

Table S1. Comparison of the sequence from isolates with sequences deposited in databases ("blast").

Isolated		%	E	Identification	Access number	Final Identification
		Similarity	Value			
1	2AO	99.76	0.0	Fusarium camptoceras	KJ782411.1	Fusarium sp.
		99.76	0.0	Fusarium equiseti	AB425996.1	
		99.76	0.0	Fusarium oxysporum	JN400714.1	
		99.76	0.0	Fusarium sp.	HQ023205.1	
		99.52	0.0	Fusarium equiseti	KP026922.1	
2	2AQ	99.19	0.0	Trametes polyzona	JN164979.1	Trametes sp.
		98.49	0.0	Trametes polyzona	MH855813.1	
		98.48	0.0	Trametes polyzona	KT120039.1	
		98.39	0.0	Trametes polyzona	JN164978.1	
3	2AN	97.19	0.0	Fusarium sp.	HQ023200.1	Fusarium sp.
		96.68	0.0	Fusarium sp.	U85548.1	
		95.62	0.0	Fusarium robustum	NR_159851.1	
		95.62	0.0	Fusarium robustum	MH861018.1	
		95.56	0.0	Fusarium sp.	HQ023198.1	
4	2BD	99.39	0.0	Fusarium equiseti	KP026922.1	Fusarium sp.
		99.59	0.0	Fusarium equiseti	AB425996.1	
		99.59	0.0	Fusarium oxysporum	JN400714.1	
		99.38	0.0	Fusarium equiseti	GQ365157.1	
5	2M	98.41	0.0	Aspergillus caelatus	MH862672.1	Aspergillus sp.
		98.27	0.0	Aspergillus caelatus	KM613140.1	
		98.12	0.0	Aspergillus pseudotamarii	MH862673.1	
		98.12	0.0	Aspergillus pseudotamarii	DQ467987.1	
		98.12	0.0	Aspergillus pseudotamarii	DQ467986.1	
6	2S	99.79	0.0	Phoma sp.	KP966098.1	Phoma sp.
		99.79	0.0	Epicoccum sorghinum	MN215627.1	
		99.79	0.0	Epicoccum sorghinum	MN215624.1	
		99.79	0.0	Epicoccum sorghinum	MN215623.1	
		99.79	0.0	Epicoccum sorghinum	MN215622.1	
7	2AM	100.00	0.0	Alternaria doliconidium	NR_158361.1	Alternaria sp.
		100.00	0.0	Alternaria doliconidium	MG828864.1	
		100.00	0.0	Alternaria alternata	LN808867.1	
		100.00	0.0	Alternaria sp.	HQ649963.1	
		99.81	0.0	Alternaria sp.	MG209668.1	

8	2E	99.79	0.0	Aspergillus sydowii	KP973600.1	Aspergillus sp.
		99.37	0.0	Aspergillus sydowii	LN898721.1	
		99.37	0.0	Aspergillus sydowii	LN898720.1	
		99.37	0.0	Aspergillus sydowii	LN898719.1	
		99.37	0.0	Aspergillus sydowii	LN898716.1	
9	2AJ	99.57	0.0	Fusarium oxysporum	MG356946.1	Fusarium sp.
		99.57	0.0	Fusarium oxysporum	KU382653.1	
		99.57	0.0	Fusarium sp.	GQ365161.1	
		99.57	0.0	Fusarium oxysporum	GQ365156.1	
		99.57	0.0	Fusarium sp.	EU363511.1	
10	1BH	99.79	0.0	Gibberella moniliformis	EU151467.1	Fusarium sp.
		99.79	0.0	Gibberella fujikuroi	AY188916.1	
		99.79	0.0	Melanospora zamiae	KY628680.1	
		99.58	0.0	Gibberella moniliformis	EU151476.1	
		99.58	0.0	Fusarium denticulatum	U61680.1	
11	2BH	99.42	0.0	Bipolaris gossypina	MH864410.1	Bipolaris sp.
		99.02	0.0	Bipolaris sorokiniana	KR812216.1	
		98.65	0.0	Curvularia sp.	KR093890.1	
		98.23	0.0	Bipolaris salviniae	MH864690.1	
		98.23	0.0	Bipolaris salviniae	MH864687.1	
12	2L	99.78	0.0	Curvularia sp.	MG976352.1	Curvularia sp.
		99.78	0.0	Curvularia soli	NR_152503.1	
		99.78	0.0	Curvularia soli	KY905679.1	
		99.78	0.0	Curvularia sp.	KT777714.1	
		99.78	0.0	Curvularia sp.	KR093913.1	
13	2BB	100.00	0.0	Aspergillus sp.	KU751784.1	Aspergillus sp.
		92.11	2e-151	Aspergillus indologenus	MH861245.1	
		92.11	2e-151	Aspergillus aculeatus	AY585557.1	
		91.71	2e-151	Aspergillus uvarum	MH863120.1	
14	1AI	99.77	0.0	Colletotrichum acidiae	MG996506.1	Colleototrichum sp.
		99.77	0.0	Colletotrichum acidiae	NR_160841.1	
		97.81	0.0	Colletotrichum capsici	EU056738.1	
15	2AL	99.37	0.0	Fusarium oxysporum	MG356946.1	Fusarium sp.
		99.37	0.0	Fusarium oxysporum	KU382653.1	
		99.37	0.0	Fusarium sp.	GQ365161.1	
		99.37	0.0	Fusarium oxysporum	GQ365156.1	
		99.37	0.0	Fusarium sp.	EU363511.1	

