

Additional file 1. *T. rubrum* EST database

GenBank Accession No.	Similarity		Organism	E-value
	Sequence ID	Gene product		
FE524602	-	No significant similarity	-	-
FE524603	-	No significant similarity	-	-
FE524604	-	No significant similarity	-	-
FE524605	-	No significant similarity	-	-
FE524606	ref XP_001241782.1	conserved hypothetical protein	<i>Coccidioides immitis</i> RS	6e-22
FE524607	-	No significant similarity	-	-
FE524608	-	No significant similarity	-	-
FE524609	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	2e-24
FE524610	emb CAK40436.1	unnamed protein product	<i>Aspergillus niger</i>	7e-08
FE524611	-	No significant similarity	-	-
FE524612	ref XP_001264707.1	S-adenosylmethionine synthetase	<i>Neosartorya fischeri</i> NRRL 181	8e-22
FE524613	-	No significant similarity	-	-
FE524614	-	No significant similarity	-	-
FE524615	-	No significant similarity	-	-
FE524616	-	No significant similarity	-	-
FE524617	-	No significant similarity	-	-
FE524618	-	No significant similarity	-	-
FE524619	-	No significant similarity	-	-
FE524620	sp P10255 CYPH_NEUCR	Peptidyl-prolyl cis-trans isomerase, mitochondrial precursor (PPIase) (Rotamase) (Cyclophilin) (Cyclosporin A-binding protein) (CPH) gb AAA33584.1 cyclophilin precursor emb CAA35681.1 cyclophilin (mitochondrial precursor)	<i>Neurospora crassa</i>	3e-17
FE524621	-	No significant similarity	-	-
FE524622	-	No significant similarity	-	-
FE524623	-	No significant similarity	-	-
FE524624	ref XP_001246797.1	polyubiquitin	<i>Coccidioides immitis</i> RS	3e-32
FE524625	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	1e-13
FE524626	-	No significant similarity	-	-
FE524627	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE524628	-	No significant similarity	-	-
FE524629	ref XP_001267336.1	outer mitochondrial membrane protein porin	<i>Neosartorya fischeri</i> NRRL 181	9e-18

FE524630	-	No significant similarity	-	-
FE524631	ref XP_001247117.1	60S ribosomal protein L44	<i>Coccidioides immitis</i> RS	2e-53
FE524632	gb AAO52807.1	hypothetical protein	<i>Bacillus megaterium</i>	1e-04
FE524633	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	1e-11
FE524634	-	No significant similarity	-	-
FE524635	-	No significant similarity	-	-
FE524636	-	No significant similarity	-	-
FE524637	dbj BAB12047.1	alpha-crystallin-related protein	<i>Arthroderma benhamiae</i>	1e-18
FE524638	-	No significant similarity	-	-
FE524639	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	2e-11
FE524640	-	No significant similarity	-	-
FE524641	-	No significant similarity	-	-
FE524642	-	No significant similarity	-	-
FE524643	ref XP_729762.1	senescence-associated protein	<i>Plasmodium yoelii yoelii</i> <i>str. 17XNL</i>	5e-06
FE524644	gb AAO52807.1	hypothetical protein	<i>Bacillus megaterium</i>	2e-04
FE524645	ref XP_001267438.1	hypothetical protein NFIA_043590	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-11
FE524646	ref XP_001240361.1	hypothetical protein CIMG_07524	<i>Coccidioides immitis</i> RS	3e-06
FE524647	-	No significant similarity	-	-
FE524648	-	No significant similarity	-	-
FE524649	-	No significant similarity	-	-
FE524650	-	No significant similarity	-	-
FE524651	-	No significant similarity	-	-
FE524652	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	3e-12
FE524653	-	No significant similarity	-	-
FE524654	-	No significant similarity	-	-
FE524655	-	No significant similarity	-	-
FE524656	-	No significant similarity	-	-
FE524657	-	No significant similarity	-	-
FE524658	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE524659	ref XP_001217109.1	methionyl-tRNA synthetase	<i>Aspergillus terreus</i> <i>NIH2624</i>	4e-04
FE524660	gb AAP05987.3	70 kDa heat shock protein	<i>Paracoccidioides</i> <i>brasiliensis</i>	8e-77
FE524661	-	No significant similarity	-	-
FE524662	sp P0C2C8 ATP7_ASPTN	ATP synthase D chain, mitochondrial precursor	<i>Aspergillus terreus</i> <i>NIH2624</i>	1e-55

FE524663	ref XP_001268300.1	chaperone/heat shock protein Hsp12, putative	<i>Aspergillus clavatus</i> NRRL 1	5e-20
FE524664	ref XP_001211604.1	predicted protein	<i>Aspergillus terreus</i> NIH2624	1e-05
FE524665	ref XP_001273409.1	dihydrolipoamide acetyltransferase component of pyruvate dehydrogenase	<i>Aspergillus clavatus</i> NRRL 1	1e-11
FE524666	-	No significant similarity	-	-
FE524667	ref XP_001273409.1	dihydrolipoamide acetyltransferase component of pyruvate dehydrogenase	<i>Aspergillus clavatus</i> NRRL 1	2e-05
FE524668	-	No significant similarity	-	-
FE524669	-	No significant similarity	-	-
FE524670	-	No significant similarity	-	-
FE524671	-	No significant similarity	-	-
FE524672	ref XP_001216088.1	splicing factor U2AF 23 kDa subunit	<i>Aspergillus terreus</i> NIH2624	4e-08
FE524673	-	No significant similarity	-	-
FE524674	-	No significant similarity	-	-
FE524675	-	No significant similarity	-	-
FE524676	sp P09188 PGK_PENCH	Phosphoglycerate kinase emb CAA31756.1 PGK protein	<i>Penicillium</i> <i>chrysogenum</i>	8e-15
FE524677	-	No significant similarity	-	-
FE524678	-	No significant similarity	-	-
FE524679	gb AAF04257.1	subtilisin-like serine protease	<i>Neospora caninum</i>	7e-05
FE524680	gb AAB05810.1	super cysteine rich protein; SCRP	<i>Homo sapiens</i>	4e-10
FE524681	-	No significant similarity	-	-
FE524682	dbj BAB12047.1	alpha-crystallin-related protein	<i>Arthroderma benhamiae</i>	3e-09
FE524683	-	No significant similarity	-	-
FE524684	ref XP_001211720.1	60S ribosomal protein L12	<i>Aspergillus terreus</i> NIH2624	6e-14
FE524685	-	No significant similarity	-	-
FE524686	-	No significant similarity	-	-
FE524687	-	No significant similarity	-	-
FE524688	-	No significant similarity	-	-
FE524689	gb EDJ96333.1	ubiquitin fusion protein	<i>Magnaporthe grisea</i> 70-15	7e-35
FE524690	-	No significant similarity	-	-
FE524691	-	No significant similarity	-	-
FE524692	-	No significant similarity	-	-

FE524693	-	No significant similarity	-	-
FE524694	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	3e-09
FE524695	ref XP_752703.1	Arp2/3 complex subunit (Arp2), putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	2e-69
FE524696	ref XP_001241781.1	conserved hypothetical protein	<i>Coccidioides immitis</i> RS	2e-19
FE524697	-	No significant similarity	-	-
FE524698	-	No significant similarity	-	-
FE524699	ref XP_001259633.1	ATP synthase F1, beta subunit, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	5e-38
FE524700	ref XP_746929.1	vacuolar protein sorting protein, putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	2e-12
FE524701	-	No significant similarity	-	-
FE524702	-	No significant similarity	-	-
FE524703	ref XP_001263333.1	proteasome subunit alpha type	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	5e-26
FE524704	ref XP_001222064.1	40S ribosomal protein S13	<i>Chaetomium globosum</i> <i>CBS 148.51</i>	7e-52
FE524705	-	No significant similarity	-	-
FE524706	-	No significant similarity	-	-
FE524707	ref XP_001244930.1	predicted protein	<i>Coccidioides immitis</i> RS	1e-08
FE524708	-	No significant similarity	-	-
FE524709	-	No significant similarity	-	-
FE524710	-	No significant similarity	-	-
FE524711	-	No significant similarity	-	-
FE524712	ref XP_755632.1	cytochrome c oxidase polypeptide vib	<i>Aspergillus fumigatus</i> <i>Af293</i>	3e-31
FE524713	-	No significant similarity	-	-
FE524714	gb EAL88345.2	60S ribosomal protein L27e	<i>Aspergillus fumigatus</i> <i>Af293</i>	1e-18
FE524715	-	No significant similarity	-	-
FE524716	-	No significant similarity	-	-
FE524717	-	No significant similarity	-	-
FE524718	-	No significant similarity	-	-
FE524719	-	No significant similarity	-	-
FE524720	-	No significant similarity	-	-
FE524721	-	No significant similarity	-	-
FE524722	-	No significant similarity	-	-
FE524723	pir B24264	proline-rich protein MP3 - mouse (fragment)	<i>Mus musculus</i>	1e-05

FE524724	-	No significant similarity	-	-
FE524725	-	No significant similarity	-	-
FE524726	-	No significant similarity	-	-
FE524727	ref XP_001267666.1	hypothetical protein NFIA_061330	<i>Neosartorya fischeri</i> NRRL 181	3e-19
FE524728	ref XP_001245526.1	hypothetical protein CIMG_04967	<i>Coccidioides immitis</i> RS	3e-28
FE524729	-	No significant similarity	-	-
FE524730	-	No significant similarity	-	-
FE524731	-	No significant similarity	-	-
FE524732	-	No significant similarity	-	-
FE524733	-	No significant similarity	-	-
FE524734	gb AAV23172.1	cytochrome c oxidase polypeptide VIa	<i>Penicillium</i> <i>chrysogenum</i>	3e-08
FE524735	-	No significant similarity	-	-
FE524736	-	No significant similarity	-	-
FE524737	-	No significant similarity	-	-
FE524738	-	No significant similarity	-	-
FE524739	gb ABI52743.1	10 kDa putative secreted protein	<i>Argas monolakensis</i>	2e-08
FE524740	-	No significant similarity	-	-
FE524741	ref XP_001248928.1	elongation factor 1-gamma	<i>Coccidioides immitis</i> RS	8e-32
FE524742	-	No significant similarity	-	-
FE524743	-	No significant similarity	-	-
FE524744	emb CAL46260.1	putative mitochondrial inner membrane protein 1	<i>Botryotinia fuckeliana</i>	6e-13
FE524745	ref XP_001213975.1	protein kinase dsk1	<i>Aspergillus terreus</i> NIH2624	3e-17
FE524746	sp Q9P8I0 PUT2_EMENI	Delta-1-pyrroline-5-carboxylate dehydrogenase, mitochondrial precursor (P5C dehydrogenase) gb AAF72527.1 AF252630_1 delta-1-pyrroline-5-carboxylate dehydrogenase	<i>Aspergillus nidulans</i>	8e-17
FE524747	-	No significant similarity	-	-
FE524748	-	No significant similarity	-	-
FE524749	-	No significant similarity	-	-
FE524750	-	No significant similarity	-	-
FE524751	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	6e-09
FE524752	dbj BAE64245.1	unnamed protein product	<i>Aspergillus oryzae</i>	5e-22
FE524753	-	No significant similarity	-	-
FE524754	dbj BAE65103.1	unnamed protein product	<i>Aspergillus oryzae</i>	2e-07
FE524755	-	No significant similarity	-	-
FE524756	gb AAW69337.1	ERV2 protein-like protein	<i>Magnaporthe grisea</i>	4e-17

FE524757	-	No significant similarity	-	-
FE524758	-	No significant similarity	-	-
FE524759	-	No significant similarity	-	-
FE524760	-	No significant similarity	-	-
FE524761	-	No significant similarity	-	-
FE524762	-	No significant similarity	-	-
FE524763	-	No significant similarity	-	-
FE524764	ref XP_001245839.1	arginyl-tRNA synthetase	<i>Coccidioides immitis</i> RS	1e-53
FE524765	-	No significant similarity	-	-
FE524766	-	No significant similarity	-	-
FE524767	gb AAR24348.1	14-3-3-like protein 2	<i>Paracoccidioides brasiliensis</i>	3e-81
FE524768	-	No significant similarity	-	-
FE524769	ref XP_001246797.1	polyubiquitin	<i>Coccidioides immitis</i> RS	8e-93
FE524770	ref XP_001267666.1	hypothetical protein NFIA_061330	<i>Neosartorya fischeri</i> NRRL 181	3e-07
FE524771	ref XP_001241781.1	conserved hypothetical protein	<i>Coccidioides immitis</i> RS	8e-17
FE524772	ref XP_001271667.1	60S ribosomal protein L11	<i>Aspergillus clavatus</i> NRRL 1	3e-63
FE524773	-	No significant similarity	-	-
FE524774	-	No significant similarity	-	-
FE524775	-	No significant similarity	-	-
FE524776	-	No significant similarity	-	-
FE524777	-	No significant similarity	-	-
FE524778	-	No significant similarity	-	-
FE524779	-	No significant similarity	-	-
FE524780	-	No significant similarity	-	-
FE524781	-	No significant similarity	-	-
FE524782	ref XP_001263817.1	UBX domain protein	<i>Neosartorya fischeri</i> NRRL 181	1e-21
FE524783	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE524784	-	No significant similarity	-	-
FE524785	-	No significant similarity	-	-
FE524786	-	No significant similarity	-	-
FE524787	-	No significant similarity	-	-
FE524788	-	No significant similarity	-	-
FE524789	-	No significant similarity	-	-
FE524790	gb AAX27763.1	SJCHGC01957 protein	<i>Schistosoma japonicum</i>	4e-16

FE524791	-	No significant similarity	-	-
FE524792	ref XP_001028186.1	hypothetical protein TTHERM_02641280	<i>Tetrahymena thermophila</i> SB210	1e-11
FE524793	ref XP_001242478.1	hypothetical protein CIMG_06374	<i>Coccidioides immitis</i> RS	3e-06
FE524794	dbj BAD90390.1	mFLJ00348 protein	<i>Mus musculus</i>	4e-04
FE524795	-	No significant similarity	-	-
FE524796	-	No significant similarity	-	-
FE524797	gb AAL76232.1 AF408429_1	metallothionein	<i>Microsporum canis</i>	2e-04
FE524798	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	2e-11
FE524799	-	No significant similarity	-	-
FE524800	-	No significant similarity	-	-
FE524801	ref XP_001243274.1	hypothetical protein CIMG_07170	<i>Coccidioides immitis</i> RS	5e-39
FE524802	-	No significant similarity	-	-
FE524803	ref XP_001270912.1	proteasome regulatory particle subunit (RpnL), putative	<i>Aspergillus clavatus</i> NRRL 1	2e-61
FE524804	ref XP_001214565.1	succinate-semialdehyde dehydrogenase	<i>Aspergillus terreus</i> NIH2624	1e-05
FE524805	-	No significant similarity	-	-
FE524806	ref XP_750061.1	ribosomal protein L29/heparin/heparan sulfate interacting protein	<i>Aspergillus fumigatus</i> Af293	3e-20
FE524807	-	No significant similarity	-	-
FE524808	gb AAC13689.1	ubiquitin fusion protein	<i>Magnaporthe grisea</i>	3e-47
FE524809	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	2e-14
FE524810	-	No significant similarity	-	-
FE524811	ref XP_001271742.1	acyl-CoA dehydrogenase, putative	<i>Aspergillus clavatus</i> NRRL 1	6e-23
FE524812	-	No significant similarity	-	-
FE524813	-	No significant similarity	-	-
FE524814	ref XP_750122.1	UPF0136 domain protein	<i>Aspergillus fumigatus</i> Af293	8e-25
FE524815	-	No significant similarity	-	-
FE524816	-	No significant similarity	-	-
FE524817	-	No significant similarity	-	-
FE524818	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	2e-09
FE524819	-	No significant similarity	-	-
FE524820	-	No significant similarity	-	-
FE524821	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	1e-13
FE524822	-	No significant similarity	-	-

FE524823	-	No significant similarity	-	-
FE524824	ref XP_750146.1	Rhodanese domain protein	<i>Aspergillus fumigatus</i> <i>Af293</i>	5e-37
FE524825	-	No significant similarity	-	-
FE524826	-	No significant similarity	-	-
FE524827	-	No significant similarity	-	-
FE524828	-	No significant similarity	-	-
FE524829	-	No significant similarity	-	-
FE524830	ref XP_001271126.1	ubiquitin	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	7e-16
FE524831	-	No significant similarity	-	-
FE524832	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	2e-09
FE524833	ref XP_001247270.1	hypothetical protein CIMG_01041	<i>Coccidioides immitis</i> RS	3e-12
FE524834	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE524835	-	No significant similarity	-	-
FE524836	-	No significant similarity	-	-
FE524837	-	No significant similarity	-	-
FE524838	ref XP_001241990.1	tubulin gamma chain	<i>Coccidioides immitis</i> RS	1e-95
FE524839	-	No significant similarity	-	-
FE524840	-	No significant similarity	-	-
FE524841	ref XP_001210407.1	elongation factor 1-beta	<i>Aspergillus terreus</i> <i>NIH2624</i>	4e-07
FE524842	-	No significant similarity	-	-
FE524843	-	No significant similarity	-	-
FE524844	-	No significant similarity	-	-
FE524845	ref XP_001275472.1	60S ribosomal protein L3	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	8e-25
FE524846	-	No significant similarity	-	-
FE524847	-	No significant similarity	-	-
FE524848	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE524849	sp P39457 PLB1_PENCH	Lysophospholipase precursor (Phospholipase B) emb CAA42906.1 lysophospholipase	<i>Penicillium</i> <i>chrysogenum</i>	4e-04
FE524850	ref XP_747457.1	DUF895 domain membrane protein	<i>Aspergillus fumigatus</i> <i>Af293</i>	3e-25
FE524851	-	No significant similarity	-	-
FE524852	-	No significant similarity	-	-
FE524853	-	No significant similarity	-	-
FE524854	-	No significant similarity	-	-

