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*Chaperone*              *(HSPs)*              *Heat Shock Proteins)*              :

*HSP60*

*HSP60*              (*highly conserved*)  
*PCR*              (*template*)              *DNA*

*HSP60*

*McHSP60*

*C.immitis*    %              *HSP60*  
*HSP60*              *S.cerevisiae*    %      *Aspergillus fumigatus*    %  
*McHSP60*

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***HSP***              ***PCR***              :

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( )

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( )      **HSP**

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*ajahangirnejad@yahoo.com* :

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DNA .( )  
 EDTA (PH : ) Tris-HCl  
 K % - β % SDS (Chaperone) HSP  
 ( mg/ml) oC  
 × g .( )  
 RNase-H HSP  
 RNA ( mg/ml)  
 ATP  
 ( : : ) HSP .( )  
 DNA  
 × g .( )  
 %

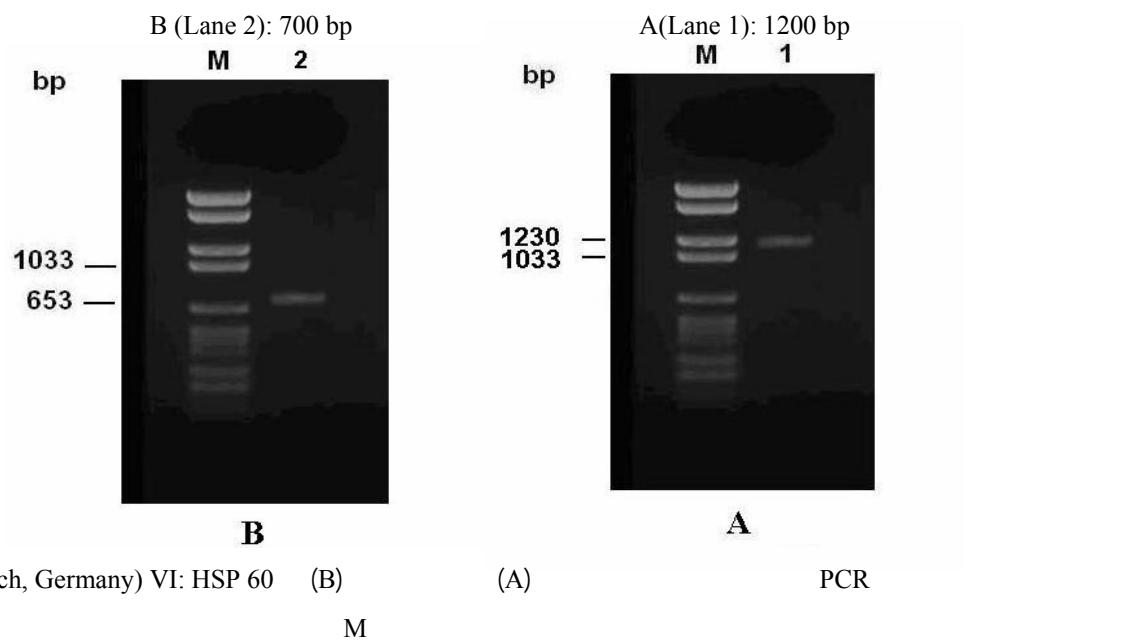
**PCR**

Gene Runner

MWG-Biotech

10X PCR Buffer .( )  
 DNA MgCl<sub>2</sub> dNTPmix •  
 / 40ng/μl DNA Choi

$^{\circ}\text{C}$  ( )  $^{\circ}\text{C}$  ( ) / (10pmol) 10ng/ $\mu\text{l}$  ( )  
 .( ) / Taq  
 PCR  
 . % ( ) PCR  
 .  $^{\circ}\text{C}$  ( )  $^{\circ}\text{C}$  : PCR  
 . PCR ( )  $^{\circ}\text{C}$  ( )  
 . Qiagen DNA ( )  
 Dye Terminator Cycle PCR :  
 . ( MWG ) 10X PCR Buffer  
 (NCBI - NIH) DNA MgCl<sub>2</sub> dNTPmix  
 . / 40ng/ $\mu\text{l}$   
 (10pmol) 10ng/ $\mu\text{l}$  ( )  
 Taq  
 PCR  
 (A) (B) : PCR  
 VI ( )  $^{\circ}\text{C}$  ( )  $^{\circ}\text{C}$  : PCR  
 . McHSP60  $^{\circ}\text{C}$  ( )  $^{\circ}\text{C}$  oC  
 . Roche