16	2AH	99.79	0.0	Fusarium chlamydo- sporum	KM076600.1	Fusarium sp.
		99.58	0.0	Fusarium equiseti	AY213655.1	
		99.37	0.0	Fusarium sp.	MH582470.1	
		99.37	0.0	Fusarium sp.	MH582467.1	
		99.37	0.0	Fusarium sp.	MH582450.1	
17	1L	99.61	0.0	Aspergillus giganteus	MH857126.1	Aspergillus sp.
		99.59	0.0	Aspergillus giganteus	KR296897.1	
		99.59	0.0	Aspergillus giganteus	NR_135403.1	
		99.59	0.0	Aspergillus giganteus	EF669987.1	
		99.59	0.0	Aspergillus giganteus	EF669928.1	
18	2F	99.59	0.0	Aspergillus fumigatus	MH864623.1	Aspergillus sp.
		99.59	0.0	Aspergillus fumigatus	MF476024.1	
		99.59	0.0	Aspergillus fumigatus	KJ863511.1	
		99.59	0.0	Aspergillus fumigatus	KP724993.1	
		99.59	0.0	Aspergillus fumigatus	KP724992.1	
19	2AC	99.00	0.0	Fusarium subglutinans	EU117221.1	Fusarium sp.
		99.00	0.0	Fusarium proliferatum	EF589878.1	
		99.00	0.0	Fusarium sp.	KM104592.1	
20	2AU	99.58	0.0	Colletotrichum fusiforme	NR_160841.1	Colletotrichum sp.
		99.38	0.0	Colletotrichum acidiae	MG996506.1	
		97.62	0.0	Colletotrichum capsici	EU056738.1	
		97.43	0.0	Colletotrichum capsici	LC152972.1	
		97.43	0.0	Glomerella truncata	FJ172230.1	
21	2AR	98.96	0.0	Colletotrichum acidiae	NR_160841.1	Colletotrichum sp.
		98.76	0.0	Colletotrichum acidiae	MG996506.1	
		97.24	0.0	Colletotrichum capsici	EU056738.1	
		97.05	0.0	Colletotrichum capsici	LC152972.1	
22	1AN	98.39	0.0	Fusarium incarnatum	MF356578.1	Fusarium sp.
		98.20	0.0	Fusarium chlamydo- sporum	KM076600.1	
		98.20	0.0	Fusarium sp.	GQ352485.1	
		98.20	0.0	Fusarium incarnatum	MH865893.1	
23	1BL	99.56	0.0	Fusarium proliferatum	MG228402.1	Fusarium sp.
		99.56	0.0	Fusarium proliferatum	MG228401.1	
		99.56	0.0	Fusarium sp.	MK432970.1	
		99.13	0.0	fusarium oxysporum	JN400715.1	
		99.13	0.0	fusarium oxysporum	JN400698.1	