FE524855	ref XP_001212274.1	40S ribosomal protein S19	<i>Aspergillus terreus</i> <i>NIH2624</i>	1e-48
FE524856	-	No significant similarity	-	-
FE524857	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	2e-13
FE524858	-	No significant similarity	-	-
FE524859	-	No significant similarity	-	-
FE524860	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	2e-13
FE524861	-	No significant similarity	-	-
FE524862	gb AAM54368.1	elongation factor 1-alpha	<i>Trichophyton rubrum</i>	3e-47
FE524863	ref XP_001247280.1	alternative oxidase, mitochondrial precursor	<i>Coccidioides immitis</i> RS	4e-11
FE524864	-	No significant similarity	-	-
FE524865	-	No significant similarity	-	-
FE524866	-	No significant similarity	-	-
FE524867	-	No significant similarity	-	-
FE524868	-	No significant similarity	-	-
FE524869	-	No significant similarity	-	-
FE524870	-	No significant similarity	-	-
FE524871	-	No significant similarity	-	-
FE524872	-	No significant similarity	-	-
FE524873	-	No significant similarity	-	-
FE524874	-	No significant similarity	-	-
FE524875	gb AAO52807.1	hypothetical protein	<i>Bacillus megaterium</i>	1e-04
FE524876	ref XP_001274544.1	N-acetylglucosaminyl-phosphatidylinositol deacetylase, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-20
FE524877	ref XP_001129064.1	PREDICTED: hypothetical protein	<i>Homo sapiens</i>	8e-04
FE524878	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	5e-13
FE524879	-	No significant similarity	-	-
FE524880	ref XP_828313.1	hypothetical protein Tb11.18.0001	<i>Trypanosoma brucei</i> <i>TREU927</i>	9e-21
FE524881	ref XP_385039.1	hypothetical protein FG04863.1	<i>Gibberella zeae</i> PH-1	1e-06
FE524882	-	No significant similarity	-	-
FE524883	gb ABB20530.1	3-hydroxyphenylacetate 6 hydroxylase	<i>Emericella nidulans</i>	6e-12
FE524884	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE524885	-	No significant similarity	-	-
FE524886	-	No significant similarity	-	-
FE524887	emb CAE72981.1	Hypothetical protein CBG20323	<i>Caenorhabditis briggsae</i>	5e-16
FE524888	-	No significant similarity	-	-
FE524889	-	No significant similarity	-	-

FE524890	ref XP_001257424.1	ubiquitin C-terminal hydrolase, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-08
FE524891	ref XP_001275443.1	galactose-1-phosphate uridylyltransferase	<i>Aspergillus clavatus</i> NRRL 1	2e-34
FE524892	-	No significant similarity	-	-
FE524893	-	No significant similarity	-	-
FE524894	-	No significant similarity	-	-
FE524895	gb EU076569.1	Dipeptidyl-peptidase 5 (DPP5)	<i>Trichophyton equinum</i>	2e-35
FE524896	-	No significant similarity	-	-
FE524897	ref XP_749924.1	t-complex protein 1, beta subunit, putative	<i>Aspergillus fumigatus</i> Af293	6e-52
FE524898	-	No significant similarity	-	-
FE524899	ref XP_001358279.1	GA19372-PA	<i>Drosophila</i> <i>pseudoobscura</i>	3e-10
FE524900	-	No significant similarity	-	-
FE524901	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	4e-14
FE524902	ref XP_001267666.1	hypothetical protein NFIA_061330	<i>Neosartorya fischeri</i> NRRL 181	9e-31
FE524903	-	No significant similarity	-	-
FE524904	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	2e-09
FE524905	-	No significant similarity	-	-
FE524906	ref XP_001247816.1	superoxide dismutase	<i>Coccidioides immitis</i> RS	1e-08
FE524907	-	No significant similarity	-	-
FE524908	-	No significant similarity	-	-
FE524909	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	4e-08
FE524910	-	No significant similarity	-	-
FE524911	ref XP_001264967.1	30S ribosomal subunit S4, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-28
FE524912	pir T02955	probable cytochrome P450 monooxygenase - maize (fragment	<i>probable cytochrome</i> <i>P450 monooxygenas</i>	5e-06
FE524913	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	2e-13
FE524914	-	No significant similarity	-	-
FE524915	gb ABH10636.1	elongation factor 2	<i>Coccidioides posadasii</i>	1e-33
FE524916	-	No significant similarity	-	-
FE524917	-	No significant similarity	-	-
FE524918	sp P14728 YAV2_XANCV	Hypothetical 82 kDa avirulence protein in avrBs3 region	<i>Xanthomonas</i> <i>euvesicatoria</i>	6e-10
FE524919	ref XP_001268551.1	conidial hydrophobin RodB	<i>Aspergillus clavatus</i>	1e-04

FE524920	ref XP_001208527.1	cytochrome c oxidase assembly protein COX11, mitochondrial precursor	NRRL 1 <i>Aspergillus terreus</i> NIH2624	1e-26
FE524921	ref XP_001267676.1	hypothetical protein NFIA_043490	<i>Neosartorya fischeri</i> NRRL 181	4e-08
FE524922	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	3e-04
FE524923	-	No significant similarity	-	-
FE524924	-	No significant similarity	-	-
FE524925	-	No significant similarity	-	-
FE524926	gb AAX82553.1	120 kDa pistil extensin-like protein	<i>Nicotiana langsdorfii</i>	2e-04
FE524927	-	No significant similarity	-	-
FE524928	ref XP_571345.1	ketol-acid reductoisomerase	<i>Cryptococcus</i> <i>neoformans</i> var. <i>neoformans</i> JEC21	7e-13
FE524929	-	No significant similarity	-	-
FE524930	gb EAL88350.2	translation elongation factor EF-1 alpha subunit , putative	<i>Aspergillus fumigatus</i> Af293	2e-22
FE524931	ref XP_001243230.1	40S ribosomal protein S23	<i>Coccidioides immitis</i> RS	5e-19
FE524932	-	No significant similarity	-	-
FE524933	-	No significant similarity	-	-
FE524934	-	No significant similarity	-	-
FE524935	-	No significant similarity	-	-
FE524936	gb EDJ97592.1	sodium P-type ATPase, putative	<i>Magnaporthe grisea</i> 70-15	6e-04
FE524937	ref XP_001258284.1	polyketide synthase, putative	<i>Neosartorya fischeri</i> NRRL 181	5e-14
FE524938	ref XP_001244923.1	hypothetical protein CIMG_04364	<i>Coccidioides immitis</i> RS	2e-04
FE524939	-	No significant similarity	-	-
FE524940	gb EAW96235.1	hCG2043380	<i>Homo sapiens</i>	1e-06
FE524941	-	No significant similarity	-	-
FE524942	-	No significant similarity	-	-
FE524943	-	No significant similarity	-	-
FE524944	-	No significant similarity	-	-
FE524945	-	No significant similarity	-	-
FE524946	pir T02955	probable cytochrome P450 monooxygenase - maize (fragment)	<i>Zea mays</i>	6e-06
FE524947	-	No significant similarity	-	-
FE524948	-	No significant similarity	-	-
FE524949	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	2e-09

FE524950	gb EAL88703.2	adenylate cyclase AcyA	<i>Aspergillus fumigatus</i> <i>Af293</i>	3e-13
FE524951	ref XP_001264585.1	COPII-coated vesicle protein SurF4/Erv29, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-27
FE524952	-	No significant similarity	-	-
FE524953	gb AAV66614.1	putative secreted protein	<i>Ixodes scapularis</i>	5e-05
FE524954	pir T02955	probable cytochrome P450 monooxygenase - maize (fragment	<i>Zea mays</i>	6e-07
FE524955	ref XP_001267666.1	hypothetical protein NFIA_061330	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-17
FE524956	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE524957	ref XP_001257681.1	actin	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	3e-42
FE524958	-	No significant similarity	-	-
FE524959	sp P14728 YAV2_XANCV	Hypothetical 82 kDa avirulence protein in avrBs3 region	<i>Xanthomonas</i> <i>euvesicatoria</i>	6e-10
FE524960	ref XP_001248928.1	elongation factor 1-gamma	<i>Coccidioides immitis</i> RS	2e-04
FE524961	ref XP_001276223.1	WD domain protein	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-23
FE524962	-	No significant similarity	-	-
FE524963	-	No significant similarity	-	-
FE524964	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE524965	ref XP_001242010.1	60S ribosomal protein L5	<i>Coccidioides immitis</i> RS	2e-46
FE524966	dbj BAA10929.1	cytochrome P450 like_TBP	<i>Nicotiana tabacum</i>	1e-21
FE524967	-	No significant similarity	-	-
FE524968	-	No significant similarity	-	-
FE524969	gb AAF23950.1	NTR	<i>Cercopithecine</i> <i>herpesvirus 12</i>	1e-05
FE524970	-	No significant similarity	-	-
FE524971	-	No significant similarity	-	-
FE524972	-	No significant similarity	-	-
FE524973	-	No significant similarity	-	-
FE524974	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE524975	-	No significant similarity	-	-
FE524976	ref XP_001274835.1	60S ribosomal protein L7	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	3e-49
FE524977	-	No significant similarity	-	-
FE524978	ref XP_001274835.1	60S ribosomal protein L7	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-49

FE524979	-	No significant similarity	-	-
FE524980	ref XP_001212080.1	enolase	<i>Aspergillus terreus</i> NIH2624	7e-04
FE524981	ref XP_747833.1	60s ribosomal protein L24, putative	<i>Aspergillus fumigatus</i> Af293	2e-52
FE524982	ref XP_001271153.1	FHA domain protein SNIP1, putative	<i>Aspergillus clavatus</i> NRRL 1	1e-09
FE524983	-	No significant similarity	-	-
FE524984	-	No significant similarity	-	-
FE524985	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	2e-09
FE524986	ref XP_001265902.1	alkaline serine protease	<i>Neosartorya fischeri</i> NRRL 181	2e-12
FE524987	-	No significant similarity	-	-
FE524988	ref XP_001265902.1	alkaline serine protease	<i>Neosartorya fischeri</i> NRRL 181	2e-32
FE524989	-	No significant similarity	-	-
FE524990	-	No significant similarity	-	-
FE524991	-	No significant similarity	-	-
FE524992	ref XP_752433.1	Vacuolar ATP synthase subunit H, putative	<i>Aspergillus fumigatus</i> Af293	2e-39
FE524993	ref XP_001268628.1	40S ribosomal protein S25, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-25
FE524994	ref XP_001274344.1	1,4-alpha-glucan branching enzyme	<i>Aspergillus clavatus</i> NRRL 1	3e-19
FE524995	ref XP_001248308.1	60S ribosomal protein L28	<i>Coccidioides immitis</i> RS	1e-48
FE524996	-	No significant similarity	-	-
FE524997	ref XP_001248308.1	60S ribosomal protein L28	<i>Coccidioides immitis</i> RS	2e-44
FE524998	-	No significant similarity	-	-
FE524999	gb EEQ29198.1	glyceraldehyde-3-phosphate dehydrogenase	<i>Microsporum canis</i> CBS113480	8e-44
FE525000	ref XP_001267428.1	NADH-ubiquinone oxidoreductase 21 kDa subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	8e-41
FE525001	ref XP_001268300.1	chaperone/heat shock protein Hsp12, putative	<i>Aspergillus clavatus</i> NRRL 1	1e-23
FE525002	dbj BAA10929.1	cytochrome P450 like_TBP	<i>Nicotiana tabacum</i>	4e-25
FE525003	-	No significant similarity	-	-
FE525004	emb CAD22154.1	pherophorin-dz1 protein	<i>Volvox carteri</i> f. nagariensis	2e-13

FE525005	-	No significant similarity	-	-
FE525006	-	No significant similarity	-	-
FE525007	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	6e-12
FE525008	-	No significant similarity	-	-
FE525009	-	No significant similarity	-	-
FE525010	-	No significant similarity	-	-
FE525011	-	No significant similarity	-	-
FE525012	-	No significant similarity	-	-
FE525013	-	No significant similarity	-	-
FE525014	-	No significant similarity	-	-
FE525015	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	3e-09
FE525016	-	No significant similarity	-	-
FE525017	-	No significant similarity	-	-
FE525018	ref XP_001248361.1	histone H2A	<i>Coccidioides immitis</i> RS	6e-04
FE525019	ref XP_001254275.1	PREDICTED: hypothetical protein	<i>Bos taurus</i>	3e-15
FE525020	-	No significant similarity	-	-
FE525021	-	No significant similarity	-	-
FE525022	ref XP_001266185.1	nucleoside diphosphate kinase	<i>Neosartorya fischeri</i> NRRL 181	2e-52
FE525023	dbj BAD26579.1	cytochrome P450 like_TBP	<i>Citrullus lanatus</i>	2e-07
FE525024	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	3e-11
FE525025	-	No significant similarity	-	-
FE525026	-	No significant similarity	-	-
FE525027	gb AAC13689.1	ubiquitin fusion protein	<i>Magnaporthe grisea</i>	1e-38
FE525028	-	No significant similarity	-	-
FE525029	-	No significant similarity	-	-
FE525030	-	No significant similarity	-	-
FE525031	ref XP_001258304.1	benzoate 4-monooxygenase cytochrome P450	<i>Neosartorya fischeri</i> NRRL 181	3e-09
FE525032	ref XP_001259888.1	ABC metal ion transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-15
FE525033	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	4e-14
FE525034	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	3e-12
FE525035	ref XP_001261370.1	Ribosomal protein S28e	<i>Neosartorya fischeri</i> NRRL 181	9e-24
FE525036	-	No significant similarity	-	-
FE525037	ref XP_001257579.1	2-methylcitrate dehydratase, putative	<i>Neosartorya fischeri</i> NRRL 181	5e-42

FE525038	-	No significant similarity	-	-
FE525039	sp Q9C3Z6 RLA0_PODAN	60S acidic ribosomal protein P0 gb AAK11262.1 AF331714_1 ribosomal protein P0	<i>Podospora anserina</i>	1e-05
FE525040	gb ABH10639.1	fructose biphosphate aldolase	<i>Coccidioides posadasii</i>	1e-59
FE525041	-	No significant similarity	-	-
FE525042	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE525043	-	No significant similarity	-	-
FE525044	-	No significant similarity	-	-
FE525045	-	No significant similarity	-	-
FE525046	-	No significant similarity	-	-
FE525047	ref XP_001264877.1	heavy metal ion transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-10
FE525048	ref XP_001247280.1	alternative oxidase, mitochondrial precursor	<i>Coccidioides immitis</i> RS	1e-04
FE525049	-	No significant similarity	-	-
FE525050	ref XP_001244689.1	hypothetical protein CIMG_04130	<i>Coccidioides immitis</i> RS	8e-05
FE525051	-	No significant similarity	-	-
FE525052	-	No significant similarity	-	-
FE525053	ref NP_173553.1	ATEXT3 (EXTENSIN 3); structural constituent of cell wall	<i>Arabidopsis thaliana</i>	4e-05
FE525054	-	No significant similarity	-	-
FE525055	-	No significant similarity	-	-
FE525056	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	1e-13
FE525057	ref XP_001267676.1	hypothetical protein NFIA_043490	<i>Neosartorya fischeri</i> NRRL 181	2e-06
FE525058	-	No significant similarity	-	-
FE525059	ref XP_001265868.1	ribosomal protein L34 protein, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-07
FE525060	-	No significant similarity	-	-
FE525061	-	No significant similarity	-	-
FE525062	-	No significant similarity	-	-
FE525063	gb EEQ28799.1	alpha 1,6 mannosyltransferase	<i>Microsporum canis</i> CBS 113480	2e-18
FE525064	-	No significant similarity	-	-
FE525065	-	No significant similarity	-	-
FE525066	-	No significant similarity	-	-
FE525067	ref XP_001261287.1	tetratricopeptide repeat domain protein	<i>Neosartorya fischeri</i> NRRL 181	2e-05
FE525068	-	No significant similarity	-	-
FE525069	-	No significant similarity	-	-

FE525070	-	No significant similarity	-	-
FE525071	-	No significant similarity	-	-
FE525072	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	3e-12
FE525073	-	No significant similarity	-	-
FE525074	-	No significant similarity	-	-
FE525075	ref XP_001272337.1	AIF-like mitochondrial oxidoreductase (Nfrl), putative	<i>Aspergillus clavatus</i> NRRL 1	3e-17
FE525076	-	No significant similarity	-	-
FE525077	dbj BAB12232.1	fructose 1,6-bisphosphate aldolase	<i>Aspergillus oryzae</i>	2e-09
FE525078	ref XP_001275347.1	40S ribosomal protein S9	<i>Aspergillus clavatus</i> NRRL 1	8e-41
FE525079	ref XP_001268220.1	TBC domain protein, putative	<i>Aspergillus clavatus</i> NRRL 1	1e-12
FE525080	-	No significant similarity	-	-
FE525081	-	No significant similarity	-	-
FE525082	-	No significant similarity	-	-
FE525083	-	No significant similarity	-	-
FE525084	-	No significant similarity	-	-
FE525085	-	No significant similarity	-	-
FE525086	ref XP_001264756.1	adenosylhomocysteinase	<i>Neosartorya fischeri</i> NRRL 181	1e-60
FE525087	-	No significant similarity	-	-
FE525088	-	No significant similarity	-	-
FE525089	-	No significant similarity	-	-
FE525090	-	No significant similarity	-	-
FE525091	-	No significant similarity	-	-
FE525092	-	No significant similarity	-	-
FE525093	-	No significant similarity	-	-
FE525094	-	No significant similarity	-	-
FE525095	-	No significant similarity	-	-
FE525096	ref XP_001246644.1	protein translation factor SUI1	<i>Coccidioides immitis</i> RS	2e-41
FE525097	-	No significant similarity	-	-
FE525098	-	No significant similarity	-	-
FE525099	-	No significant similarity	-	-
FE525100	-	No significant similarity	-	-
FE525101	-	No significant similarity	-	-
FE525102	ref XP_001263624.1	eukaryotic translation initiation factor 4, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-25

