

S5 Table. Top ten sequences similar to NCU05950*

NCBI/Gene bank sequence reference number	Protein function and name	Organism	E value
XP_009854653.1	Hypothetical protein NEUTE1DRAFT_69638	<i>Neurospora tetrasperma</i>	5e-114
XP_003349138.1	Hypothetical protein SMAC_06974	<i>Sordaria macrospora</i>	1e-103
XP_003650216.1	Hypothetical protein THITE_2063314	<i>Thielavia terrestris</i>	2e-60
KXX83062.1	Hypothetical protein MMYC01_200469	<i>Madurella mycetomatis</i>	6e-59
XP_001227334.1	Hypothetical protein CHGG_09407	<i>Chaetomium globosum</i>	8e-57
XP_006696736.1	Hypothetical protein CTHT_0064310	<i>Chaetomium thermophilum</i>	2e-56
XP_003663911.1	Hypothetical protein MYCTH_2315477	<i>Myceliophthora thermophile</i>	1e-55
XP_001906340.1	Hypothetical protein	<i>Podospora anserina</i>	3e-48
XP_008095402.1	Hypothetical protein GLRG_06526	<i>Colletotrichum graminicola</i>	4e-39
KZL63082.1	Hypothetical protein CI238_05869	<i>Colletotrichum incanum</i>	2e-38

*The NCU05950 protein sequence was used as the query for a protein BLAST search in the NCBI database of non-redundant coding sequences from GenBank.