24	1K	99.16	0.0	Penicillium citrinum	KM613148.1	Penicillium sp.
		98.57	0.0	Penicillium citrinum	KX066022.1	
		98.57	0.0	Penicillium citrinum	KX066021.1	
		98.37	0.0	Penicillium sp.	GU017521.1	
		98.37	0.0	Penicillium griseofulvum	FJ717687.1	
25	2AS	99.58	0.0	Volutella lini	JQ693169.1	Volutella sp.
		99.41	0.0	Volutella lini	JQ647452.1	
		93.35	0.0	volutella consors	KM231768.1	
26	2AV	98.96	0.0	Fusarium sp.	HQ023200.1	Fusarium sp.
		98.16	0.0	Fusarium robustum	NR_159851.1	
		98.16	0.0	Fusarium robustum	MH861018.1	
		97.96	0.0	Fusarium robustum	U85539.1	
		97.69	0.0	Gibberella pulicaris	DQ026011.1	
27	2AZ	93.33	0.0	Gibberella moniliformis	EU151482.1	NI
		93.26	0.0	Fusarium sp.	KU382552.1	
28	2BA	98.13	0.0	Fusarium equiseti	HQ649909.1	Fusarium sp.
		98.13	0.0	Fusarium equiseti	HQ649904.1	
		98.13	0.0	Fusarium equiseti	HQ649906.1	
		98.13	0.0	Fusarium equiseti	HQ649905.1	
		97.93	0.0	Fusarium caatingaense	MH668833.1	
29	2BF	98.38	0.0	Fusarium sp.	GQ352485.1	Fusarium sp.
		98.38	0.0	Fusarium chlamydo- sporum	MF072670.1	
		98.38	0.0	Fusarium incarnatum	MF356578.1	
30	2N	98.25	0.0	Curvularia sp.	HQ909079.1	Curvularia sp.
		98.24	0.0	Curvularia clavata	MN592977.1	
		98.24	0.0	Curvularia sp.	MG976352.1	
		98.24	0.0	Curvularia soli	NR_152503.1	
		98.24	0.0	Curvularia soli	KY905679.1	
31	2Y	94.91	0.0	Byssochlamys spectabilis	MH856887.1	NI
		94.91	0.0	Paecilomyces variotii	JF766675.1	
		94.91	0.0	Paecilomyces variotii	GU968674.1	
		94.91	0.0	Paecilomyces variotii	FJ389930.1	
		94.91	0.0	Paecilomyces variotii	AF033395.1	
32	2O	98.90	0.0	Dothideales sp.	HQ608008.1	Epicoccum sp.
		98.90	0.0	Epicoccum sp.	MG976442.1	
		98.90	0.0	Epicoccum sp.	MG976431.1	
		98.90	0.0	Epicoccum sp.	LT592959.1	

		98.90	0.0	Epicoccum sp.	LT592927.1	
33	2R	95.63	0.0	Montagnula cirsii	KX274242.1	Montagnula sp.
		95.49	0.0	Montagnula scabiosae	NR_155378	
		95.49	0.0	Montagnula scabiosae	KT443907	
34	2B	99.29	0.0	Trichoderma hamatum	MH864073.1	Trichoderma sp.
		99.12	0.0	Trichoderma harzianum	KR868288.1	
		99.12	0.0	Hypocrea lixii	HQ011501.1	
		99.12	0.0	Trichoderma harzianum	KR868336.1	
		99.12	0.0	Trichoderma harzianum	KR868321.1	
35	2J	99.08	0.0	Fusarium oxysporum	DQ068983.1	Fusarium sp.
		98.87	0.0	Fusarium oxysporum	KX786247.1	
		98.87	0.0	Fusarium oxysporum	KX786244.1	
		98.87	0.0	Fusarium oxysporum	KX786241.1	
		98.87	0.0	Fusarium oxysporum	LN828174.1	
36	1AS	99.31	0.0	Fusarium solani	MH348949.1	Fusarium sp.
		99.31	0.0	Fusarium solani	MH348946.1	
		99.31	0.0	Fusarium solani	MH348944.1	
		99.30	0.0	Fusarium solani	LN828130.1	
		98.85	0.0	Fusarium solani	MH865999.1	
37	2AW	98.88	0.0	Clonostachys sp.	KY413708.1	Clonostachys sp.
		98.88	0.0	Clonostachys sp.	KY413698.1	
		98.62	0.0	Gliocladium sp.	AF139857.1	
		98.42	0.0	Clonostachys sp.	MH267864.1	
		98.42	0.0	Clonostachys rogersoni- ana	KR812215.1	
38	2A	98.99	0.0	Cladosporium cladospori- oides	MF044039.1	Cladosporium sp.
		98.99	0.0	Cladosporium tenuissi- mum	MF473296.1	
		98.99	0.0	Cladosporium tenuissi- mum	MF473282.1	
		98.99	0.0	Cladosporium tenuissi- mum	MF473290.1	
		98.99	0.0	Cladosporium tenuissi- mum	LN834398.1	
39	1G	99.77	0.0	Aspergillus sydowii	MH858253.1	Aspergillus sp.
		99.77	0.0	Aspergillus sydowii	MH854859.1	
		99.77	0.0	Aspergillus sp.	MG976613.1	
		99.77	0.0	Aspergillus sydowii	LN898735.1	
		99.77	0.0	Aspergillus sydowii	LN898734.1	