FE525103	gb AAQ10290.1	class V chitin synthase	<i>Coccidioides posadasii</i>	5e-06
FE525104	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	8e-13
FE525105	-	No significant similarity	-	-
FE525106	ref XP_001268551.1	conidial hydrophobin RodB	<i>Aspergillus clavatus</i> NRRL 1	5e-05
FE525107	ref XP_001276223.1	WD domain protein	<i>Aspergillus clavatus</i> NRRL 1	8e-25
FE525108	-	No significant similarity	-	-
FE525109	-	No significant similarity	-	-
FE525110	ref XP_747995.1	nucleoside diphosphate kinase	<i>Aspergillus fumigatus</i> Af293	2e-32
FE525111	gb EAW96235.1	hCG2043380	<i>Homo sapiens</i>	1e-06
FE525112	-	No significant similarity	-	-
FE525113	-	No significant similarity	-	-
FE525114	ref XP_001243230.1	40S ribosomal protein S23	<i>Coccidioides immitis</i> RS	1e-19
FE525115	-	No significant similarity	-	-
FE525116	ref XP_001274770.1	60S ribosomal protein L13	<i>Aspergillus clavatus</i> NRRL 1	4e-12
FE525117	-	No significant similarity	-	-
FE525118	ref YP_293958.1	putative membrane protein	<i>Emiliania huxleyi</i> virus 86	8e-08
FE525119	ref XP_001240069.1	ATP-citrate synthase subunit 1	<i>Coccidioides immitis</i> RS	3e-28
FE525120	-	No significant similarity	-	-
FE525121	-	No significant similarity	-	-
FE525122	-	No significant similarity	-	-
FE525123	-	No significant similarity	-	-
FE525124	gb AAO52807.1	hypothetical protein	<i>Bacillus megaterium</i>	2e-04
FE525125	ref XP_001260256.1	replication factor A 1, rfa1	<i>Neosartorya fischeri</i> NRRL 181	1e-22
FE525126	-	No significant similarity	-	-
FE525127	ref XP_001271667.1	60S ribosomal protein L11	<i>Aspergillus clavatus</i> NRRL 1	3e-67
FE525128	ref XP_001246273.1	ADP-ribosylation factor 6	<i>Coccidioides immitis</i> RS	5e-21
FE525129	-	No significant similarity	-	-
FE525130	-	No significant similarity	-	-
FE525131	ref XP_001210992.1	homocitrate synthase, mitochondrial precursor	<i>Aspergillus terreus</i> NIH2624	1e-80
FE525132	-	No significant similarity	-	-

FE525133	-	No significant similarity	-	-
FE525134	gb AAL09828.1	beta-glucosidase 4	<i>Coccidioides immitis</i>	1e-16
FE525135	-	No significant similarity	-	-
FE525136	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	7e-12
FE525137	gb AAX33296.1	heat shock protein 90	<i>Paracoccidioides brasiliensis</i>	1e-31
FE525138	emb CAK47870.1	unnamed protein product	<i>Aspergillus niger</i>	3e-10
FE525139	-	No significant similarity	-	-
FE525140	gb EAT87923.1	hypothetical protein SNOG_04163	<i>Phaeosphaeria nodorum</i> SN15	3e-14
FE525141	-	No significant similarity	-	-
FE525142	-	No significant similarity	-	-
FE525143	ref XP_001276036.1	sulfatase domain protein	<i>Aspergillus clavatus</i> NRRL 1	9e-18
FE525144	gb ABA33785.1	RACK1-like protein	<i>Paracoccidioides brasiliensis</i>	2e-42
FE525145	-	No significant similarity	-	-
FE525146	-	No significant similarity	-	-
FE525147	-	No significant similarity	-	-
FE525148	-	No significant similarity	-	-
FE525149	-	No significant similarity	-	-
FE525150	-	No significant similarity	-	-
FE525151	ref XP_001269061.1	glycine cleavage system H protein	<i>Aspergillus clavatus</i> NRRL 1	5e-27
FE525152	emb CAM67156.1	Hypothetical repeat protein	<i>Leishmania infantum</i>	5e-06
FE525153	-	No significant similarity	-	-
FE525154	-	No significant similarity	-	-
FE525155	ref XP_729762.1	senescence-associated protein	<i>Plasmodium yoelii yoelii</i> str. 17XNL	4e-12
FE525156	-	No significant similarity	-	-
FE525157	-	No significant similarity	-	-
FE525158	-	No significant similarity	-	-
FE525159	-	No significant similarity	-	-
FE525160	-	No significant similarity	-	-
FE525161	-	No significant similarity	-	-
FE525162	-	No significant similarity	-	-
FE525163	-	No significant similarity	-	-
FE525164	-	No significant similarity	-	-

FE525165	ref XP_001214063.1	cytochrome c1, mitochondrial precursor	<i>Aspergillus terreus</i> NIH2624	8e-14
FE525166	ref XP_755078.1	fimbrin	<i>Aspergillus fumigatus</i> Af293	1e-13
FE525167	ref XP_001248022.1	threonyl-tRNA synthetase, cytoplasmic	<i>Coccidioides immitis</i> RS	7e-09
FE525168	-	No significant similarity	-	-
FE525169	-	No significant similarity	-	-
FE525170	-	No significant similarity	-	-
FE525171	-	No significant similarity	-	-
FE525172	ref XP_001269707.1	Ribosomal L18ae protein family	<i>Aspergillus clavatus</i> NRRL 1	5e-05
FE525173	-	No significant similarity	-	-
FE525174	-	No significant similarity	-	-
FE525175	gb AAK13589.1	rRNA intron-encoded homing endonuclease	<i>Oryza sativa</i>	3e-05
FE525176	-	No significant similarity	-	-
FE525177	-	No significant similarity	-	-
FE525178	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	1e-09
FE525179	gb EDJ94901.1	nucleoside diphosphate kinase	<i>Magnaporthe grisea</i> 70-15	3e-20
FE525180	ref XP_001272046.1	splicing factor 3a subunit 2, putative	<i>Aspergillus clavatus</i> NRRL 1	1e-10
FE525181	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-08
FE525182	-	No significant similarity	-	-
FE525183	-	No significant similarity	-	-
FE525184	-	No significant similarity	-	-
FE525185	ref XP_623110.1	PREDICTED: similar to CG3195-PA, isoform A	<i>Apis mellifera</i>	3e-21
FE525186	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	3e-12
FE525187	gb AAC78304.1	ubiquitin/ribosomal fusion protein	<i>Schistosoma japonicum</i>	3e-20
FE525188	ref XP_001243274.1	hypothetical protein CIMG_07170	<i>Coccidioides immitis</i> RS	1e-08
FE525189	-	No significant similarity	-	-
FE525190	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	7e-13
FE525191	ref XP_001239918.1	hypothetical protein CIMG_09539	<i>Coccidioides immitis</i> RS	8e-35
FE525192	gb ABA40770.1	SJCHGC01393 protein	<i>Schistosoma japonicum</i>	6e-04
FE525193	emb CAB51041.1	putative transcription factor	<i>Periplaneta americana</i>	4e-14
FE525194	-	No significant similarity	-	-
FE525195	-	No significant similarity	-	-
FE525196	-	No significant similarity	-	-
FE525197	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	5e-18

FE525198	dbj BAE65360.1	unnamed protein product	<i>Aspergillus oryzae</i>	4e-04
FE525199	-	No significant similarity	-	-
FE525200	-	No significant similarity	-	-
FE525201	ref XP_001257727.1	TOM complex component Tom7, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-13
FE525202	-	No significant similarity	-	-
FE525203	ref XP_001273715.1	acyl-CoA dehydrogenase, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-17
FE525204	-	No significant similarity	-	-
FE525205	ref XP_001269192.1	proteasome regulatory particle subunit (Nas6), putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-10
FE525206	ref XP_001265788.1	UDP-glucose:sterol glycosyltransferase	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	4e-05
FE525207	ref XP_965645.1	ATP synthase alpha chain, mitochondrial precursor	<i>Neurospora crassa</i> <i>OR74A</i>	4e-28
FE525208	ref XP_001274770.1	60S ribosomal protein L13	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-10
FE525209	-	No significant similarity	-	-
FE525210	sp P0C2C8 ATP7_ASPTN	ATP synthase D chain, mitochondrial precursor	<i>Aspergillus terreus</i> <i>NIH2624</i>	4e-11
FE525211	-	No significant similarity	-	-
FE525212	ref XP_001213339.1	glutathione peroxidase	<i>Aspergillus terreus</i> <i>NIH2624</i>	5e-12
FE525213	-	No significant similarity	-	-
FE525214	-	No significant similarity	-	-
FE525215	ref XP_001267676.1	hypothetical protein NFIA_043490	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	4e-08
FE525216	-	No significant similarity	-	-
FE525217	-	No significant similarity	-	-
FE525218	-	No significant similarity	-	-
FE525219	-	No significant similarity	-	-
FE525220	-	No significant similarity	-	-
FE525221	-	No significant similarity	-	-
FE525222	ref XP_568855.1	glucose transporter	<i>Cryptococcus</i> <i>neoformans</i> var. <i>neoformans JEC21</i>	2e-22
FE525223	-	No significant similarity	-	-
FE525224	-	No significant similarity	-	-

FE525225	-	No significant similarity	-	-
FE525226	-	No significant similarity	-	-
FE525227	ref XP_001214668.1	acetolactate synthase small subunit, mitochondrial precursor	<i>Aspergillus terreus</i> <i>NIH2624</i>	3e-05
FE525228	-	No significant similarity	-	-
FE525229	-	No significant similarity	-	-
FE525230	-	No significant similarity	-	-
FE525231	-	No significant similarity	-	-
FE525232	-	No significant similarity	-	-
FE525233	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	2e-09
FE525234	ref XP_001268551.1	conidial hydrophobin RodB	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-04
FE525235	-	No significant similarity	-	-
FE525236	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-27
FE525237	-	No significant similarity	-	-
FE525238	-	No significant similarity	-	-
FE525239	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	2e-13
FE525240	-	No significant similarity	-	-
FE525241	-	No significant similarity	-	-
FE525242	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	2e-09
FE525243	-	No significant similarity	-	-
FE525244	-	No significant similarity	-	-
FE525245	-	No significant similarity	-	-
FE525246	sp P56205 CYC_ASPNG	Cytochrome c	<i>Aspergillus niger</i>	1e-11
FE525247	gb EAL85176.2	glucosamine-6-phosphate deaminase, putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	6e-25
FE525248	-	No significant similarity	-	-
FE525249	tpd FAA00317.1	TPA: polyubiquitin	<i>Aspergillus nidulans</i> <i>FGSC A4</i>	1e-18
FE525250	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	2e-09
FE525251	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	1e-05
FE525252	ref XP_752345.1	stomatin family protein	<i>Aspergillus fumigatus</i> <i>Af293</i>	2e-06
FE525253	ref XP_001246620.1	40S ribosomal protein S4	<i>Coccidioides immitis</i> RS	3e-14
FE525254	ref XP_001264282.1	transcription factor TFIIA complex subunit Toa1, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-05
FE525255	ref XP_001266185.1	nucleoside diphosphate kinase	<i>Neosartorya fischeri</i>	2e-16

FE525256	-	No significant similarity	<i>NRRL 181</i>	-
FE525257	-	No significant similarity	-	-
FE525258	-	No significant similarity	-	-
FE525259	-	No significant similarity	-	-
FE525260	-	No significant similarity	-	-
FE525261	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	1e-08
FE525262	ref XP_001132700.1	PREDICTED: hypothetical protein	<i>Homo sapiens</i>	4e-17
FE525263	ref XP_001267666.1	hypothetical protein NFIA_061330	<i>Neosartorya fischeri</i>	6e-08
			<i>NRRL 181</i>	
FE525264	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	2e-08
FE525265	ref XP_001217121.1	60S ribosomal protein L9-B	<i>Aspergillus terreus</i>	2e-12
			<i>NIH2624</i>	
FE525266	-	No significant similarity	-	-
FE525267	-	No significant similarity	-	-
FE525268	-	No significant similarity	-	-
FE525269	-	No significant similarity	-	-
FE525270	-	No significant similarity	-	-
FE525271	-	No significant similarity	-	-
FE525272	ref XP_001266191.1	PHD transcription factor (Rum1), putative	<i>Neosartorya fischeri</i>	6e-17
			<i>NRRL 181</i>	
FE525273	-	No significant similarity	-	-
FE525274	ref XP_001267282.1	60S ribosomal protein L36	<i>Neosartorya fischeri</i>	7e-05
			<i>NRRL 181</i>	
FE525275	-	No significant similarity	-	-
FE525276	-	No significant similarity	-	-
FE525277	emb CAD56870.1	BclA protein	<i>Bacillus anthracis</i>	2e-06
FE525278	gb EAT89258.1	hypothetical protein SNOG_04053	<i>Phaeosphaeria nodorum</i>	2e-11
			<i>SN15</i>	
FE525279	-	No significant similarity	-	-
FE525280	-	No significant similarity	-	-
FE525281	-	No significant similarity	-	-
FE525282	-	No significant similarity	-	-
FE525283	-	No significant similarity	-	-
FE525284	-	No significant similarity	-	-
FE525285	gb ABH10645.1	ATP synthase beta chain	<i>Coccidioides posadasii</i>	2e-15
FE525286	-	No significant similarity	-	-
FE525287	-	No significant similarity	-	-

FE525288	-	No significant similarity	-	-
FE525289	-	No significant similarity	-	-
FE525290	-	No significant similarity	-	-
FE525291	-	No significant similarity	-	-
FE525292	-	No significant similarity	-	-
FE525293	-	No significant similarity	-	-
FE525294	-	No significant similarity	-	-
FE525295	-	No significant similarity	-	-
FE525296	ref XP_001246341.1	probable cation-transporting ATPase	<i>Coccidioides immitis</i> RS	6e-24
FE525297	-	No significant similarity	-	-
FE525298	-	No significant similarity	-	-
FE525299	-	No significant similarity	-	-
FE525300	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-10
FE525301	-	No significant similarity	-	-
FE525302	-	No significant similarity	-	-
FE525303	-	No significant similarity	-	-
FE525304	ref XP_001264792.1	aldehyde reductase (AKR1), putative	<i>Neosartorya fischeri</i> NRRL 181	1e-25
FE525305	-	No significant similarity	-	-
FE525306	-	No significant similarity	-	-
FE525307	-	No significant similarity	-	-
FE525308	-	No significant similarity	-	-
FE525309	-	No significant similarity	-	-
FE525310	-	No significant similarity	-	-
FE525311	-	No significant similarity	-	-
FE525312	-	No significant similarity	-	-
FE525313	ref XP_001261298.1	sugar transporter	<i>Neosartorya fischeri</i> NRRL 181	4e-24
FE525314	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	7e-15
FE525315	-	No significant similarity	-	-
FE525316	-	No significant similarity	-	-
FE525317	ref XP_001240403.1	proteasome component precursor	<i>Coccidioides immitis</i> RS	5e-17
FE525318	-	No significant similarity	-	-
FE525319	ref XP_001268304.1	BolA domain protein	<i>Aspergillus clavatus</i> NRRL 1	9e-15
FE525320	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	2e-07
FE525321	-	No significant similarity	-	-
FE525322	-	No significant similarity	-	-

FE525323	-	No significant similarity	-	-
FE525324	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE525325	ref XP_001209805.1	2-methylcitrate synthase, mitochondrial precursor	<i>Aspergillus terreus</i> NIH2624	2e-19
FE525326	gb EDK06386.1	40S ribosomal protein S3	<i>Magnaporthe grisea</i> 70-15	2e-17
FE525327	-	No significant similarity	-	-
FE525328	ref XP_001259371.1	chorismate mutase	<i>Neosartorya fischeri</i> NRRL 181	7e-08
FE525329	-	No significant similarity	-	-
FE525330	gb ABB18373.1	chitinase	<i>Coccidioides posadasii</i>	4e-14
FE525331	-	No significant similarity	-	-
FE525332	ref XP_001244490.1	ADP,ATP carrier protein	<i>Coccidioides immitis</i> RS	4e-22
FE525333	-	No significant similarity	-	-
FE525334	ref XP_001271628.1	5-methyltetrahydropteroyltriglutamate--homocysteine S-methyltransferase	<i>Aspergillus clavatus</i> NRRL 1	9e-05
FE525335	ref XP_001265922.1	IMP dehydrogenase, putative	<i>Neosartorya fischeri</i> NRRL 181	4e-26
FE525336	-	No significant similarity	-	-
FE525337	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	2e-09
FE525338	-	No significant similarity	-	-
FE525339	-	No significant similarity	-	-
FE525340	-	No significant similarity	-	-
FE525341	-	No significant similarity	-	-
FE525342	-	No significant similarity	-	-
FE525343	-	No significant similarity	-	-
FE525344	ref XP_001239885.1	60S ribosomal protein L32	<i>Coccidioides immitis</i> RS	1e-15
FE525345	ref XP_001151718.1	PREDICTED: hypothetical protein isoform 2	<i>Pan troglodytes</i>	5e-08
FE525346	-	No significant similarity	-	-
FE525347	-	No significant similarity	-	-
FE525348	gb ABI52743.1	10 kDa putative secreted protein	<i>Argas monolakensis</i>	3e-05
FE525349	-	No significant similarity	-	-
FE525350	-	No significant similarity	-	-
FE525351	sp Q9P4E9 GSP1_CANAL	GTP-binding nuclear protein GSP1/Ran gb AAF78478.1 AF190700_1 small G-protein Gsp1p	<i>Candida albicans</i>	7e-24
FE525352	gb EAT91215.1	predicted protein	<i>Phaeosphaeria nodorum</i> SN15	5e-09
FE525353	-	No significant similarity	-	-

