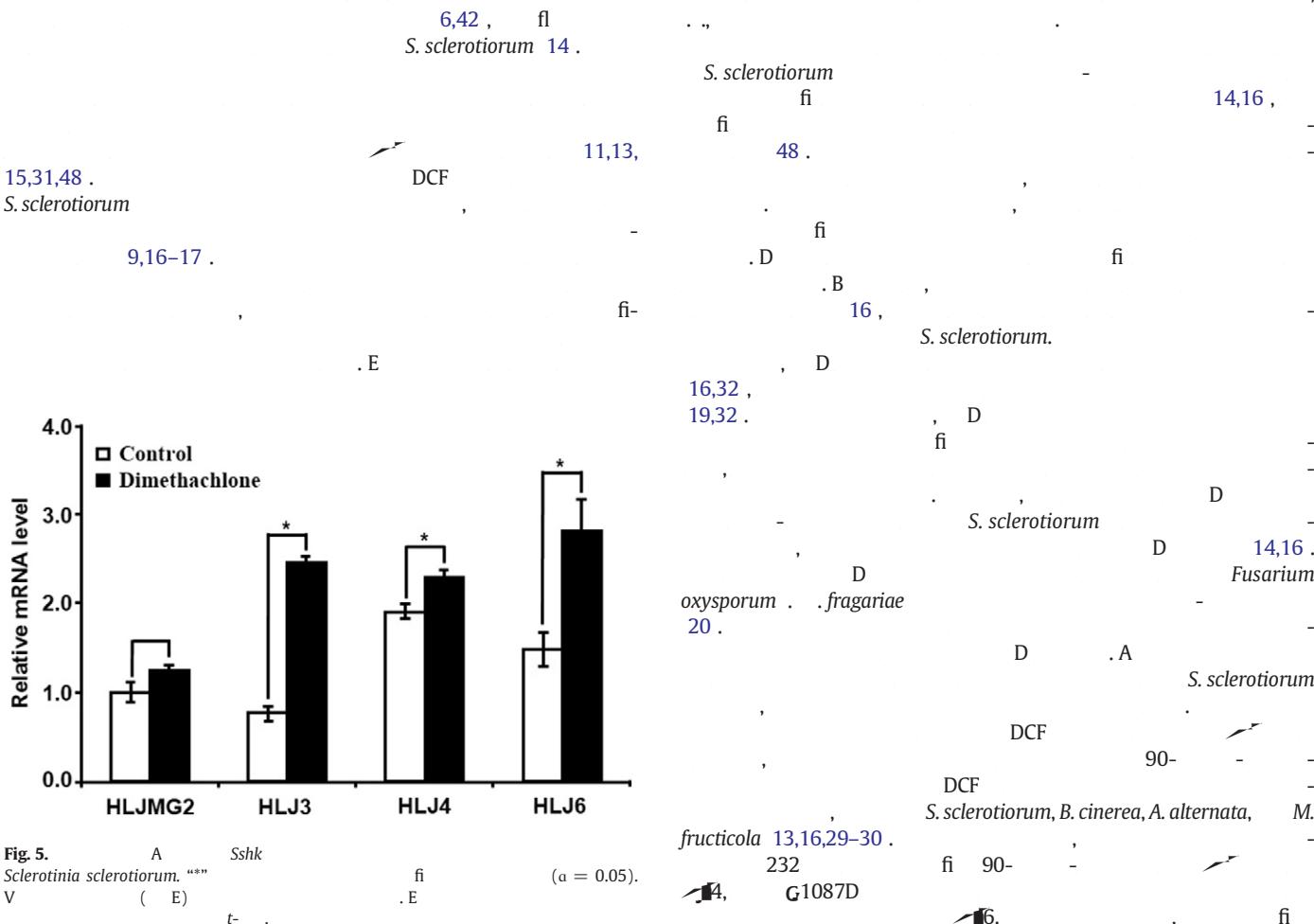


### 4. Discussion





**Fig. 4.** C Sshk ( 1G 12694.3) .(A) D A .(A) 14 .16 Sshk ( .(C) Sshk ( SsPbs ).(B) B 96 Sshk B A .(B) 14 .16



<i>S. sclerotiorum</i> , A 1040 ( <i>Bos1</i> ) <i>B. cinerea</i> 31. <i>SsYpd</i> , <i>Sssk1</i> , A	CGA DCF -	GA -	6 <b>C</b> , 7 ..C., .C. 8 C.C., .C. 9 .C., .C., <b>V</b> . 10 <b>C</b> , .C., .C., D. 11 <b>V</b> .C., .E.B.	<b>C</b> , B.G., -	<b>V</b> , -	- -	<i>Sclerotinia sclerotiorum</i> C 2008. .C 2 (2003) 1151–1161. .117 (1998) 723–731. <i>Sclerotinia sclerotiorum</i> C 32 (2010) 534–539. <i>Alternaria brassicicola</i> , ( <i>Botrytis cinerea</i> ), F <b>G</b> , B	<i>Sclerotinia sclerotiorum</i> C 2008. .C 2 (2003) 1151–1161. .117 (1998) 723–731. <i>Sclerotinia sclerotiorum</i> C 32 (2010) 534–539. <i>Alternaria brassicicola</i> , ( <i>Botrytis cinerea</i> ), F <b>G</b> , B
( D 1)  <i>Sshk</i>  <i>sclerotiorum</i> , A. longipes sclerotiorum 17.  <i>S. sclerotiorum</i> .	fi Sshk V238A	21 G2 Sshk	fi Sshk	21 A	fi Sshk	21 Sshk	fi Sshk	21 Sshk
<b>G1087D)</b>  <i>S. sclerotiorum</i> . sclerotiorum 16,20,29–31 .	Sshk	D	( 232	S.	S.	S.	S.	S.
  <i>S. sclerotiorum</i> . sclerotiorum	  fi Sshk	  Sshk	  D, ( )	  S.	  D, ( )	  13,	  A	  <i>S. sclerotiorum</i>

## Competing interests

## Acknowledgements

C (31371964).

## References

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