40	1M	99.60	0.0	Aspergillus sydowii	MH854859.1	Aspergillus sp.
		99.60	0.0	Aspergillus sp.	MG976613.1	
		99.60	0.0	Aspergillus sydowii	LN898735.1	
		99.60	0.0	Aspergillus sydowii	LN898734.1	
		99.60	0.0	Aspergillus sydowii	KT826624.1	
41	1J	96.30	0.0	Cladosporium cladosporioides	MF072635.1	Cladosporium sp.
		96.30	0.0	Cladosporium cladosporioides	MF072642.1	
		96.30	0.0	Cladosporium sp.	LN809014.1	
		96.30	0.0	Cladosporium cladosporioides	LN809009.1	
		96.30	0.0	Cladosporium cladosporioides	LN809000.1	
42	1AE	99.22	0.0	Aspergillus versicolor	AM883155.1	Aspergillus sp.
		99.04	0.0	Aspergillus tabacinus	LN898739.1	
		99.04	0.0	Aspergillus tabacinus	LN898738.1	
		99.04	0.0	Aspergillus tabacinus	LN898737.1	
		99.04	0.0	Aspergillus tabacinus	LN898736.1	
43	1AM	98.23	0.0	Fusarium chlamydosporum	AY213655.1	Fusarium sp.
		98.01	0.0	Fusarium chlamydosporum	KM076600.1	
		97.79	0.0	Fusarium incarnatum	MH865897.1	
		97.79	0.0	Fusarium incarnatum	MH865895.1	
		97.79	0.0	Fusarium camptoceras	MH858537.1	
44	2AA	97.94	0.0	Trichoderma spirale	FJ861463.1	Trichoderma sp.
		97.38	0.0	Trichoderma harzianum	FJ884153.1	
		97.37	0.0	Trichoderma spirale	KR296909.1	
		97.22	0.0	Trichoderma spirale	KT278911.1	
		97.05	0.0	Trichoderma spirale	KP263614.1	
45	1AG	95.82	0.0	Bionectria sp.	HM849058.1	NI
		95.80	0.0	Clonostachys rosea	KJ619987.1	
		95.80	0.0	Bionectria ochroleuca	AY669327.1	
		95.76	0.0	Bionectria sp.	AB734437.1	
46	2X	97.79	0.0	Roussoella solani	LC195220.1	Roussoella sp.
		97.79	0.0	Roussoella solani	LC195219.1	
		97.79	0.0	Roussoella solani	LC195218.1	
		97.78	0.0	Neoroussoella leucaenae	NR_165226.1	
		97.78	0.0	Neoroussoella entadae	MK347786.1	