FE525354	gb AAF23950.1	NTR	<i>Cercopithecine herpesvirus 12</i>	4e-09
FE525355	-	No significant similarity	-	-
FE525356	-	No significant similarity	-	-
FE525357	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE525358	-	No significant similarity	-	-
FE525359	ref XP_001214052.1	GTP-binding protein ypt3	<i>Aspergillus terreus</i> NIH2624	1e-05
FE525360	-	No significant similarity	-	-
FE525361	ref XP_001193920.1	PREDICTED: hypothetical protein, partial	<i>Strongylocentrotus purpuratus</i>	3e-05
FE525362	-	No significant similarity	-	-
FE525363	-	No significant similarity	-	-
FE525364	-	No significant similarity	-	-
FE525365	-	No significant similarity	-	-
FE525366	ref XP_751123.1	12 kDa heat shock protein	<i>Aspergillus fumigatus</i> Af293	5e-10
FE525367	ref XP_001241783.1	hypothetical protein CIMG_05679	<i>Coccidioides immitis</i> RS	5e-06
FE525368	ref XP_001243753.1	60S ribosomal protein L17	<i>Coccidioides immitis</i> RS	4e-24
FE525369	ref XP_001266324.1	UPF0047 domain protein	<i>Neosartorya fischeri</i> NRRL 181	2e-23
FE525370	-	No significant similarity	-	-
FE525371	dbj BAE73006.1	hypothetical protein	<i>Macaca fascicularis</i>	1e-06
FE525372	-	No significant similarity	-	-
FE525373	-	No significant similarity	-	-
FE525374	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	1e-12
FE525375	-	No significant similarity	-	-
FE525376	-	No significant similarity	-	-
FE525377	gb AAO73810.2	heat shock protein CLPA	<i>Paracoccidioides brasiliensis</i>	7e-06
FE525378	-	No significant similarity	-	-
FE525379	-	No significant similarity	-	-
FE525380	-	No significant similarity	-	-
FE525381	-	No significant similarity	-	-
FE525382	ref XP_001270964.1	extracellular thaumatin domain protein, putative	<i>Aspergillus clavatus</i> NRRL 1	1e-08
FE525383	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	1e-11
FE525384	-	No significant similarity	-	-

FE525385	ref XP_001246338.1	eukaryotic translation initiation factor 2 alpha subunit (eIF-2-alpha)	<i>Coccidioides immitis</i> RS	1e-18
FE525386	-	No significant similarity	-	-
FE525387	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	1e-04
FE525388	ref XP_001268377.1	60S ribosomal protein L31e	<i>Aspergillus clavatus</i> NRRL 1	4e-27
FE525389	-	No significant similarity	-	-
FE525390	-	No significant similarity	-	-
FE525391	-	No significant similarity	-	-
FE525392	-	No significant similarity	-	-
FE525393	ref XP_001259633.1	ATP synthase F1, beta subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	9e-22
FE525394	gb ABF13477.1	DRK1 histidine kinase	<i>Ajellomyces dermatitidis</i>	8e-19
FE525395	ref XP_755487.1	transmembrane domain-containing protein	<i>Aspergillus fumigatus</i> Af293	1e-11
FE525396	-	No significant similarity	-	-
FE525397	-	No significant similarity	-	-
FE525398	-	No significant similarity	-	-
FE525399	-	No significant similarity	-	-
FE525400	-	No significant similarity	-	-
FE525401	ref XP_001247329.1	tyrosyl-tRNA synthetase, cytoplasmic	<i>Coccidioides immitis</i> RS	9e-06
FE525402	ref XP_753079.1	long chain fatty alcohol oxidase, putative	<i>Aspergillus fumigatus</i> Af293	2e-07
FE525403	-	No significant similarity	-	-
FE525404	gb ABF13597.1	dermatan-binding protein PA5541	<i>Propionibacterium</i> <i>acnes</i>	7e-06
FE525405	ref XP_001209812.1	malate synthase	<i>Aspergillus terreus</i> NIH2624	8e-32
FE525406	-	No significant similarity	-	-
FE525407	gb EAL84560.2	protein kinase activator Bem1, putative	<i>Aspergillus fumigatus</i> Af293	6e-06
FE525408	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	3e-09
FE525409	ref XP_001212958.1	40S ribosomal protein S8-B	<i>Aspergillus terreus</i> NIH2624	7e-09
FE525410	gb AAM54368.1	elongation factor 1-alpha	<i>Trichophyton rubrum</i>	1e-46
FE525411	-	No significant similarity	-	-
FE525412	-	No significant similarity	-	-
FE525413	-	No significant similarity	-	-
FE525414	-	No significant similarity	-	-

FE525415	ref XP_001267676.1	hypothetical protein NFIA_043490	<i>Neosartorya fischeri</i> NRRL 181	1e-04
FE525416	-	No significant similarity	-	-
FE525417	-	No significant similarity	-	-
FE525418	-	No significant similarity	-	-
FE525419	-	No significant similarity	-	-
FE525420	-	No significant similarity	-	-
FE525421	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	2e-09
FE525422	ref XP_001267290.1	5-methyltetrahydropteroyltriglutamate--homocysteine S-methyltransferase	<i>Neosartorya fischeri</i> NRRL 181	6e-34
FE525423	ref XP_001241783.1	hypothetical protein CIMG_05679	<i>Coccidioides immitis</i> RS	3e-07
FE525424	-	No significant similarity	-	-
FE525425	ref XP_001271804.1	cytochrome c peroxidase Ccp1, putative	<i>Aspergillus clavatus</i> NRRL 1	4e-37
FE525426	ref XP_001266962.1	transcription regulator BDF1, putative	<i>Neosartorya fischeri</i> NRRL 181	4e-08
FE525427	ref YP_288338.1	Surface protein from Gram-positive cocci, anchor region	<i>Thermobifida fusca</i> YX	4e-04
FE525428	ref XP_568822.1	UDP-glucose:glycoprotein glucosyltransferase	<i>Cryptococcus</i> <i>neoformans</i> var. <i>neoformans</i> JEC21	9e-38
FE525429	-	No significant similarity	-	-
FE525430	-	No significant similarity	-	-
FE525431	-	No significant similarity	-	-
FE525432	-	No significant similarity	-	-
FE525433	-	No significant similarity	-	-
FE525434	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	2e-09
FE525435	-	No significant similarity	-	-
FE525436	-	No significant similarity	-	-
FE525437	-	No significant similarity	-	-
FE525438	ref XP_001215933.1	asparaginyl-tRNA synthetase	<i>Aspergillus terreus</i> NIH2624	2e-50
FE525439	-	No significant similarity	-	-
FE525440	-	No significant similarity	-	-
FE525441	-	No significant similarity	-	-
FE525442	ref XP_868852.1	uricase	<i>Aspergillus nidulans</i> FGSC A4	6e-35
FE525443	-	No significant similarity	-	-
FE525444	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	9e-07

FE525445	-	No significant similarity	-	-
FE525446	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	2e-09
FE525447	-	No significant similarity	-	-
FE525448	ref XP_001267336.1	outer mitochondrial membrane protein porin	<i>Neosartorya fischeri</i> NRRL 181	5e-11
FE525449	-	No significant similarity	-	-
FE525450	-	No significant similarity	-	-
FE525451	-	No significant similarity	-	-
FE525452	gb AAO52807.1	hypothetical protein	<i>Bacillus megaterium</i>	9e-05
FE525453	ref XP_001267336.1	outer mitochondrial membrane protein porin	<i>Neosartorya fischeri</i> NRRL 181	3e-09
FE525454	-	No significant similarity	-	-
FE525455	-	No significant similarity	-	-
FE525456	ref XP_001247752.1	hypothetical protein CIMG_01523	<i>Coccidioides immitis</i> RS	6e-20
FE525457	-	No significant similarity	-	-
FE525458	-	No significant similarity	-	-
FE525459	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	4e-13
FE525460	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	3e-12
FE525461	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	4e-23
FE525462	-	No significant similarity	-	-
FE525463	-	No significant similarity	-	-
FE525464	-	No significant similarity	-	-
FE525465	-	No significant similarity	-	-
FE525466	-	No significant similarity	-	-
FE525467	ref XP_001245555.1	eukaryotic translation initiation factor 5A-2	<i>Coccidioides immitis</i> RS	2e-05
FE525468	-	No significant similarity	-	-
FE525469	-	No significant similarity	-	-
FE525470	-	No significant similarity	-	-
FE525471	ref XP_001258207.1	short chain dehydrogenase/reductase family	<i>Neosartorya fischeri</i> NRRL 181	5e-05
FE525472	ref XP_001210407.1	elongation factor 1-beta	<i>Aspergillus terreus</i> NIH2624	7e-12
FE525473	-	No significant similarity	-	-
FE525474	-	No significant similarity	-	-
FE525475	-	No significant similarity	-	-
FE525476	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE525477	-	No significant similarity	-	-
FE525478	-	No significant similarity	-	-

FE525479	-	No significant similarity	-	-
FE525480	-	No significant similarity	-	-
FE525481	ref XP_001257958.1	ubiquinol cytochrome-c reductase subunit 9, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-17
FE525482	ref XP_001260275.1	serine/threonine protein kinase (Ark1), putative	<i>Neosartorya fischeri</i> NRRL 181	8e-35
FE525483	ref XP_001229506.1	60S ribosomal protein L29	<i>Chaetomium globosum</i> CBS 148.51	7e-28
FE525484	ref NP_941181.1	chloramphenicol acetyltransferase	<i>Serratia marcescens</i>	4e-05
FE525485	-	No significant similarity	-	-
FE525486	-	No significant similarity	-	-
FE525487	-	No significant similarity	-	-
FE525488	-	No significant similarity	-	-
FE525489	ref XP_001240451.1	aspartyl aminopeptidase, putative	<i>Coccidioides immitis</i> RS	1e-08
FE525490	ref XP_752593.1	mitochondrial hypoxia responsive domain protein	<i>Aspergillus fumigatus</i> Af293	3e-30
FE525491	ref XP_001264032.1	conidial hydrophobin RodB	<i>Neosartorya fischeri</i> NRRL 181	7e-07
FE525492	-	No significant similarity	-	-
FE525493	-	No significant similarity	-	-
FE525494	-	No significant similarity	-	-
FE525495	-	No significant similarity	-	-
FE525496	-	No significant similarity	-	-
FE525497	gb AAL31950.1	CDH1-D	<i>Gallus gallus</i>	3e-11
FE525498	-	No significant similarity	-	-
FE525499	ref XP_749092.1	integral peroxisomal membrane protein	<i>Aspergillus fumigatus</i> Af293	4e-24
FE525500	-	No significant similarity	-	-
FE525501	ref XP_001265155.1	COPII-coated vesicle membrane protein Erv46, putative	<i>Neosartorya fischeri</i> NRRL 181	9e-33
FE525502	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	2e-13
FE525503	ref XP_001241599.1	predicted protein	<i>Coccidioides immitis</i> RS	6e-11
FE525504	-	No significant similarity	-	-
FE525505	-	No significant similarity	-	-
FE525506	pir T02955	probable cytochrome P450 monooxygenase - maize (fragment)	<i>Zea mays</i>	6e-07
FE525507	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	5e-14
FE525508	-	No significant similarity	-	-
FE525509	ref XP_001269838.1	6-phosphogluconolactonase	<i>Aspergillus clavatus</i>	2e-34

FE525510	-	No significant similarity
FE525511	ref XP_001241788.1	hypothetical protein CIMG_05684
FE525512	ref XP_001267282.1	60S ribosomal protein L36
FE525513	ref XP_453838.1	unnamed protein product
FE525514	-	No significant similarity
FE525515	-	No significant similarity
FE525516	gb AAN63566.1 AF429823_1	septin 3
FE525517	-	No significant similarity
FE525518	-	No significant similarity
FE525519	-	No significant similarity
FE525520	-	No significant similarity
FE525521	gb AAO52807.1	hypothetical protein
FE525522	-	No significant similarity
FE525523	-	No significant similarity
FE525524	-	No significant similarity
FE525525	-	No significant similarity
FE525526	-	No significant similarity
FE525527	ref XP_660794.1	hypothetical protein AN3190.2
FE525528	-	No significant similarity
FE525529	-	No significant similarity
FE525530	-	No significant similarity
FE525531	-	No significant similarity
FE525532	-	No significant similarity
FE525533	-	No significant similarity
FE525534	-	No significant similarity
FE525535	-	No significant similarity
FE525536	ref XP_001242406.1	O-acetylhomoserine
FE525537	-	No significant similarity
FE525538	-	No significant similarity
FE525539	-	No significant similarity
FE525540	-	No significant similarity
FE525541	-	No significant similarity
FE525542	-	No significant similarity
FE525543	ref XP_001241788.1	hypothetical protein CIMG_05684
FE525544	ref XP_001241788.1	hypothetical protein CIMG_05684

NRRL 1

-	-
<i>Coccidioides immitis</i> RS	1e-12
<i>Neosartorya fischeri</i>	7e-26
NRRL 181	
<i>Kluyveromyces lactis</i>	5e-06
-	-
-	-
<i>Coccidioides immitis</i>	5e-08
-	-
-	-
-	-
-	-
<i>Bacillus megaterium</i>	2e-04
-	-
-	-
-	-
-	-
-	-
<i>Aspergillus nidulans</i>	7e-23
FGSC A4	
-	-
-	-
-	-
-	-
-	-
-	-
-	-
<i>Coccidioides immitis</i> RS	5e-21
-	-
-	-
-	-
-	-
-	-
-	-
<i>Coccidioides immitis</i> RS	5e-14
<i>Coccidioides immitis</i> RS	5e-14

FE525545	-	No significant similarity	-	-
FE525546	-	No significant similarity	-	-
FE525547	ref XP_001247210.1	NAD-specific glutamate dehydrogenase (NAD-GDH)	<i>Coccidioides immitis</i> RS	3e-34
FE525548	gb ABG74714.1	Ac1147-like protein	<i>Diaphorina citri</i>	4e-21
FE525549	ref XP_001241788.1	hypothetical protein CIMG_05684	<i>Coccidioides immitis</i> RS	8e-14
FE525550	-	No significant similarity	-	-
FE525551	-	No significant similarity	-	-
FE525552	-	No significant similarity	-	-
FE525553	ref XP_001266925.1	alpha-aminoadipate reductase large subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	4e-11
FE525554	-	No significant similarity	-	-
FE525555	ref XP_001241783.1	hypothetical protein CIMG_05679	<i>Coccidioides immitis</i> RS	3e-05
FE525556	emb CAM19749.1	novel KRAB box containing protein	<i>Mus musculus</i>	1e-08
FE525557	-	No significant similarity	-	-
FE525558	-	No significant similarity	-	-
FE525559	-	No significant similarity	-	-
FE525560	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	3e-09
FE525561	-	No significant similarity	-	-
FE525562	-	No significant similarity	-	-
FE525563	-	No significant similarity	-	-
FE525564	-	No significant similarity	-	-
FE525565	-	No significant similarity	-	-
FE525566	-	No significant similarity	-	-
FE525567	-	No significant similarity	-	-
FE525568	ref XP_001276134.1	vacuolar aspartyl aminopeptidase Lap4, putative	<i>Aspergillus clavatus</i> NRRL 1	1e-10
FE525569	gb AAB05810.1	super cysteine rich protein; SCRP	<i>Homo sapiens</i>	9e-15
FE525570	ref XP_001239110.1	hypothetical protein CIMG_10132	<i>Coccidioides immitis</i> RS	1e-16
FE525571	-	No significant similarity	-	-
FE525572	ref XP_001261472.1	40S ribosomal protein S13	<i>Neosartorya fischeri</i> NRRL 181	2e-10
FE525573	ref XP_001275347.1	40S ribosomal protein S9	<i>Aspergillus clavatus</i> NRRL 1	1e-11
FE525574	-	No significant similarity	-	-
FE525575	-	No significant similarity	-	-
FE525576	-	No significant similarity	-	-
FE525577	-	No significant similarity	-	-
FE525578	-	No significant similarity	-	-