1	K	G	R	N	V	L	I	E	S	S	Y	G	S	P	K	I	T	T	K		18		
1	aag	gga	agg	aat	gtt	ttg	att	gag	tct	tca	tac	ggc	tcc	cca	aaa	att	act	aaa	g		55		
56	<b>gtatgccgtcaatttgcgcatactctacttaccgcggatagctaactccaaatata</b>																				114		
19	D	G	V	T	V	A	K	A	I	S	L	Q	D	K	F	E	N	L	G	A	38		
115	ac	gg	gt	c	ac	gg	gt	cc	aa	g	ct	tc	tt	ca	aa	g	aa	tt	g	at	cc	173	
39	R	L	L	Q	D	V	A	S	K	T	N	E	V	A	G	D	G	T	T	T	58		
174	cgt	ctt	ctc	caa	gac	gtt	gct	tcc	aag	aca	aac	gag	gtc	gcc	gg	gt	gac	gg	tt	ac	aca	233	
59	A	T	V	L	A	R	A	I	F	S	E	T	V	K	N	V	A	A	G	C	78		
234	gcg	acc	gt	ctt	gca	cgt	gct	atc	ttt	tcc	gag	acc	gtc	aag	aat	gtt	gct	gct	ggc	tgc	293		
79	N	P	M	D	L	R	R	G	I	Q	A	A	V	D	S	V	V	E	Y	L	98		
294	aa	cc	at	tg	gac	tt	aga	aga	gg	att	cag	gg	cc	tt	gac	tcc	gtc	gtc	gaa	ta	tt	353	
99	Q	A	N	K	R	E	I	T	T	S	E	E	I	A	Q	V	A	T	I	S	118		
354	caa	gca	aa	aag	aga	gag	atc	acc	acc	agc	gaa	gag	att	g	cg	ca	gt	gct	ac	atc	tct	413	
119	A	N	G	D	T	H	I	G	K	L	I	S	N	A	M	E	R	V	G	K	138		
414	gct	aa	gg	gac	ac	cat	atc	gg	aag	tt	atc	tcc	aa	gca	at	g	aa	ga	aa	gtt	gg	aa	473
139	E	G	V	I	T	V	K	D	G	K	T	I	E	D	E	L	E	V	T	E	158		
474	gaa	gg	gt	tt	ac	g	tt	aa	g	ac	cc	att	gaa	gac	g	ag	ctt	gag	tt	ac	gg	533	
159	G	M	R	F	D	R	G	Y	V	S	P	Y	F	I	T	D	P	K	T	Q	178		
534	gg	at	g	ca	ttt	gac	cgc	gg	tt	tcc	cct	tac	ttt	atc	ac	gac	ccc	aa	act	cag	593		
179	K	V	E	F	E	K	P	L	I	L	L	S	E	K	K	I	S	A	V	Q	198		
594	aa	g	tt	g	aa	g	aa	cct	tt	atc	ctc	ctc	tct	gag	aa	g	aa	atc	tct	gg	cag	563	
199	D	I	I	P	A	L	E	A	S	T	T	L	R	R	P	L	V	I	I	A	218		
654	gat	at	atc	cc	cc	gg	ctt	gag	gg	cc	tct	acc	cc	ctc	cg	ca	ct	gt	atc	tt	gct	713	
219	E	D	I	E	G	E	A	L	A	V	C	I	L	N	K	L	R	G	Q	L	238		
714	gag	gat	at	gag	gg	gag	gt	ctc	gca	gtc	tgc	att	ctc	aat	aa	ctg	cgt	gg	ca	ctt	773		
239	Q	V	A	A	V	K	A	P	G	F	G	D	N	R	K	S	I	L	G	D	258		
774	caa	gt	cc	gt	cc	aa	g	gt	cc	gg	tt	g	gt	aa	cc	cg	aa	ag	atc	ctt	gg	833	
259	I	A	V	L	T	N	G	T	V	F	T	D	E	L	D	M	K	L	D	K	278		
834	at	gg	cc	gt	tt	ac	aa	tt	gg	ac	cc	gt	tt	ac	aa	gt	tt	at	aa	gg	aa	893	
279	A	T	P	D	M	L	G	S	T	G	S	I	T	I	T	K	E	D	T	I	298		