47	1E	99.81	0.0	Curvularia sp.	MG976352.1	Curvularia sp.
		99.81	0.0	Curvularia soli	NR_152503.1	
		99.81	0.0	Curvularia soli	KY905679.1	
		99.81	0.0	Curvularia sp.	KR093913.1	
		99.81	0.0	Curvularia senegalensis	HG779001.1	
48	1AU	100.00	0.0	Fusarium camptoceras	KJ782411.1	Fusarium sp.
		99.59	0.0	Fusarium equiseti	AB425996.1	
		99.59	0.0	Fusarium oxysporum	JN400714.1	
		99.59	0.0	Fusarium sp.	HQ023205.1	
		99.39	0.0	Fusarium sp.	EF120420.1	
49	2I	99.82	0.0	Mucor sp.	JF723592.1	Mucor sp.
		99.65	0.0	Mucoraceae sp.	KX148752.1	
		99.33	0.0	Mucor fragilis	FN650655.1	
		98.64	0.0	Mucor fragilis	KY047150.1	
		98.64	0.0	Mucor fragilis	KY047147.1	
50	2G	100.00	0.0	Penicillium sp.	GU017521.1	Penicillium sp.
		100.00	0.0	Penicillium sp.	GU017519.1	
		100.00	0.0	Penicillium griseofulvum	FJ717687.1	
		100.00	0.0	Penicillium citrinum	MH864240.1	
		100.00	0.0	Penicillium citrinum	MH858380.1	
51	1AH	99.46	0.0	Pleosporales sp.	KP306958.1	NI
		99.29	0.0	Roussoella solani	LC195220.1	
		99.28	0.0	Pleosporales sp.	KP263105.1	
		99.28	0.0	Ascomycota sp.	JX966643.1	
		99.25	0.0	Roussoella solani	MH923251.1	
52	1AY	97.89	0.0	Myrothecium roridum	KJ815095.1	Myrothecium sp.
		97.73	0.0	Myrothecium roridum	KJ174523.1	
		97.57	0.0	Paramyrothecium roridum	MH856665.1	
		97.57	0.0	Paramyrothecium roridum	KU846301.1	
		97.57	0.0	Myrothecium roridum	JF343832.1	
53	1BG	95.16	0.0	Myrothecium inundatum	KU856653.1	Myrothecium sp.
		94.58	0.0	Myrothecium inundatum	LC387258.1	
		94.58	0.0	Myrothecium inundatum	JN851023.1	
54	1R	97.93	0.0	Colletotrichum dematium	AB046608.1	Colletotrichum sp.
		97.93	0.0	Colletotrichum capsici	EU056740.1	
		97.76	0.0	Colletotrichum dematium	AB042310.1	

		97.60	0.0	Colletotrichum truncatum	KC460308.1	
		97.60	0.0	Colletotrichum truncatum	AJ301945.1	
55	1U	99.30	0.0	Aspergillus tabacinus	LN898739.1	Aspergillus sp.
		99.30	0.0	Aspergillus sp.	FJ770064.1	
		99.29	0.0	Aspergillus versicolor	AM883155.1	
		99.29	0.0	Aspergillus tabacinus	LN898736.1	
		99.28	0.0	Aspergillus tabacinus	FR733851.1	
56	1W	97.59	0.0	Colletotrichum dematium	AB046608.1	Colletotrichum sp.
		97.59	0.0	Colletotrichum capsici	EU056740.1	
		97.42	0.0	Colletotrichum dematium	Ab042310.1	
		97.26	0.0	Colletotrichum truncatum	MG677594.1	
57	1Z	99.35	0.0	Nigrograna mackinnonii	NR_132037.1	NI
		99.18	0.0	Biatriospora mackinnonii	LN626660.1	
		98.79	0.0	Nigrograna mackinnonii	LN626659.1	
		98.70	0.0	Pleosporales sp.	KP263091.1	
		98.70	0.0	Nigrograna mackinnonii	KC288117.1	
58	2Q	99.28	0.0	Ascomycota sp.	KP306958.1	Roussoella sp.
		99.11	0.0	Roussoella solani	LC195220.1	
		99.11	0.0	Roussoella solani	LC195219.1	
		99.11	0.0	Roussoella solani	LC195218.1	
		99.11	0.0	Pleosporales sp.	KP263105.1	
59	2V	98.79	0.0	Fusarium chlamydo- sporum	MF072670.1	Fusarium sp.
		98.42	0.0	Fusarium chlamydo- sporum	AY213655.1	
		98.42	0.0	Fusarium equiseti	HQ649909.1	
		98.41	0.0	Fusarium equiseti	HQ649905.1	
		98.23	0.0	Fusarium incarnatum	MH857950.1	
60	2T	97.18	0.0	Colletotrichum siamense	KP703406.1	Colletotrichum sp.
		97.18	0.0	Colletotrichum siamense	KP703404.1	
		97.18	0.0	Colletotrichum siamense	KP703403.1	
		97.18	0.0	Colletotrichum siamense	KP703402.1	
		97.18	0.0	Colletotrichum siamense	KP703385.1	
61	2AI	98.64	0.0	Eutypella sp.	KY083415.1	Eutypella sp.
		98.63	0.0	Eutypella scoparia	JQ922162.1	
		98.60	0.0	Eutypella sp.	JQ411308.1	
		98.44	0.0	Peroneutypa scoparia	MG873478.1	
		98.44	0.0	Eutypella sp.	GQ293963.1	

