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	diasis is the leading fungal blood	stream infection, and its in	ncidence has been on t	he rise. Recently, Ca	ndida parapsilosis has en	nerged as an increasingly
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nice. Mass spec f Candida albie	ns extracted from C.parapsilosis ctrometry analysis of the 32 imm cans, whereas Idh2p was identifi city was verified by immunoblot	unoreactive protein spots ed for the first time as an i	resulted in the identifi	cation of 12 distinct	proteins. Among them, 1	1 proteins were known antigen
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his work repre	esents the first immunoproteomic			insights into host-pa	thogen interactions and p	pathogenesis of C.parapsilosis.
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