894	g	o	c	cc	a	g	at	g	tc	gg	cc	tcc	ac	cc	tc	cc	aa	g	ac	act	tt	953	
299	I	L	N	G	E	G	S	K	D	A	I	A	Q	R	C	E	Q	I	S	G	318		
954	atc	ctg	aa	gg	gg	cc	tcc	aa	g	at	gg	cc	tt	g	ct	gg	ca	aa	tt	agc	gg	1013	
319	I	I	A	D	P	A	T	S	E	Y	E	K	E	K	L	Q	E	R	L	A	338		
1014	atc	at	g	ct	cc	gg	cc	aa	tcc	gaa	ta	c	g	aa	g	at	ctt	cag	gg	ct	gct	1073	
339	K	L	S	G	G	V	A	V	I	K	V	G	G	A	S	E	V	E	V	G	358		
1074	aa	aa	ct	tc	tt	gg	gt	gt	gt	gt	gt	cc	gg	gt	ct	t	gaa	gtt	gaa	gtt	gg	1133	
359	E	K	K	D	R	V	V	D	A	L	N	A	T	R	A	A	V	E	E	G	378		
1134	gag	aa	aa	g	ac	cgt	tt	gg	tt	gat	gg	cc	tc	ac	cc	cg	cc	gt	tt	gag	gg	1193	
379	I	L	P	G	G	T	A	L	L	K	A	S	A	N	G	L	K	D	V	398			
1194	at	ct	cc	gg	gg	gt	ac	gg	cc	tt	gg	ct	cc	aa	gt	tt	gg	aa	gac	gtc	1253		
399	K	P	A	N	F	D	Q	Q	L	G	V	S	I	V	K	N	A	I	Q	R	418		
1254	aag	cc	aa	cc	ttt	gac	cag	cag	ctg	gg	gt	cc	ac	cc	cc	gt	tt	gag	gg	gg	gt	1313	
419	P	A	R	T	I	V	E	N	A	G	L	E	G	S	V	I	V	G	K	L	438		
1314	cct	gt	cgt	act	ttt	gtt	gg	aa	t	gt	gg	tt	gg	gt	cc	at	gt	gg	cc	at	tt	1373	
439	T	D	E	F	A	D	D	F	N	R	G	F	D	S	A	K	G	E	Y	V	458		
1374	aca	gat	aa	ttt	gg	gac	gt	tt	aa	g	gg	cc	tt	gt	ac	gg	aa	gg	aa	tg	tt	1433	
459	D	M	I	Q	A	G	I	V	D	P	L	K	V	V	R	T	A	L	V	D	478		
1434	gat	at	g	c	ttt	gg	aa	ttt	gg	cc	aa	cc	cc	gt	cc	cc	gt	tt	gg	cc	at	1493	
479	A	S	G	V	A	S	L	L	G	T	T	E	V	A	I	V	E	A	P		497		
1494	gg	cc	gg	ttt	gg	cc	cc	tt	gg	cc	cc	gg	cc	gg	tt	gg	cc	aa	gt	cc		1550	

/ \_\_\_\_\_  
  
 McHSP60  
  
 .( )  
 HSP  
  
 Genbank  
 HSP  
 %  
 %  
 Coccidioides immitis  
 Aspergillus fumigatus  
  
 .( )  
 HSP  
 .( )  
 HSP60  
 DNA  
 HSP60  
 HSP60 .( ) HSP 60  
 Paracoccidioides brasiliensis (NCBI)  
 DQ981834  
 .( )  
 HSP60 Roska  
 T.mentagrophytes  
 (NCBI , NIH : AF199024 ) Epidermophyton Microsporum Trichophyton  
 McHSP60  
  
 M.canis  
  
 .( )  
 PCR  
 II DNA PCR-RFLP  
 % PCR .( )  
 Coccidioides immitis (Inter-single-sequence-repeat-PCR) ISSR-PCR  
 Aspergillus nidulans .( ) M.canis  
 HSP

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