FE525579	dbj BAE58435.1	unnamed protein product	<i>Aspergillus oryzae</i>	2e-06
FE525580	ref XP_001258253.1	ABC transporter, putative	<i>Neosartorya fischeri</i>	1e-21
			NRRL 181	
FE525581	ref XP_001248139.1	hypothetical protein CIMG_01910	<i>Coccidioides immitis</i> RS	2e-25
FE525582	-	No significant similarity	-	-
FE525583	ref XP_001246822.1	hypothetical protein CIMG_00593	<i>Coccidioides immitis</i> RS	3e-40
FE525584	-	No significant similarity	-	-
FE525585	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	3e-21
FE525586	ref XP_001258253.1	ABC transporter, putative	<i>Neosartorya fischeri</i>	3e-21
			NRRL 181	
FE525587	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	2e-07
FE525588	ref YP_840033.1	Glyoxalase/bleomycin resistance protein/dioxygenase	<i>Burkholderia</i>	3e-11
			<i>cenocepacia</i> HI2424	
FE525589	ref XP_001258253.1	ABC transporter, putative	<i>Neosartorya fischeri</i>	1e-21
			NRRL 181	
FE525590	-	No significant similarity	-	-
FE525591	-	No significant similarity	-	-
FE525592	ref YP_840033.1	Glyoxalase/bleomycin resistance protein/dioxygenase	<i>Burkholderia</i>	2e-07
			<i>cenocepacia</i> HI2424	
FE525593	gb EAL84327.2	short chain oxidoreductase (CsgA), putative	<i>Aspergillus fumigatus</i>	2e-04
			Af293	
FE525594	ref XP_001272698.1	Dopey, N-terminal domain protein	<i>Aspergillus clavatus</i>	2e-09
			NRRL 1	
FE525595	-	No significant similarity	-	-
FE525596	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	2e-43
FE525597	ref XP_001266096.1	Pumilio-family RNA binding repeat protein	<i>Neosartorya fischeri</i>	3e-10
			NRRL 181	
FE525598	-	No significant similarity	-	-
FE525599	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	5e-07
FE525600	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	3e-21
FE525601	-	No significant similarity	-	-
FE525602	ref YP_840033.1	Glyoxalase/bleomycin resistance protein/dioxygenase	<i>Burkholderia</i>	3e-11
			<i>cenocepacia</i> HI2424	
FE525603	-	No significant similarity	-	-
FE525604	-	No significant similarity	-	-
FE525605	gb EEQ28523.1	Salicylate hydroxylase (Salicylate 1-monooxygenase)	<i>Microsporum canis</i> CBS	6e-43
			113480	
FE525606	ref XP_001247286.1	hypothetical protein CIMG_01057	<i>Coccidioides immitis</i> RS	1e-05

FE525607	ref XP_001258253.1	ABC transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-21
FE525608	gb EAL84327.2	short chain oxidoreductase (CsgA), putative	<i>Aspergillus fumigatus</i> Af293	7e-04
FE525609	-	No significant similarity	-	-
FE525610	ref XP_001268233.1	ABC transporter, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-17
FE525611	ref XP_662910.1	hypothetical protein AN5306.2	<i>Aspergillus nidulans</i> FGSC A4	6e-11
FE525612	ref XP_001229436.1	hypothetical protein CHGG_02920	<i>Chaetomium globosum</i> CBS 148.51	3e-04
FE525613	ref XP_001258253.1	ABC transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-21
FE525614	-	No significant similarity	-	-
FE525615	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE525616	-	No significant similarity	-	-
FE525617	-	No significant similarity	-	-
FE525618	ref XP_750058.1	carboxylic ester hydrolase (Ppe1), putative	<i>Aspergillus fumigatus</i> Af293	2e-24
FE525619	ref YP_673529.1	Glyoxalase/bleomycin resistance protein/dioxygenase	<i>Mesorhizobium</i> sp. BNC1	4e-09
FE525620	-	No significant similarity	-	-
FE525621	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	5e-15
FE525622	ref XP_662910.1	hypothetical protein AN5306.2	<i>Aspergillus nidulans</i> FGSC A4	2e-10
FE525623	ref XP_001266096.1	Pumilio-family RNA binding repeat protein	<i>Neosartorya fischeri</i> NRRL 181	4e-09
FE525624	ref XP_001267191.1	DNA mismatch repair protein Msh6, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-33
FE525625	-	No significant similarity	-	-
FE525626	ref XP_367322.1	hypothetical protein MG07247.4	<i>Magnaporthe grisea</i> 70-15	2e-06
FE525627	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE525628	ref XP_001262368.1	DEAD helicases superfamily protein (Aquarius), putative	<i>Neosartorya fischeri</i> NRRL 181	1e-19
FE525629	-	No significant similarity	-	-
FE525630	-	No significant similarity	-	-
FE525631	-	No significant similarity	-	-

FE525632	ref XP_001211183.1	DNA polymerase gamma	<i>Aspergillus terreus</i> <i>NIH2624</i>	4e-22
FE525633	-	No significant similarity	-	-
FE525634	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE525635	gb AAP78735.1	nonribosomal peptide synthase	<i>Alternaria brassicae</i>	8e-06
FE525636	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	3e-06
FE525637	ref XP_001211183.1	DNA polymerase gamma	<i>Aspergillus terreus</i> <i>NIH2624</i>	4e-22
FE525638	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	3e-21
FE525639	-	No significant similarity	-	-
FE525640	ref XP_380291.1	hypothetical protein FG00115.1	<i>Gibberella zeae</i> PH-1	5e-06
FE525641	gb AAP78735.1	nonribosomal peptide synthase	<i>Alternaria brassicae</i>	4e-04
FE525642	ref XP_001272698.1	Dopey, N-terminal domain protein	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	4e-09
FE525643	gb AAP78735.1	nonribosomal peptide synthase	<i>Alternaria brassicae</i>	5e-06
FE525644	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	1e-42
FE525645	ref XP_001386445.1	Salicylate hydroxylase (Salicylate 1-monooxygenase)	<i>Pichia stipitis</i> CBS 6054	1e-16
FE525646	-	No significant similarity	-	-
FE525647	-	No significant similarity	-	-
FE525648	-	No significant similarity	-	-
FE525649	-	No significant similarity	-	-
FE525650	-	No significant similarity	-	-
FE525651	ref XP_001272698.1	Dopey, N-terminal domain protein	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-08
FE525652	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE525653	-	No significant similarity	-	-
FE525654	-	No significant similarity	-	-
FE525655	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-18
FE525656	ref XP_662910.1	hypothetical protein AN5306.2	<i>Aspergillus nidulans</i> <i>FGSC A4</i>	3e-11
FE525657	ref XP_001275274.1	PB1 domain protein, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	8e-06
FE525658	ref XP_662910.1	hypothetical protein AN5306.2	<i>Aspergillus nidulans</i> <i>FGSC A4</i>	3e-11
FE525659	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	9e-21
FE525660	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE525661	ref XP_001272698.1	Dopey, N-terminal domain protein	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	4e-09

FE525662	ref XP_001272698.1	Dopey, N-terminal domain protein	<i>Aspergillus clavatus</i> NRRL 1	4e-09
FE525663	-	No significant similarity	-	-
FE525664	gb AAP78735.1	nonribosomal peptide synthase	<i>Alternaria brassicae</i>	5e-06
FE525665	ref XP_001258253.1	ABC transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-21
FE525666	ref XP_001226392.1	hypothetical protein CHGG_08465	<i>Chaetomium globosum</i> CBS 148.51	3e-09
FE525667	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Neosartorya fischeri</i> NRRL 181	2e-24
FE525668	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Neosartorya fischeri</i> NRRL 181	2e-24
FE525669	ref XP_001248139.1	hypothetical protein CIMG_01910	<i>Coccidioides immitis</i> RS	7e-26
FE525670	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Neosartorya fischeri</i> NRRL 181	2e-24
FE525671	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Neosartorya fischeri</i> NRRL 181	2e-24
FE525672	gb AAP78735.1	nonribosomal peptide synthase	<i>Alternaria brassicae</i>	5e-06
FE525673	ref YP_840033.1	Glyoxalase/bleomycin resistance protein/dioxygenase	<i>Burkholderia</i> <i>cenocepacia</i> HI2424	5e-11
FE525674	ref XP_001386445.1	Salicylate hydroxylase (Salicylate 1-monooxygenase)	<i>Pichia stipitis</i> CBS 6054	2e-18
FE525675	ref XP_001211183.1	DNA polymerase gamma	<i>Aspergillus terreus</i> NIH2624	4e-22
FE525676	ref XP_001211183.1	DNA polymerase gamma	<i>Aspergillus terreus</i> NIH2624	4e-22
FE525677	-	No significant similarity	-	-
FE525678	-	No significant similarity	-	-
FE525679	ref XP_662910.1	hypothetical protein AN5306.2	<i>Aspergillus nidulans</i> FGSC A4	3e-11
FE525680	-	No significant similarity	-	-
FE525681	-	No significant similarity	-	-
FE525682	ref XP_001215759.1	woronin body major protein	<i>Aspergillus terreus</i> NIH2624	2e-26
FE525683	ref XP_001272343.1	indoleamine 2,3-dioxygenase family protein	<i>Aspergillus clavatus</i> NRRL 1	1e-27
FE525684	-	No significant similarity	-	-
FE525685	-	No significant similarity	-	-
FE525686	-	No significant similarity	-	-

FE525687	ref XP_001269335.1	AAA family ATPase/60S ribosome export protein Rix7, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-12
FE525688	-	No significant similarity	-	-
FE525689	ref XP_001268941.1	methyltransferase small domain protein	<i>Aspergillus clavatus</i> NRRL 1	7e-23
FE525690	ref XP_001262277.1	alpha/beta fold family hydrolase, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-06
FE525691	ref XP_001272129.1	LYR family protein	<i>Aspergillus clavatus</i> NRRL 1	2e-07
FE525692	ref XP_001228475.1	40s ribosomal protein S5	<i>Chaetomium globosum</i> CBS 148.51	2e-46
FE525693	ref XP_001268941.1	methyltransferase small domain protein	<i>Aspergillus clavatus</i> NRRL 1	2e-13
FE525694	-	No significant similarity	-	-
FE525695	-	No significant similarity	-	-
FE525696	ref XP_001264172.1	3-isopropylmalate dehydrogenase Leu2A	<i>Neosartorya fischeri</i> NRRL 181	6e-36
FE525697	-	No significant similarity	-	-
FE525698	dbj BAB33421.1	putative senescence-associated protein	<i>Pisum sativum</i>	3e-32
FE525699	gb AAB05810.1	super cysteine rich protein; SCRP	<i>Homo sapiens</i>	2e-14
FE525700	-	No significant similarity	-	-
FE525701	-	No significant similarity	-	-
FE525702	-	No significant similarity	-	-
FE525703	gb AAL78196.1 AF126048_1	UDP-N-acetylglucosamine:dolichyl phosphate N-acetylglucosamine-1-phosphate transferase; GPT	<i>Aspergillus niger</i>	5e-63
FE525704	ref XP_748912.1	small nuclear ribonucleoprotein SmE, putative	<i>Aspergillus fumigatus</i> Af293	1e-05
FE525705	-	No significant similarity	-	-
FE525706	-	No significant similarity	-	-
FE525707	-	No significant similarity	-	-
FE525708	ref XP_001244644.1	hypothetical protein CIMG_04085	<i>Coccidioides immitis</i> RS	4e-04
FE525709	-	No significant similarity	-	-
FE525710	ref XP_001213511.1	type II proteins geranylgeranyltransferase beta subunit	<i>Aspergillus terreus</i> NIH2624	4e-35
FE525711	ref XP_001247006.1	hypothetical protein CIMG_00777	<i>Coccidioides immitis</i> RS	1e-08
FE525712	-	No significant similarity	-	-
FE525713	-	No significant similarity	-	-
FE525714	ref XP_748019.1	pyruvate dehydrogenase kinase, putative	<i>Aspergillus fumigatus</i>	5e-56

FE525715	-	No significant similarity	Af293	-
FE525716	dbj BAB12047.1	alpha-crystallin-related protein	<i>Arthroderma benhamiae</i>	3e-10
FE525717	dbj BAE62191.1	unnamed protein product	<i>Aspergillus oryzae</i>	2e-16
FE525718	-	No significant similarity	-	-
FE525719	ref XP_001269249.1	adenosylhomocysteinase	<i>Aspergillus clavatus</i>	1e-37
			NRRL 1	
FE525720	-	No significant similarity	-	-
FE525721	dbj BAB12047.1	alpha-crystallin-related protein	<i>Arthroderma benhamiae</i>	8e-20
FE525722	ref XP_001260869.1	kinase activator (Atg17), putative	<i>Neosartorya fischeri</i>	3e-09
			NRRL 181	
FE525723	ref XP_001215759.1	woronin body major protein	<i>Aspergillus terreus</i>	5e-27
			NIH2624	
FE525724	-	No significant similarity	-	-
FE525725	ref XP_001258025.1	VHS domain protein	<i>Neosartorya fischeri</i>	1e-06
			NRRL 181	
FE525726	-	No significant similarity	-	-
FE525727	-	No significant similarity	-	-
FE525728	-	No significant similarity	-	-
FE525729	ref XP_001246822.1	hypothetical protein CIMG_00593	<i>Coccidioides immitis</i> RS	7e-20
FE525730	ref XP_001248599.1	eukaryotic translation initiation factor 3 subunit 7 homolog	<i>Coccidioides immitis</i> RS	1e-24
FE525731	-	No significant similarity	-	-
FE525732	-	No significant similarity	-	-
FE525733	gb EDK04135.1	conserved hypothetical protein	<i>Magnaporthe grisea</i> 70-15	3e-04
FE525734	dbj BAB12047.1	alpha-crystallin-related protein	<i>Arthroderma benhamiae</i>	9e-18
FE525735	dbj BAB12047.1	alpha-crystallin-related protein	<i>Arthroderma benhamiae</i>	7e-20
FE525736	-	No significant similarity	-	-
FE525737	-	No significant similarity	-	-
FE525738	gb ABB18373.1	chitobiase	<i>Coccidioides posadasii</i>	4e-50
FE525739	-	No significant similarity	-	-
FE525740	ref XP_001268837.1	fibrillarin	<i>Aspergillus clavatus</i>	3e-36
			NRRL 1	
FE525741	-	No significant similarity	-	-
FE525742	-	No significant similarity	-	-
FE525743	ref XP_001276345.1	histone H2A	<i>Aspergillus clavatus</i>	4e-19
			NRRL 1	
FE525744	ref XP_001261116.1	cyanate hydratase, putative	<i>Neosartorya fischeri</i>	9e-21

FE525745	ref XP_750031.1	mRNA cleavage and polyadenylation specificity factor complex subunit, putative	NRRL 181 <i>Aspergillus fumigatus</i> Af293	1e-06
FE525746	gb AAB05810.1	super cysteine rich protein; SCRP	<i>Homo sapiens</i>	4e-14
FE525747	gb AAB05810.1	super cysteine rich protein; SCRP	<i>Homo sapiens</i>	2e-14
FE525748	ref XP_001266066.1	60S ribosomal protein L23	<i>Neosartorya fischeri</i> NRRL 181	2e-34
FE525749	-	No significant similarity	-	-
FE525750	ref XP_001274417.1	oxidoreductase	<i>Aspergillus clavatus</i> NRRL 1	4e-44
FE525751	-	No significant similarity	-	-
FE525752	ref XP_001272850.1	mitochondrial ATPase inhibitor, putative	<i>Aspergillus clavatus</i> NRRL 1	1e-05
FE525753	ref XP_001274092.1	cysteinyl-tRNA synthetase	<i>Aspergillus clavatus</i> NRRL 1	4e-51
FE525754	-	No significant similarity	-	-
FE525755	dbj BAB12047.1	alpha-crystallin-related protein	<i>Arthroderma benhamiae</i>	7e-20
FE525756	gb EAL84586.2	DNA repair protein (Tof1), putative	<i>Aspergillus fumigatus</i> Af293	1e-24
FE525757	ref XP_001269377.1	DUF814 domain protein	<i>Aspergillus clavatus</i> NRRL 1	2e-05
FE525758	-	No significant similarity	-	-
FE525759	dbj BAB12047.1	alpha-crystallin-related protein	<i>Arthroderma benhamiae</i>	1e-16
FE525760	-	No significant similarity	-	-
FE525761	ref XP_752890.1	conserved hypothetical protein	<i>Aspergillus fumigatus</i> Af293	4e-13
FE525762	ref XP_750435.1	protein phosphatase 2C family protein	<i>Aspergillus fumigatus</i> Af293	3e-30
FE525763	ref XP_001244432.1	hypothetical protein CIMG_03873	<i>Coccidioides immitis</i> RS	8e-56
FE525764	ref XP_001272790.1	small nucleolar ribonucleoprotein complex subunit, putative	<i>Aspergillus clavatus</i> NRRL 1	4e-14
FE525765	-	No significant similarity	-	-
FE525766	ref NP_983024.1	ABR078Cp	<i>Ashbya gossypii</i> ATCC 10895	6e-09
FE525767	-	No significant similarity	-	-
FE525768	ref XP_001266177.1	mitochondrial large ribosomal subunit protein L1, putative	<i>Neosartorya fischeri</i> NRRL 181	3e-22
FE525769	emb CAK42954.1	unnamed protein product	<i>Aspergillus niger</i>	4e-24