62	2AF	98.65	0.0	Trichoderma asperellum	KC993073.1	Trichoderma sp.
		98.64	0.0	Trichoderma lieckfeldtia	DQ109529.1	
		98.64	0.0	Trichoderma asperellum	KX510127.1	
		98.65	0.0	Trichoderma asperellum	MH153605.1	
		98.65	0.0	Trichoderma sp.	MK044000.1	
63	1O	99.19	0.0	Aspergillus tabacinus	AM883155.1	Aspergillus sp.
		99.00	0.0	Aspergillus tabacinus	LN898739.1	
		99.00	0.0	Aspergillus tabacinus	LN898738.1	
		99.00	0.0	Aspergillus tabacinus	LN898737.1	
		99.00	0.0	Aspergillus tabacinus	FR733851.1	
64	1F	98.01	0.0	Hypocrea sp.	HQ657316.1	Trichoderma sp.
		98.01	0.0	Trichoderma hamatum	MH864073.1	
		98.01	0.0	Trichoderma inhamatum	MH857107.1	
		98.01	0.0	Trichoderma harzianum	KT278882.1	
		98.01	0.0	Trichoderma harzianum	KR868358.1	
65	2D	99.38	00.0	Penicillium sp.	KX148648.1	Penicillium sp.
		99.17	0.0	Penicillium sp.	GU017521.1	
		99.17	0.0	Penicillium sp.	GU017519.1	
		99.17	0.0	Penicillium griseofulvum	FJ717687.1	
		99.17	0.0	Penicillium citrinum	MH864240	
66	2AG	99.16	0.0	Lasiodiplodia goni-biensis	KM006443.1	Lasiodiplodia sp.
		99.16	0.0	Lasiodiplodia goni-biensis	KF766191.1	
		99.16	0.0	Lasiodiplodia goni-biensis	NR_111218.1	
		99.13	0.0	Lasiodiplodia goni-biensis	AY639594.1	
		98.95	0.0	Lasiodiplodia goni-biensis	MH057180.1	
67	1BJ	99.08	0.0	Trichoderma spirale	KT278877.1	Trichoderma sp.
		99.08	0.0	Trichoderma sp.	LC229678.1	
		98.90	0.0	Trichoderma spirale	KT278886.1	
		98.90	0.0	Trichoderma spirale	KR093830.1	
		98.90	0.0	Trichoderma spirale	EU280082.1	
68	2K	98.85	0.0	Pestalotiopsis microspora	KT459350.1	Pestalotiopsis sp.
		98.85	0.0	Pestalotiopsis microspora	DQ456865.1	
		98.85	0.0	Pestalotiopsis microspora	AY924279.1	
		98.85	0.0	Pestalotiopsis sp.	HQ909077.1	
		98.62	0.0	Pestalotiopsis microspora	AY924271.1	

