

Supplemental file 1

Table S1. The results of molecular identification and GenBank accession numbers of DNA sequences included in this study.

Isolate	Molecular identification	GenBank accession number	Isolate	Molecular identification	GenBank accession number
	(IT'S gene)			(ITS ^a gene)	
SUB10203970 seq1	<i>Neoscytalidium dimidiatum</i>	MZ882261	SUB10203970 seq31	<i>Trichophyton mentagrophytes</i>	MZ882291
SUB10203970 seq2	<i>Candida albicans</i>	MZ882262	SUB10203970 seq32	<i>Trichophyton mentagrophytes</i>	MZ882292
SUB10203970 seq3	<i>Candida parapsilosis</i>	MZ882263	SUB10203970 seq33	<i>Trichophyton tonsurans</i>	MZ882293
SUB10203970 seq4	<i>Candida albicans</i>	MZ882264	SUB10203970 seq34	<i>Trichophyton rubrum</i>	MZ882294
SUB10203970 seq5	<i>Penicillium glabrum</i>	MZ882265	SUB10203970 seq35	<i>Trichophyton mentagrophytes</i>	MZ882295
SUB10203970 seq6	<i>Fusarium fujikuroi</i>	MZ882266	SUB10203970 seq36	<i>Trichophyton mentagrophytes</i>	MZ882296
SUB10203970 seq7	<i>Aspergillus flavus</i>	MZ882267	SUB10203970 seq37	<i>Trichophyton rubrum i</i>	MZ882297
SUB10203970 seq8	<i>Alternaria infectoria</i>	MZ882268	SUB10203970 seq38	<i>Candida albicans</i>	MK793223
SUB10203970 seq9	<i>Trichophyton tonsurans</i>	MZ882269	SUB10203970 seq39	<i>Candida tropicalis</i>	MK793225

SUB10203970 seq10	<i>Trichophyton rubrum</i>	MZ882270			SUB10203970 seq40	<i>Trichosporon asahii</i>	AB018013
SUB10203970 seq11	<i>Trichophyton rubrum</i>	MZ882271			SUB10203970 seq41	<i>Candida tropicalis</i>	MT772046
SUB10203970 seq12	<i>Aspergillus niger</i>	MZ882271			SUB10203970 seq42	<i>Candida glabrata</i>	MT772052
SUB10203970 seq13	<i>Trichophyton mentagrophytes</i>	MZ882273			SUB10203970 seq43	<i>Candida albicans</i>	MK138363
SUB10203970 seq14	<i>Penicillium polonicum</i>	MZ882274			SUB10203970 seq44	<i>Candida albicans</i>	MT772067
SUB10203970 seq15	<i>Aspergillus niger</i>	MZ882275			SUB10203970 seq45	<i>Candida parapsilosis</i>	EU564209
SUB10203970 seq16	<i>Candida albicans</i>	MZ882276			SUB10203970 seq46	<i>Fusarium oxysporum</i>	GQ922558
SUB10203970 seq17	<i>Aspergillus flavus</i>	MZ882277			SUB10203970 seq47	<i>Aspergillus flavus</i>	MK119732
SUB10203970 seq18	<i>Candida parapsilosis</i>	MZ882278			SUB10203970 seq48	<i>Aspergillus flavus</i>	MK119733
SUB10203970 seq19	<i>Cladosporium herbarum</i>	MZ882279			SUB10203970 seq49	<i>Aspergillus flavus</i>	MK119734
SUB10203970 seq20	<i>Candida parapsilosis</i>	MZ882280			SUB10203970 seq50	<i>Aspergillus niger</i>	LC387867
SUB10203970 seq21	<i>Aspergillus flavus</i>	MZ882281			SUB10203970 seq51	<i>Aspergillus niger</i>	LC387868
SUB10203970 seq22	<i>Candida parapsilosis</i>	MZ882282			SUB10203970 seq52	<i>Aspergillus terreus</i> <i>Trichophyton mentagrophytes</i>	GQ461911 AB520842
SUB10203970 seq23	<i>Rhodotorula mucilaginosa</i>	MZ882283			SUB10203970 seq53	<i>Trichophyton rubrum</i>	FJ746657
SUB10203970 seq24	<i>Candida tropicalis</i>	MZ882284			SUB10203970 seq54	<i>Trichophyton rubrum</i>	DW005357