FE525770	ref XP_001266773.1	DNA repair protein Rad50	<i>Neosartorya fischeri</i> NRRL 181	1e-19
FE525771	dbj BAB12047.1	alpha-crystallin-related protein	<i>Arthroderma benhamiae</i>	7e-20
FE525772	-	No significant similarity	-	-
FE525773	-	No significant similarity	-	-
FE525774	dbj BAB12047.1	alpha-crystallin-related protein	<i>Arthroderma benhamiae</i>	2e-19
FE525775	ref XP_001263008.1	heat shock protein Hsp30/Hsp42, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-09
FE525776	-	No significant similarity	-	-
FE525777	-	No significant similarity	-	-
FE525778	-	No significant similarity	-	-
FE525779	ref XP_748881.1	NACHT domain protein, putative	<i>Aspergillus fumigatus</i> Af293	6e-04
FE525780	-	No significant similarity	-	-
FE525781	-	No significant similarity	-	-
FE525782	-	No significant similarity	-	-
FE525783	-	No significant similarity	-	-
FE525784	ref XP_001246947.1	glycolipid 2-alpha-mannosyltransferase	<i>Coccidioides immitis</i> RS	2e-06
FE525785	ref XP_868845.1	hypothetical protein AN9463.2	<i>Aspergillus nidulans</i> FGSC A4	8e-04
FE525786	ref XP_001269539.1	DUF887 domain protein	<i>Aspergillus clavatus</i> NRRL 1	3e-12
FE525787	-	No significant similarity	-	-
FE525788	-	No significant similarity	-	-
FE525789	ref XP_001268953.1	regulator of nonsense transcripts, putative	<i>Aspergillus clavatus</i> NRRL 1	5e-25
FE525790	-	No significant similarity	-	-
FE525791	ref XP_001213531.1	proteasome component PRE3 precursor	<i>Aspergillus terreus</i> NIH2624	2e-30
FE525792	ref XP_001267735.1	short chain dehydrogenase/reductase family protein	<i>Aspergillus clavatus</i> NRRL 1	6e-20
FE525793	ref XP_001258400.1	anthranilate synthase component I, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-07
FE525794	ref XP_001270328.1	DUF803 domain protein	<i>Aspergillus clavatus</i> NRRL 1	6e-32
FE525795	-	No significant similarity	-	-
FE525796	gb AAX33296.1	heat shock protein 90	<i>Paracoccidioides</i> <i>brasiliensis</i>	7e-21

FE525797	ref XP_001268331.1	proteasome regulatory particle subunit (RpnI), putative	<i>Aspergillus clavatus</i> NRRL 1	1e-17
FE525798	-	No significant similarity	-	-
FE525799	dbj BAE62495.1	unnamed protein product	<i>Aspergillus oryzae</i>	6e-14
FE525800	-	No significant similarity	-	-
FE525801	ref XP_001247141.1	60S ribosomal protein L19	<i>Coccidioides immitis</i> RS	8e-06
FE525802	ref XP_001274718.1	amino acid permease, putative	<i>Aspergillus clavatus</i> NRRL 1	3e-12
FE525803	-	No significant similarity	-	-
FE525804	-	No significant similarity	-	-
FE525805	-	No significant similarity	-	-
FE525806	-	No significant similarity	-	-
FE525807	ref XP_749497.1	Ras GTPase activating protein, putative	<i>Aspergillus fumigatus</i> Af293	3e-04
FE525808	ref XP_001260419.1	60S ribosomal protein L10	<i>Neosartorya fischeri</i> NRRL 181	7e-39
FE525809	ref XP_001265142.1	DNA-directed DNA polymerase theta, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-27
FE525810	ref XP_001269539.1	DUF887 domain protein	<i>Aspergillus clavatus</i> NRRL 1	3e-12
FE525811	-	No significant similarity	-	-
FE525812	ref XP_682122.1	eukaryotic peptide chain release factor subunit 1	<i>Aspergillus nidulans</i> FGSC A4	1e-70
FE525813	-	No significant similarity	-	-
FE525814	ref XP_001248599.1	eukaryotic translation initiation factor 3 subunit 7 homolog	<i>Coccidioides immitis</i> RS	2e-22
FE525815	ref XP_001248599.1	eukaryotic translation initiation factor 3 subunit 7 homolog	<i>Coccidioides immitis</i> RS	8e-22
FE525816	ref XP_001212104.1	serine/threonine-protein kinase ssp1	<i>Aspergillus terreus</i> NIH2624	1e-03
FE525817	ref XP_754663.1	DNA polymerase epsilon, catalytic subunit A/POL2, putative	<i>Aspergillus fumigatus</i> Af293	1e-20
FE525818	ref XP_754663.1	DNA polymerase epsilon, catalytic subunit A/POL2, putative	<i>Aspergillus fumigatus</i> Af293	8e-06
FE525819	-	No significant similarity	-	-
FE525820	dbj BAE62495.1	unnamed protein product	<i>Aspergillus oryzae</i>	1e-03
FE525821	ref XP_754663.1	DNA polymerase epsilon, catalytic subunit A/POL2, putative	<i>Aspergillus fumigatus</i> Af293	2e-19
FE525822	ref XP_001267336.1	outer mitochondrial membrane protein porin	<i>Neosartorya fischeri</i> NRRL 181	8e-41

FE525823	ref XP_001258439.1	60S ribosomal protein L7a	<i>Neosartorya fischeri</i> NRRL 181	1e-08
FE525824	-	No significant similarity	-	-
FE525825	ref XP_001214623.1	eukaryotic translation initiation factor 2 gamma subunit	<i>Aspergillus terreus</i> NIH2624	6e-41
FE525826	-	No significant similarity	-	-
FE525827	ref ZP_00344940.1	COG0458: Carbamoylphosphate synthase large subunit (split gene in MJ)	<i>Nostoc punctiforme</i> PCC 73102	8e-12
FE525828	ref XP_001264383.1	C2H2 transcription factor (Rpn4), putative	<i>Neosartorya fischeri</i> NRRL 181	3e-54
FE525829	-	No significant similarity	-	-
FE525830	-	No significant similarity	-	-
FE525831	ref XP_001270884.1	eukaryotic translation initiation factor 3 subunit 2i, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-42
FE525832	ref XP_001265914.1	sugar transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	4e-21
FE525833	-	No significant similarity	-	-
FE525834	-	No significant similarity	-	-
FE525835	-	No significant similarity	-	-
FE525836	-	No significant similarity	-	-
FE525837	ref XP_001270884.1	eukaryotic translation initiation factor 3 subunit 2i, putative	<i>Aspergillus clavatus</i> NRRL 1	5e-42
FE525838	ref XP_001273963.1	MFS transporter, putative	<i>Aspergillus clavatus</i> NRRL 1	4e-11
FE525839	gb AAB06687.1	complement fixation antigen gb AAA96515.1 complement-fixation antigen	<i>Coccidioides posadasii</i>	6e-11
FE525840	ref XP_001258400.1	anthranilate synthase component I, putative	<i>Neosartorya fischeri</i> NRRL 181	4e-05
FE525841	-	No significant similarity	-	-
FE525842	ref XP_755151.1	NEDD8-like protein (RubA), putative	<i>Aspergillus fumigatus</i> Af293	3e-12
FE525843	ref XP_755086.1	prolidase pepP, putative	<i>Aspergillus fumigatus</i> Af293	1e-40
FE525844	-	No significant similarity	-	-
FE525845	ref XP_001271354.1	glutamate-cysteine ligase Gcs1, putative	<i>Aspergillus clavatus</i> NRRL 1	8e-14
FE525846	ref XP_001270884.1	eukaryotic translation initiation factor 3 subunit 2i, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-43
FE525847	ref XP_808355.1	formin	<i>Trypanosoma cruzi</i>	2e-04

FE525848	ref XP_001272561.1	heat shock Hsp30-like protein, putative	strain CL Brener <i>Aspergillus clavatus</i> NRRL 1	4e-12
FE525849	ref XP_001211399.1	acetoacetyl-CoA reductase	<i>Aspergillus terreus</i> NIH2624	4e-30
FE525850	-	No significant similarity	-	-
FE525851	-	No significant similarity	-	-
FE525852	-	No significant similarity	-	-
FE525853	-	No significant similarity	-	-
FE525854	-	No significant similarity	-	-
FE525855	ref XP_001260488.1	actin cytoskeleton protein (VIP1), putative	<i>Neosartorya fischeri</i> NRRL 181	3e-10
FE525856	ref XP_001258534.1	acyltransferase, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-14
FE525857	-	No significant similarity	-	-
FE525858	ref XP_001259741.1	ABC transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-22
FE525859	-	No significant similarity	-	-
FE525860	-	No significant similarity	-	-
FE525861	-	No significant similarity	-	-
FE525862	-	No significant similarity	-	-
FE525863	-	No significant similarity	-	-
FE525864	-	No significant similarity	-	-
FE525865	-	No significant similarity	-	-
FE525866	-	No significant similarity	-	-
FE525867	-	No significant similarity	-	-
FE525868	-	No significant similarity	-	-
FE525869	ref XP_754941.1	aminoalcoholphosphotransferase	<i>Aspergillus fumigatus</i> Af293	2e-14
FE525870	ref XP_001257402.1	C2HC5 finger protein	<i>Neosartorya fischeri</i> NRRL 181	7e-14
FE525871	-	No significant similarity	-	-
FE525872	ref XP_001257402.1	C2HC5 finger protein	<i>Neosartorya fischeri</i> NRRL 181	5e-13
FE525873	ref ZP_00344940.1	COG0458: Carbamoylphosphate synthase large subunit (split gene in MJ)	<i>Nostoc punctiforme</i> PCC 73102	8e-12
FE525874	ref XP_001248278.1	hypothetical protein CIMG_02049	<i>Coccidioides immitis</i> RS	1e-10
FE525875	ref XP_001209708.1	conserved hypothetical protein	<i>Aspergillus terreus</i>	2e-18

FE525876	ref XP_001262831.1	conserved hypothetical protein	NIH2624 <i>Neosartorya fischeri</i> NRRL 181	8e-06
FE525877	-	No significant similarity	-	-
FE525878	ref XP_001269644.1	N-acetyltransferase (Nat5), putative	<i>Aspergillus clavatus</i> NRRL 1	3e-21
FE525879	ref XP_754941.1	aminoalcoholphosphotransferase	<i>Aspergillus fumigatus</i> Af293	2e-14
FE525880	dbj BAA81686.1	expressed in cucumber hypocotyls	<i>Cucumis sativus</i>	1e-04
FE525881	-	No significant similarity	-	-
FE525882	-	No significant similarity	-	-
FE525883	sp P00048 CYC_NEUCR	Cytochrome c emb CAA29050.1 cytochrome c	<i>Neurospora crassa</i>	5e-11
FE525884	-	No significant similarity	-	-
FE525885	ref XP_001246545.1	phosphatidylethanolamine N-methyltransferase	<i>Coccidioides immitis</i> RS	1e-04
FE525886	ref XP_001275966.1	MFS multidrug transporter, putative	<i>Aspergillus clavatus</i> NRRL 1	3e-31
FE525887	-	No significant similarity	-	-
FE525888	-	No significant similarity	-	-
FE525889	ref XP_001213531.1	proteasome component PRE3 precursor	<i>Aspergillus terreus</i> NIH2624	6e-31
FE525890	-	No significant similarity	-	-
FE525891	-	No significant similarity	-	-
FE525892	-	No significant similarity	-	-
FE525893	ref XP_001247301.1	proteasome component	<i>Coccidioides immitis</i> RS	7e-24
FE525894	-	No significant similarity	-	-
FE525895	ref XP_721980.1	putative S-adenosyl-L-homocysteine hydrolase	<i>Candida albicans</i> SC5314	8e-08
FE525896	gb ABK60177.1	putative reverse transcriptase	<i>Zingiber officinale</i>	1e-03
FE525897	ref XP_721980.1	putative S-adenosyl-L-homocysteine hydrolase	<i>Candida albicans</i> SC5314	2e-10
FE525898	ref XP_001260488.1	actin cytoskeleton protein (VIP1), putative	<i>Neosartorya fischeri</i> NRRL 181	1e-08
FE525899	-	No significant similarity	-	-
FE525900	ref XP_001266911.1	acetyl-coenzyme A synthetase FacA	<i>Neosartorya fischeri</i> NRRL 181	9e-26
FE525901	-	No significant similarity	-	-
FE525902	-	No significant similarity	-	-
FE525903	ref XP_001257912.1	PH domain protein	<i>Neosartorya fischeri</i>	2e-10

FE525904	ref XP_001268018.1	PH domain protein	NRRL 181 <i>Aspergillus clavatus</i>	5e-11
FE525905	ref XP_001258439.1	60S ribosomal protein L7a	NRRL 1 <i>Neosartorya fischeri</i>	1e-08
FE525906	-	No significant similarity	NRRL 181	-
FE525907	ref ZP_00344940.1	COG0458: Carbamoylphosphate synthase large subunit (split gene in MJ)	- <i>Nostoc punctiforme</i> PCC 73102	7e-15
FE525908	ref XP_001264383.1	C2H2 transcription factor (Rpn4), putative	<i>Neosartorya fischeri</i> NRRL 181	1e-35
FE525909	gb ABG56823.1	hypothetical protein	<i>Klebsiella pneumoniae</i>	1e-05
FE525910	ref XP_001270884.1	eukaryotic translation initiation factor 3 subunit 2i, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-42
FE525911	-	No significant similarity	-	-
FE525912	-	No significant similarity	-	-
FE525913	-	No significant similarity	-	-
FE525914	ref XP_001263576.1	eukaryotic translation initiation factor 3 subunit 2i, putative	<i>Neosartorya fischeri</i> NRRL 181	5e-27
FE525915	-	No significant similarity	-	-
FE525916	ref XP_755151.1	NEDD8-like protein (RubA), putative	<i>Aspergillus fumigatus</i> Af293	1e-11
FE525917	-	No significant similarity	-	-
FE525918	ref XP_001263576.1	eukaryotic translation initiation factor 3 subunit 2i, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-27
FE525919	-	No significant similarity	-	-
FE525920	gb ABF82266.1	heat shock protein 30	<i>Penicillium marneffeii</i>	2e-11
FE525921	ref XP_001264679.1	conserved hypothetical protein	<i>Neosartorya fischeri</i> NRRL 181	4e-31
FE525922	-	No significant similarity	-	-
FE525923	ref XP_001214623.1	eukaryotic translation initiation factor 2 gamma subunit	<i>Aspergillus terreus</i> NIH2624	2e-25
FE525924	-	No significant similarity	-	-
FE525925	ref XP_001247301.1	proteasome component	<i>Coccidioides immitis</i> RS	2e-26
FE525926	-	No significant similarity	-	-
FE525927	ref XP_721980.1	putative S-adenosyl-L-homocysteine hydrolase	<i>Candida albicans</i> SC5314	2e-13
FE525928	-	No significant similarity	-	-
FE525929	-	No significant similarity	-	-

FE525930	gb EAL86837.2	sodium/phosphate symporter, putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	2e-11
FE525931	-	No significant similarity	-	-
FE525932	ref XP_001266911.1	acetyl-coenzyme A synthetase FacA	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	5e-29
FE525933	-	No significant similarity	-	-
FE525934	-	No significant similarity	-	-
FE525935	ref XP_001257912.1	PH domain protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	9e-13
FE525936	-	No significant similarity	-	-
FE525937	-	No significant similarity	-	-
FE525938	-	No significant similarity	-	-
FE525939	ref XP_001267735.1	short chain dehydrogenase/reductase family protein	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	4e-21
FE525940	ref XP_001213531.1	proteasome component PRE3 precursor	<i>Aspergillus terreus</i> <i>NIH2624</i>	1e-30
FE525941	ref XP_001260315.1	short chain dehydrogenase/reductase family protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-19
FE525942	ref XP_001258400.1	anthranilate synthase component I, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-07
FE525943	ref XP_001270328.1	DUF803 domain protein	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-29
FE525944	-	No significant similarity	-	-
FE525945	gb AAX33296.1	heat shock protein 90	<i>Paracoccidioides</i> <i>brasiliensis</i>	7e-21
FE525946	-	No significant similarity	-	-
FE525947	-	No significant similarity	-	-
FE525948	dbj BAE62495.1	unnamed protein product	<i>Aspergillus oryzae</i>	1e-14
FE525949	-	No significant similarity	-	-
FE525950	ref XP_001247141.1	60S ribosomal protein L19	<i>Coccidioides immitis</i> RS	8e-06
FE525951	-	No significant similarity	-	-
FE525952	-	No significant similarity	-	-
FE525953	-	No significant similarity	-	-
FE525954	ref XP_001275476.1	pyruvate dehydrogenase kinase	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-11
FE525955	-	No significant similarity	-	-
FE525956	-	No significant similarity	-	-
FE525957	-	No significant similarity	-	-

FE525958	-	No significant similarity	-	-
FE525959	ref XP_001265142.1	DNA-directed DNA polymerase theta, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-27
FE525960	ref XP_001247141.1	60S ribosomal protein L19	<i>Coccidioides immitis</i> RS	6e-11
FE525961	-	No significant similarity	-	-
FE525962	ref XP_682122.1	eukaryotic peptide chain release factor subunit 1	<i>Aspergillus nidulans</i> <i>FGSC A4</i>	3e-52
FE525963	-	No significant similarity	-	-
FE525964	ref XP_001248599.1	eukaryotic translation initiation factor 3 subunit 7 homolog	<i>Coccidioides immitis</i> RS	5e-21
FE525965	-	No significant similarity	-	-
FE525966	-	No significant similarity	-	-
FE525967	ref XP_746577.1	peroxisomal multifunctional beta-oxidation protein (MFP), putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	2e-19
FE525968	ref XP_001211399.1	acetoacetyl-CoA reductase	<i>Aspergillus terreus</i> <i>NIH2624</i>	4e-30
FE525969	ref XP_001268779.1	BAP31 domain protein, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	4e-14
FE525970	-	No significant similarity	-	-
FE525971	-	No significant similarity	-	-
FE525972	ref XP_001275966.1	MFS multidrug transporter, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	3e-31
FE525973	-	No significant similarity	-	-
FE525974	-	No significant similarity	-	-
FE525975	-	No significant similarity	-	-
FE525976	-	No significant similarity	-	-
FE525977	-	No significant similarity	-	-
FE525978	-	No significant similarity	-	-
FE525979	-	No significant similarity	-	-
FE525980	-	No significant similarity	-	-
FE525981	-	No significant similarity	-	-
FE525982	-	No significant similarity	-	-
FE525983	ref XP_754941.1	aminoalcoholphosphotransferase	<i>Aspergillus fumigatus</i> <i>Af293</i>	2e-14
FE525984	ref XP_001211399.1	acetoacetyl-CoA reductase	<i>Aspergillus terreus</i> <i>NIH2624</i>	2e-29
FE525985	-	No significant similarity	-	-
FE525986	-	No significant similarity	-	-
FE525987	ref XP_001209708.1	conserved hypothetical protein	<i>Aspergillus terreus</i>	2e-18