69	1P	100.00	0.0	Phoma sp.	JQ649376.1	Phoma sp.
		100.00	0.0	Phoma sp.	KP306990.1	
		100.00	0.0	Phoma sp.	KP306982.1	
		100.00	0.0	Phoma sp.	MN435151.1	
		100.00	0.0	Phoma sp.	KP306984.1	
70	1BD	96.79	0.0	Xenomyrothecium tongaense	MH861297.1	Myrothecium sp.
		96.43	0.0	Myrothecium inundatum	KU856653.1	
		96.22	0.0	Myrothecium inundatum	LC387258.1	
		96.22	0.0	Myrothecium inundatum	LC387257.1	
		96.22	0.0	Myrothecium inundatum	LC387254.1	
71	1S	97.39	0.0	Ramophialophora globispora	KU746700.1	NI
		96.91	0.0	Ramophialophora globispora	KU746699.1	
		95.23	0.0	Sordariomycetes sp.	KR909162.1	
72	1BM	99.60	0.0	Penicillium sp.	GU017521.1	Penicillium sp.
		99.60	0.0	Penicillium griseofulvum	FJ717687.1	
		99.60	0.0	Penicillium citrinum	MH864240.1	
		99.60	0.0	Penicillium citrinum	MH858380.1	
		99.60	0.0	Penicillium citrinum	MH856132.1	
73	1BE	98.05	0.0	Diaporthe sp.	MG976437.1	Diaporthe sp.
		97.39	0.0	Diaporthe sp.	MG976357.1	
		97.05	0.0	Diaporthe conorum	DQ116552.1	
		96.59	0.0	Diaporthe citri	KC343053.1	
		96.59	0.0	Diaporthe sp.	MG976377.1	
74	1BK	99.58	0.0	Fusarium incarnatum	KU382560.1	Fusarium sp.
		99.58	0.0	Fusarium sp.	HQ023205.1	
		98.97	0.0	Fusarium oxysporum	JN400714.1	
75	1N	98.99	0.0	Sarocladium glaucum	AB540577.1	Sarocladium sp.
		98.57	0.0	Sarocladium glaucum	NR_130686.1	
		98.39	0.0	Sarocladium glaucum	MH860868.1	
		98.38	0.0	Sarocladium glaucum	HG965020.1	
		98.09	0.0	Sarocladium glaucum	HG965019.1	
76	2AD	97.09	0.0	Fusarium sp.	HQ023200.1	Fusarium sp.
		96.42	0.0	Fusarium robustum	NR_159851.1	
		96.42	0.0	Fusarium robustum	MH861018.1	
		96.20	0.0	Fusarium robustum	U85539.1	
		95.75	0.0	Fusarium sp.	U85548.1	

77	2BC	99.79	0.0	Bionectria sp.	KF746125.1	NI
		99.57	0.0	Clonostachys pityrodes	JQ411387.1	
		99.17	0.0	Bionectria pityrodes	AF210673.1	
		98.75	0.0	Bionectria pityrodes	AF210672.1	
		98.63	0.0	Clonostachys pityrodes	MH864280.1	
78	1BA	99.15	0.0	Nigrospora pryzae	KJ159597.1	Nigrospora sp.
		99.14	0.0	Nigrospora sacchari-of-ficinarum	NR_165926.1	
		99.14	0.0	Nigrospora sp.	MN215791.1	
		98.85	0.0	Nigrospora sp.	HQ630982.1	
		98.74	0.0	Nigrospora sp.	JN207335.1	
79	1AD	99.63	0.0	Penicillium meleagrinum	HE962595.1	Penicillium sp.
		99.28	0.0	Penicillium meleagrinum	HM469412.1	
		99.28	0.0	Penicillium meleagrinum	EF198529.1	
		99.28	0.0	Penicillium meleagrinum	EF198530.1	
		99.28	0.0	Penicillium sp.	JF439496.1	
80	1D	92.90	9e-180	Mucor nidicola	JF299215.1	Mucor sp.
		92.86	0.0	Mucor sp.	FJ210509.1	
		90.02	0.0	Mucor sp.	FJ210517.1	
		89.95	0.0	Mucor nidicola	HQ913647.1	
81	2BE	98.41	0.0	Wiesneriomyces laurinus	KR822214.1	Wiesneriomyces sp.
		87.27	2e-111	Helicosporium gracile	AY916485.1	
		86.87	1e-113	Neofusiciccum mediterraneum	EU040221.1	

NI: Unable to get identification