SUB10203970 seq25	<i>Candida parapsilosis</i>	MZ882285			SUB10203970 seq55	<i>Trichophyton mentagrophytes</i>	HQ395066
SUB10203970 seq26	<i>Cladosporium pseudocladosporioides</i>	MZ882286			SUB10203970 seq56	<i>Trichophyton verrucosum</i>	AF168126
SUB10203970 seq27	<i>Trichophyton mentagrophytes</i>	MZ882287			SUB10203970 seq57	<i>Trichophyton rubrum</i>	DW005363
SUB10203970 seq28	<i>Penicillium glabrum</i>	MZ882288			SUB10203970 seq58	<i>Trichophyton rubrum</i>	DW005366
SUB10203970 seq29	<i>Rhodotorula mucilaginosa</i>	MZ882289			SUB10203970 seq59	<i>Trichophyton mentagrophytes</i>	MK045530
SUB10203970 seq30	<i>Candida albicans</i>	MZ882290			SUB10203970 seq60	<i>Trichophyton mentagrophytes</i>	MK045535

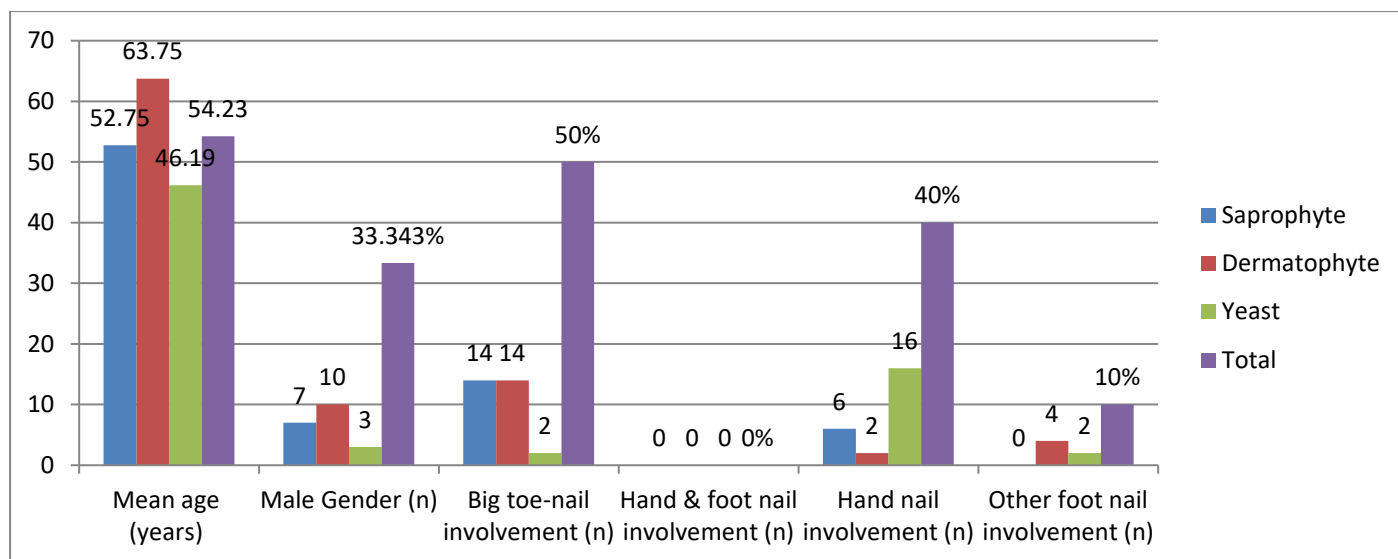
Table S2. GM of MIC concentrations for all tested isolates

GM(μ g/mL)	All	Yeast	Der	Sap
TER res	0.044	-	0.031	0.060
TER con	0.059	-	0.033	0.096
ITR res	0.56	0.435	0.529	0.756
ITR con	0.707	0.466	0.707	0.85
FLZ res	1.69	1.889	1.74	-
FLZ con	2.17	2.17	2.29	-
VOR res	0.158	0.153	0.096	0.229
VOR con	0.218	0.186	0.114	0.192
CAS res	0.567	0.567	-	-

CAS con	0.777	0.777	-	-
AMP B res	5.03	-	-	5.039
AMP B con	5.339	-	-	5.339

GM: Geometric mean; TER: terbinafine; ITR: itraconazole; FLZ: fluconazole; VOR: voriconazole; CAS: caspofungin; AMP B: amphotericin B

Figure S1. Demographic characteristics of the patients with onychomycosis included in the present study.



n: number