FE525988	ref ZP_00344940.1	COG0458: Carbamoylphosphate synthase large subunit (split gene in MJ)	<i>NIH2624</i> <i>Nostoc punctiforme</i> PCC 73102	8e-12
FE525989	-	No significant similarity	-	-
FE525990	-	No significant similarity	-	-
FE525991	-	No significant similarity	-	-
FE525992	ref XP_754941.1	aminoalcoholphosphotransferase	<i>Aspergillus fumigatus</i> Af293	2e-14
FE525993	-	No significant similarity	-	-
FE525994	ref XP_001246545.1	phosphatidylethanolamine N-methyltransferase	<i>Coccidioides immitis</i> RS	1e-04
FE525995	-	No significant similarity	-	-
FE525996	-	No significant similarity	-	-
FE525997	sp P00048 CYC_NEUCR	Cytochrome c emb CAA29050.1 cytochrome c	<i>Neurospora crassa</i>	5e-11
FE525998	ref XP_001209915.1	conserved hypothetical protein	<i>Aspergillus terreus</i> NIH2624	1e-07
FE525999	-	No significant similarity	-	-
FE526000	ref XP_001269644.1	N-acetyltransferase (Nat5), putative	<i>Aspergillus clavatus</i> NRRL 1	3e-21
FE526001	-	No significant similarity	-	-
FE526002	-	No significant similarity	-	-
FE526003	-	No significant similarity	-	-
FE526004	gb ABG67901.1	phospholipase B	<i>Trichophyton rubrum</i>	4e-19
FE526005	ref XP_751277.1	acyltransferase, putative	<i>Aspergillus fumigatus</i> Af293	4e-09
FE526006	-	No significant similarity	-	-
FE526007	ref XP_001258439.1	60S ribosomal protein L7a	<i>Neosartorya fischeri</i> NRRL 181	1e-08
FE526008	-	No significant similarity	-	-
FE526009	gb EAL90029.2	translation initiation factor EF-2 gamma subunit, putative	<i>Aspergillus fumigatus</i> Af293	4e-21
FE526010	-	No significant similarity	-	-
FE526011	-	No significant similarity	-	-
FE526012	ref ZP_00344940.1	COG0458: Carbamoylphosphate synthase large subunit (split gene in MJ)	<i>Nostoc punctiforme</i> PCC 73102	2e-11
FE526013	ref XP_001264383.1	C2H2 transcription factor (Rpn4), putative	<i>Neosartorya fischeri</i> NRRL 181	8e-51
FE526014	-	No significant similarity	-	-
FE526015	-	No significant similarity	-	-

FE526016	ref XP_001270884.1	eukaryotic translation initiation factor 3 subunit 2i, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-42
FE526017	-	No significant similarity	-	-
FE526018	-	No significant similarity	-	-
FE526019	-	No significant similarity	-	-
FE526020	-	No significant similarity	-	-
FE526021	ref XP_001270884.1	eukaryotic translation initiation factor 3 subunit 2i, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	5e-42
FE526022	-	No significant similarity	-	-
FE526023	gb AAB06687.1	complement fixation antigen gb AAA96515.1 complement-fixation antigen	<i>Coccidioides posadasii</i>	6e-11
FE526024	ref XP_755151.1	NEDD8-like protein (RubA), putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	1e-08
FE526025	-	No significant similarity	-	-
FE526026	ref XP_001263576.1	eukaryotic translation initiation factor 3 subunit 2i, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	3e-28
FE526027	-	No significant similarity	-	-
FE526028	-	No significant similarity	-	-
FE526029	emb CAM14118.1	novel protein containing SEA domains	<i>Danio rerio</i>	1e-04
FE526030	-	No significant similarity	-	-
FE526031	-	No significant similarity	-	-
FE526032	-	No significant similarity	-	-
FE526033	ref XP_001271726.1	tyrosyl-tRNA synthetase, mitochondrial precursor (tyrosine--tRNA ligase) (tyrrs)	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	1e-27
FE526034	-	No significant similarity	-	-
FE526035	gb AAO52807.1	hypothetical protein	<i>Bacillus megaterium</i>	2e-04
FE526036	-	No significant similarity	-	-
FE526037	ref XP_001270287.1	60S ribosomal protein L24	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	3e-34
FE526038	-	No significant similarity	-	-
FE526039	-	No significant similarity	-	-
FE526040	-	No significant similarity	-	-
FE526041	-	No significant similarity	-	-
FE526042	-	No significant similarity	-	-
FE526043	-	No significant similarity	-	-
FE526044	ref XP_001264535.1	MFS peptide transporter, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	3e-21
FE526045	-	No significant similarity	-	-
FE526046	-	No significant similarity	-	-

FE526047	-	No significant similarity	-	-
FE526048	ref XP_001272561.1	heat shock Hsp30-like protein, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	4e-06
FE526049	-	No significant similarity	-	-
FE526050	-	No significant similarity	-	-
FE526051	ref XP_001260474.1	oxidoreductase, 2-nitropropane dioxygenase family, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	2e-49
FE526052	ref XP_751914.1	cytochrome c peroxidase, putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	3e-04
FE526053	-	No significant similarity	-	-
FE526054	sp P59769 GRP78_ASPAW	78 kDa glucose-regulated protein homolog precursor (GRP 78) (Immunoglobulin heavy chain-binding protein homolog) (BiP) sp P83616 GRP78_ASPNG 78 kDa glucose-regulated protein homolog precursor (GRP 78) (Immunoglobulin heavy chain-binding protein homolog) (	<i>Aspergillus kawachii</i>	1e-52
FE526055	ref XP_001247978.1	30 kDa heat shock protein	<i>Coccidioides immitis</i> RS	7e-04
FE526056	-	No significant similarity	-	-
FE526057	-	No significant similarity	-	-
FE526058	-	No significant similarity	-	-
FE526059	-	No significant similarity	-	-
FE526060	ref XP_001257997.1	NADH-ubiquinone oxidoreductase 304 kDa subunit precursor	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	5e-67
FE526061	ref XP_001271404.1	heat shock protein Hsp30/Hsp42, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-08
FE526062	-	No significant similarity	-	-
FE526063	-	No significant similarity	-	-
FE526064	-	No significant similarity	-	-
FE526065	-	No significant similarity	-	-
FE526066	-	No significant similarity	-	-
FE526067	-	No significant similarity	-	-
FE526068	-	No significant similarity	-	-
FE526069	-	No significant similarity	-	-
FE526070	-	No significant similarity	-	-
FE526071	-	No significant similarity	-	-
FE526072	gb ABB96277.1	hesp-767	<i>Melampsora lini</i>	7e-05
FE526073	-	No significant similarity	-	-
FE526074	-	No significant similarity	-	-
FE526075	-	No significant similarity	-	-

FE526076	ref XP_001213531.1	proteasome component PRE3 precursor	<i>Aspergillus terreus</i> NIH2624	1e-29
FE526077	ref XP_001220639.1	proteasome subunit beta type 7 precursor	<i>Chaetomium globosum</i> CBS 148.51	2e-25
FE526078	-	No significant similarity	-	-
FE526079	gb EAL86837.2	sodium/phosphate symporter, putative	<i>Aspergillus fumigatus</i> Af293	6e-12
FE526080	ref XP_001266911.1	acetyl-coenzyme A synthetase FacA	<i>Neosartorya fischeri</i> NRRL 181	5e-29
FE526081	-	No significant similarity	-	-
FE526082	-	No significant similarity	-	-
FE526083	-	No significant similarity	-	-
FE526084	ref XP_001257912.1	PH domain protein	<i>Neosartorya fischeri</i> NRRL 181	9e-13
FE526085	ref XP_001212691.1	30 kDa heat shock protein	<i>Aspergillus terreus</i> NIH2624	1e-03
FE526086	-	No significant similarity	-	-
FE526087	-	No significant similarity	-	-
FE526088	ref XP_001272465.1	MFS multidrug transporter, putative	<i>Aspergillus clavatus</i> NRRL 1	5e-23
FE526089	-	No significant similarity	-	-
FE526090	-	No significant similarity	-	-
FE526091	ref XP_001240305.1	hypothetical protein CIMG_07468	<i>Coccidioides immitis</i> RS	2e-32
FE526092	-	No significant similarity	-	-
FE526093	ref XP_001239696.1	predicted protein	<i>Coccidioides immitis</i> RS	1e-07
FE526094	-	No significant similarity	-	-
FE526095	-	No significant similarity	-	-
FE526096	-	No significant similarity	-	-
FE526097	gb ABM92787.1	phytase	<i>Aspergillus oryzae</i>	4e-08
FE526098	-	No significant similarity	-	-
FE526099	-	No significant similarity	-	-
FE526100	-	No significant similarity	-	-
FE526101	-	No significant similarity	-	-
FE526102	-	No significant similarity	-	-
FE526103	-	No significant similarity	-	-
FE526104	-	No significant similarity	-	-
FE526105	-	No significant similarity	-	-
FE526106	gb EDJ95143.1	60S ribosomal protein L7	<i>Magnaporthe grisea</i> 70-	5e-11

FE526107	-	No significant similarity	-	-
FE526108	-	No significant similarity	-	-
FE526109	ref XP_001248014.1	predicted protein	<i>Coccidioides immitis</i> RS	7e-04
FE526110	-	No significant similarity	-	-
FE526111	-	No significant similarity	-	-
FE526112	-	No significant similarity	-	-
FE526113	-	No significant similarity	-	-
FE526114	ref XP_001271793.1	DEAD/DEAH box helicase, putative	<i>Aspergillus clavatus</i> NRRL 1	1e-48
FE526115	-	No significant similarity	-	-
FE526116	-	No significant similarity	-	-
FE526117	ref XP_001271793.1	DEAD/DEAH box helicase, putative	<i>Aspergillus clavatus</i> NRRL 1	1e-49
FE526118	-	No significant similarity	-	-
FE526119	-	No significant similarity	-	-
FE526120	-	No significant similarity	-	-
FE526121	-	No significant similarity	-	-
FE526122	-	No significant similarity	-	-
FE526123	ref XP_001239221.1	GTP-binding nuclear protein GSP1/Ran	<i>Coccidioides immitis</i> RS	4e-25
FE526124	-	No significant similarity	-	-
FE526125	-	No significant similarity	-	-
FE526126	-	No significant similarity	-	-
FE526127	-	No significant similarity	-	-
FE526128	ref XP_001276376.1	bZIP transcription factor JlbA/IDI-4	<i>Aspergillus clavatus</i> NRRL 1	8e-05
FE526129	-	No significant similarity	-	-
FE526130	-	No significant similarity	-	-
FE526131	-	No significant similarity	-	-
FE526132	-	No significant similarity	-	-
FE526133	-	No significant similarity	-	-
FE526134	-	No significant similarity	-	-
FE526135	-	No significant similarity	-	-
FE526136	-	No significant similarity	-	-
FE526137	-	No significant similarity	-	-
FE526138	-	No significant similarity	-	-
FE526139	-	No significant similarity	-	-
FE526140	ref XP_755151.1	NEDD8-like protein (RubA), putative	<i>Aspergillus fumigatus</i>	7e-12

FE526141	-	No significant similarity	-	-
FE526142	-	No significant similarity	-	-
FE526143	-	No significant similarity	-	-
FE526144	-	No significant similarity	-	-
FE526145	-	No significant similarity	-	-
FE526146	-	No significant similarity	-	-
FE526147	-	No significant similarity	-	-
FE526148	-	No significant similarity	-	-
FE526149	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-18
FE526150	ref XP_001266920.1	kynurenine aminotransferase, putative	<i>Neosartorya fischeri</i>	2e-62
			<i>NRRL 181</i>	
FE526151	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	3e-24
FE526152	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	3e-06
FE526153	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-25
FE526154	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526155	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	3e-10
			<i>NRRL 1</i>	
FE526156	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-18
FE526157	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-26
FE526158	ref XP_001266920.1	kynurenine aminotransferase, putative	<i>Neosartorya fischeri</i>	3e-63
			<i>NRRL 181</i>	
FE526159	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-26
FE526160	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	5e-26
FE526161	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	1e-20
FE526162	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-18
FE526163	-	No significant similarity	-	-
FE526164	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	5e-26
FE526165	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-26
FE526166	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526167	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	3e-10
			<i>NRRL 1</i>	
FE526168	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526169	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	4e-26
FE526170	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	3e-27
FE526171	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526172	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-18
FE526173	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	3e-21

FE526174	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i> NRRL 1	3e-10
FE526175	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526176	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526177	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	3e-06
FE526178	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Neosartorya fischeri</i> NRRL 181	2e-24
FE526179	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-26
FE526180	-	No significant similarity	-	-
FE526181	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	3e-06
FE526182	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	3e-06
FE526183	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526184	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-18
FE526185	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526186	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-18
FE526187	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526188	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	1e-20
FE526189	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526190	-	No significant similarity	-	-
FE526191	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	3e-13
FE526192	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526193	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	3e-06
FE526194	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526195	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i> NRRL 1	8e-04
FE526196	-	No significant similarity	-	-
FE526197	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526198	ref XP_001266920.1	kynurenine aminotransferase, putative	<i>Neosartorya fischeri</i> NRRL 181	4e-62
FE526199	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	3e-26
FE526200	-	No significant similarity	-	-
FE526201	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526202	-	No significant similarity	-	-
FE526203	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	2e-24
FE526204	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	5e-26
FE526205	-	No significant similarity	-	-
FE526206	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	1e-42
FE526207	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	6e-43

FE526208	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	3e-06
FE526209	ref XP_751710.1	kynurenine aminotransferase, putative	<i>Aspergillus fumigatus</i> Af293	2e-60
FE526210	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Neosartorya fischeri</i> NRRL 181	3e-25
FE526211	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	1e-20
FE526212	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	4e-13
FE526213	sp Q2URJ0 PPME1_ASPOR	Protein phosphatase methylesterase 1 (PMe-1)	<i>Aspergillus oryzae</i>	2e-25
FE526214	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	1e-06
FE526215	-	No significant similarity	-	-
FE526216	-	No significant similarity	-	-
FE526217	-	No significant similarity	-	-
FE526218	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526219	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526220	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	5e-22
FE526221	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526222	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	2e-17
FE526223	dbj BAC01275.1	cytochrome P450nor	<i>Aspergillus oryzae</i>	2e-16
FE526224	gb EEQ27465.1	copper resistance-associated P-type ATPase, putative	<i>Microsporum canis</i> CBS 113480	8e-24
FE526225	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i> NRRL 1	3e-10
FE526226	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	3e-22
FE526227	ref XP_001262830.1	sexual development protein EsdC, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-11
FE526228	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	5e-22
FE526229	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	9e-29
FE526230	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	2e-23
FE526231	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526232	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-23
FE526233	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	6e-47
FE526234	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	8e-48
FE526235	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	3e-06
FE526236	ref XP_751710.1	kynurenine aminotransferase, putative	<i>Aspergillus fumigatus</i> Af293	9e-62
FE526237	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	4e-06

FE526238	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	2e-22
FE526239	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	3e-24
FE526240	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	4e-14
FE526241	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	3e-24
FE526242	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	3e-44
FE526243	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE526244	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-23
FE526245	ref XP_751710.1	kynurenine aminotransferase, putative	<i>Aspergillus fumigatus</i>	3e-27
			<i>Af293</i>	
FE526246	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	2e-43
FE526247	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	2e-16
FE526248	dbj BAC01275.1	cytochrome P450nor	<i>Aspergillus oryzae</i>	6e-16
FE526249	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	3e-09
			<i>NRRL 1</i>	
FE526250	ref XP_001266920.1	kynurenine aminotransferase, putative	<i>Neosartorya fischeri</i>	1e-62
			<i>NRRL 181</i>	
FE526251	ref XP_001211183.1	DNA polymerase gamma	<i>Aspergillus terreus</i>	4e-22
			<i>NIH2624</i>	
FE526252	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	9e-23
FE526253	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	3e-27
FE526254	dbj BAC01275.1	cytochrome P450nor	<i>Aspergillus oryzae</i>	6e-16
FE526255	ref XP_001266920.1	kynurenine aminotransferase, putative	<i>Neosartorya fischeri</i>	8e-55
			<i>NRRL 181</i>	
FE526256	ref XP_001211183.1	DNA polymerase gamma	<i>Aspergillus terreus</i>	4e-22
			<i>NIH2624</i>	
FE526257	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Neosartorya fischeri</i>	3e-25
			<i>NRRL 181</i>	
FE526258	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	8e-06
FE526259	dbj BAC01275.1	cytochrome P450nor	<i>Aspergillus oryzae</i>	1e-15
FE526260	ref XP_001211183.1	DNA polymerase gamma	<i>Aspergillus terreus</i>	4e-22
			<i>NIH2624</i>	
FE526261	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	7e-23
FE526262	dbj BAC01275.1	cytochrome P450nor	<i>Aspergillus oryzae</i>	6e-16
FE526263	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Neosartorya fischeri</i>	2e-24
			<i>NRRL 181</i>	
FE526264	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	7e-23
FE526265	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	3e-21
FE526266	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Neosartorya fischeri</i>	2e-24

FE526267	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	NRRL 181	
			<i>Neosartorya fischeri</i>	2e-24
FE526268	gb ABF22673.1	copper-sulfate regulated protein 1	NRRL 181	
FE526269	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Ajellomyces capsulatus</i>	7e-18
FE526270	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Trichophyton rubrum</i>	2e-23
			<i>Neosartorya fischeri</i>	6e-24
			NRRL 181	
FE526271	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Neosartorya fischeri</i>	2e-24
			NRRL 181	
FE526272	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	4e-05
FE526273	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-18
FE526274	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	7e-23
FE526275	ref XP_754782.1	hypothetical protein Afu3g08320	<i>Aspergillus fumigatus</i>	3e-07
			Af293	
FE526276	ref XP_001265341.1	ribosomal protein/carboxylic ester hydrolase (Ppe1), putative	<i>Neosartorya fischeri</i>	2e-24
			NRRL 181	
FE526277	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	3e-10
			NRRL 1	
FE526278	-	No significant similarity	-	-
FE526279	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	2e-18
FE526280	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	2e-43
FE526281	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	3e-06
FE526282	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	3e-10
			NRRL 1	
FE526283	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	3e-06
FE526284	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	8e-48
			<i>Alternaria brassicae</i>	5e-06
FE526285	gb AAP78735.1	nonribosomal peptide synthase	<i>Emericella nidulans</i>	3e-25
FE526286	gb AAP23304.1	NIMA interactive protein	<i>Alternaria brassicae</i>	5e-06
FE526287	gb AAP78735.1	nonribosomal peptide synthase	<i>Aspergillus clavatus</i>	3e-10
FE526288	ref XP_001269482.1	Leucine Rich Repeat domain protein	NRRL 1	
			<i>Emericella nidulans</i>	2e-24
FE526289	gb AAP23304.1	NIMA interactive protein	<i>Alternaria brassicae</i>	5e-06
FE526290	gb AAP78735.1	nonribosomal peptide synthase	<i>Emericella nidulans</i>	2e-25
FE526291	gb AAP23304.1	NIMA interactive protein	<i>Aspergillus clavatus</i>	4e-10
FE526292	ref XP_001269482.1	Leucine Rich Repeat domain protein	NRRL 1	
			<i>Emericella nidulans</i>	6e-20
FE526293	gb AAP23304.1	NIMA interactive protein		

FE526294	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	3e-47
FE526295	gb AAP78735.1	nonribosomal peptide synthase	<i>Alternaria brassicae</i>	5e-06
FE526296	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	2e-16
FE526297	gb AAP78735.1	nonribosomal peptide synthase	<i>Alternaria brassicae</i>	4e-07
FE526298	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	3e-10
			<i>NRRL 1</i>	
FE526299	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	2e-19
FE526300	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	8e-48
FE526301	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	2e-10
FE526302	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	2e-11
FE526303	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE526304	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	3e-49
FE526305	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE526306	ref YP_001098942.1	glycerophosphodiester phosphodiesterase, cytosolic	<i>Herminiimonas arsenicoxydans</i>	1e-07
FE526307	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE526308	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE526309	ref YP_001098942.1	glycerophosphodiester phosphodiesterase, cytosolic	<i>Herminiimonas arsenicoxydans</i>	1e-07
FE526310	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE526311	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	3e-10
			<i>NRRL 1</i>	
FE526312	ref YP_001098942.1	glycerophosphodiester phosphodiesterase, cytosolic	<i>Herminiimonas arsenicoxydans</i>	1e-07
FE526313	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE526314	gb AAP78735.1	nonribosomal peptide synthase	<i>Alternaria brassicae</i>	5e-06
FE526315	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	3e-10
			<i>NRRL 1</i>	
FE526316	ref YP_001098942.1	glycerophosphodiester phosphodiesterase, cytosolic	<i>Herminiimonas arsenicoxydans</i>	1e-07
FE526317	-	No significant similarity	-	-
FE526318	ref XP_001276510.1	glucosamine-6-phosphate isomerase	<i>Aspergillus clavatus</i>	5e-46
			<i>NRRL 1</i>	
FE526319	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i>	3e-40
			<i>NRRL 181</i>	

FE526320	ref XP_001273963.1	MFS transporter, putative	<i>Aspergillus clavatus</i> NRRL 1	4e-11
FE526321	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i> <i>verrucosum</i>	8e-48
FE526322	-	No significant similarity	-	-
FE526323	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i> NRRL 181	5e-37
FE526324	-	No significant similarity	-	-
FE526325	-	No significant similarity	-	-
FE526326	-	No significant similarity	-	-
FE526327	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	2e-13
FE526328	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i> NRRL 181	3e-28
FE526329	ref XP_001245309.1	hypothetical protein CIMG_04750	<i>Coccidioides immitis</i> RS	9e-07
FE526330	-	No significant similarity	-	-
FE526331	-	No significant similarity	-	-
FE526332	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i> <i>verrucosum</i>	2e-47
FE526333	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i> NRRL 181	1e-34
FE526334	-	No significant similarity	-	-
FE526335	ref YP_270051.1	ISCps6, transposase	<i>Colwellia</i> <i>psychrerythraea</i> 34H	2e-11
FE526336	-	No significant similarity	-	-
FE526337	ref XP_001264535.1	MFS peptide transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	5e-19
FE526338	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i> <i>verrucosum</i>	8e-43
FE526339	-	No significant similarity	-	-
FE526340	-	No significant similarity	-	-
FE526341	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i> <i>verrucosum</i>	5e-50
FE526342	ref XP_752247.1	cytochrome P450 phenylacetate hydroxylase, putative	<i>Aspergillus fumigatus</i> Af293	2e-18
FE526343	ref XP_746951.1	MFS peptide transporter, putative	<i>Aspergillus fumigatus</i> Af293	1e-39
FE526344	-	No significant similarity	-	-

FE526345	-	No significant similarity	-	-
FE526346	gb AAZ32401.1	peptide transporter PTR2A	<i>Hebeloma cylindrosporum</i>	1e-11
FE526347	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	4e-49
FE526348	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-19
FE526349	gb AAR02424.1	subtilisin-like protease SUB5	<i>Trichophyton rubrum</i>	6e-46
FE526350	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-20
FE526351	ref XP_747457.1	DUF895 domain membrane protein	<i>Aspergillus fumigatus</i> Af293	6e-10
FE526352	ref XP_747214.1	glucosamine-6-phosphate deaminase, putative	<i>Aspergillus fumigatus</i> Af293	6e-49
FE526353	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	8e-45
FE526354	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	5e-47
FE526355	-	No significant similarity	-	-
FE526356	gb ABL84984.1	metalloprotease Mep3	<i>Trichophyton equinum</i>	8e-41
FE526357	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	3e-72
FE526358	ref XP_001246883.1	hypothetical protein CIMG_00654	<i>Coccidioides immitis</i> RS	3e-04
FE526359	gb AAR02424.1	subtilisin-like protease SUB5	<i>Trichophyton rubrum</i>	4e-72
FE526360	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	9e-31
FE526361	ref XP_001244025.1	hypothetical protein CIMG_03466	<i>Coccidioides immitis</i> RS	3e-29
FE526362	ref XP_001272561.1	heat shock Hsp30-like protein, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-07
FE526363	-	No significant similarity	-	-
FE526364	-	No significant similarity	-	-
FE526365	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i> NRRL 181	5e-40
FE526366	ref XP_754004.1	MFS transporter, putative	<i>Aspergillus fumigatus</i> Af293	4e-35
FE526367	-	No significant similarity	-	-
FE526368	-	No significant similarity	-	-
FE526369	ref XP_001270309.1	benzoate 4-monooxygenase cytochrome P450	<i>Aspergillus clavatus</i> NRRL 1	3e-35
FE526370	ref XP_747214.1	glucosamine-6-phosphate deaminase, putative	<i>Aspergillus fumigatus</i> Af293	2e-55

FE526371	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	6e-28
FE526372	-	No significant similarity	-	-
FE526373	ref XP_001244025.1	hypothetical protein CIMG_03466	<i>Coccidioides immitis</i> RS	4e-27
FE526374	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i>	2e-33
			NRRL 181	
FE526375	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	1e-47
FE526376	ref XP_001266791.1	conserved lysine-rich protein, putative	<i>Neosartorya fischeri</i>	8e-04
			NRRL 181	
FE526377	-	No significant similarity	-	-
FE526378	ref XP_747457.1	DUF895 domain membrane protein	<i>Aspergillus fumigatus</i>	5e-12
			Af293	
FE526379	gb AAR02424.1	subtilisin-like protease SUB5	<i>Trichophyton rubrum</i>	5e-77
FE526380	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i>	3e-20
			NRRL 181	
FE526381	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	4e-44
FE526382	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i>	5e-21
			NRRL 181	
FE526383	gb AAR02424.1	subtilisin-like protease SUB5	<i>Trichophyton rubrum</i>	3e-45
FE526384	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	5e-41
FE526385	-	No significant similarity	-	-
FE526386	-	No significant similarity	-	-
FE526387	-	No significant similarity	-	-
FE526388	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i>	4e-21
			NRRL 181	
FE526389	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	2e-48
FE526390	-	No significant similarity	-	-
FE526391	emb CAJ83813.1	CHK1 checkpoint homolog (S. pombe)	<i>Xenopus tropicalis</i>	7e-08
FE526392	-	No significant similarity	-	-
FE526393	ref XP_001246883.1	hypothetical protein CIMG_00654	<i>Coccidioides immitis</i> RS	3e-04
FE526394	ref XP_001244025.1	hypothetical protein CIMG_03466	<i>Coccidioides immitis</i> RS	1e-19
FE526395	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	3e-53
FE526396	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	3e-28

FE526397	ref XP_001244025.1	hypothetical protein CIMG_03466	<i>Coccidioides immitis</i> RS	9e-31
FE526398	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	2e-44
FE526399	-	No significant similarity	-	-
FE526400	-	No significant similarity	-	-
FE526401	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	6e-55
FE526402	ref XP_001229086.1	hypothetical protein CHGG_02570	<i>Chaetomium globosum</i> <i>CBS 148.51</i>	1e-03
FE526403	-	No significant similarity	-	-
FE526404	-	No significant similarity	-	-
FE526405	ref XP_001246883.1	hypothetical protein CIMG_00654	<i>Coccidioides immitis</i> RS	3e-04
FE526406	ref YP_080386.1	oxalate decarboxylase	<i>Bacillus licheniformis</i> <i>ATCC 14580</i>	2e-09
FE526407	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	2e-26
FE526408	-	No significant similarity	-	-
FE526409	gb AAR02424.1	subtilisin-like protease SUB5	<i>Trichophyton rubrum</i>	4e-41
FE526410	-	No significant similarity	-	-
FE526411	gb ABL84992.1	metalloprotease Mep4	<i>Trichophyton tonsurans</i>	5e-40
FE526412	-	No significant similarity	-	-
FE526413	gb AAR11462.1	subtilisin-like protease SUB3	<i>Trichophyton rubrum</i>	1e-24
FE526414	-	No significant similarity	-	-
FE526415	ref XP_001265751.1	phytanoyl-CoA dioxygenase family protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-18
FE526416	emb CAJ04955.1	hypothetical protein, unknown function	<i>Leishmania major</i>	7e-07
FE526417	gb AAN62296.1 AF440524_83	transposase domain protein	<i>Pseudomonas aeruginosa</i>	1e-22
FE526418	ref XP_001262224.1	K ⁺ /H ⁺ antiporter, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	4e-41
FE526419	ref XP_001275855.1	gamma interferon inducible lysosomal thiol reductase (GILT), putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	3e-06
FE526420	-	No significant similarity	-	-
FE526421	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	2e-36
FE526422	-	No significant similarity	-	-
FE526423	gb AAR11462.1	subtilisin-like protease SUB3	<i>Trichophyton rubrum</i>	1e-21
FE526424	-	No significant similarity	-	-
FE526425	-	No significant similarity	-	-

FE526426	-	No significant similarity	-	-
FE526427	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	7e-40
FE526428	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i> NRRL 181	1e-59
FE526429	-	No significant similarity	-	-
FE526430	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	1e-39
FE526431	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	3e-39
FE526432	gb AAR02424.1	subtilisin-like protease SUB5	<i>Trichophyton rubrum</i>	3e-47
FE526433	ref XP_448232.1	unnamed protein product	<i>Candida glabrata</i>	3e-04
FE526434	-	No significant similarity	-	-
FE526435	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	2e-74
FE526436	ref XP_001242709.1	hypothetical protein CIMG_06605	<i>Coccidioides immitis</i> RS	8e-44
FE526437	ref XP_754546.1	Ribosomal protein S7e	<i>Aspergillus fumigatus</i> Af293	9e-31
FE526438	-	No significant similarity	-	-
FE526439	emb CAE74488.1	Hypothetical protein CBG22239	<i>Caenorhabditis briggsae</i>	2e-12
FE526440	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	7e-21
FE526441	-	No significant similarity	-	-
FE526442	ref XP_001244025.1	hypothetical protein CIMG_03466	<i>Coccidioides immitis</i> RS	7e-31
FE526443	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	7e-73
FE526444	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	8e-30
FE526445	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	2e-38
FE526446	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	4e-22
FE526447	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	8e-40
FE526448	ref XP_661159.1	30 kD heat shock protein	<i>Aspergillus nidulans</i> FGSC A4	7e-07
FE526449	ref XP_001246883.1	hypothetical protein CIMG_00654	<i>Coccidioides immitis</i> RS	1e-03
FE526450	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	3e-42
FE526451	ref XP_001244025.1	hypothetical protein CIMG_03466	<i>Coccidioides immitis</i> RS	7e-31
FE526452	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	2e-21

FE526453	-	No significant similarity	-	-
FE526454	gb EEQ35176.1	urease accessory protein UreD	<i>Microsporum canis</i> CBS 113480	8e-66
FE526455	-	No significant similarity	-	-
FE526456	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	8e-22
FE526457	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	8e-22
FE526458	gb EEQ28888.1	MFS oligopeptide transporter, putative	<i>Microsporum canis</i> CBS 113480	2e-19
FE526459	-	No significant similarity	-	-
FE526460	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i> NRRL 181	2e-31
FE526461	ref NP_064450.2	formin 2	<i>Homo sapiens</i>	9e-06
FE526462	-	No significant similarity	-	-
FE526463	-	No significant similarity	-	-
FE526464	gb AAR11462.1	subtilisin-like protease SUB3	<i>Trichophyton rubrum</i>	5e-25
FE526465	-	No significant similarity	-	-
FE526466	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	5e-21
FE526467	gb AAR02424.1	subtilisin-like protease SUB5	<i>Trichophyton rubrum</i>	1e-40
FE526468	-	No significant similarity	-	-
FE526469	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	1e-35
FE526470	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	5e-21
FE526471	-	No significant similarity	-	-
FE526472	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	5e-21
FE526473	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	3e-42
FE526474	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	2e-65
FE526475	gb AAR02424.1	subtilisin-like protease SUB5	<i>Trichophyton rubrum</i>	3e-45
FE526476	ref XP_001244025.1	hypothetical protein CIMG_03466	<i>Coccidioides immitis</i> RS	2e-44
FE526477	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	5e-21
FE526478	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i> NRRL 181	9e-43
FE526479	gb AAR02424.1	subtilisin-like protease SUB5	<i>Trichophyton rubrum</i>	2e-45

FE526480	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	1e-16
FE526481	ref XP_001210481.1	protein MGM1, mitochondrial precursor	<i>Aspergillus terreus</i> NIH2624	3e-41
FE526482	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i> <i>verrucosum</i>	1e-44
FE526483	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	1e-44
FE526484	-	No significant similarity	-	-
FE526485	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	1e-28
FE526486	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	4e-20
FE526487	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i> NRRL 181	7e-67
FE526488	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i> <i>verrucosum</i>	2e-78
FE526489	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	2e-48
FE526490	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	6e-28
FE526491	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	1e-52
FE526492	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	8e-40
FE526493	gb AAR11462.1	subtilisin-like protease SUB3	<i>Trichophyton rubrum</i>	1e-24
FE526494	ref XP_001246302.1	hypothetical protein CIMG_00073	<i>Coccidioides immitis</i> RS	1e-25
FE526495	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	7e-39
FE526496	ref XP_001264535.1	MFS peptide transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-18
FE526497	gb EAX07174.1	hCG1793893	<i>Homo sapiens</i>	2e-05
FE526498	ref XP_747457.1	DUF895 domain membrane protein	<i>Aspergillus fumigatus</i> Af293	5e-26
FE526499	ref XP_001211350.1	sterol 24-C-methyltransferase	<i>Aspergillus terreus</i> NIH2624	7e-23
FE526500	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i> <i>verrucosum</i>	2e-48
FE526501	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	3e-19
FE526502	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i>	6e-29

FE526503	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>verrucosum</i> <i>Neosartorya fischeri</i> NRRL 181	3e-20
FE526504	ref XP_001276510.1	glucosamine-6-phosphate isomerase	<i>Aspergillus clavatus</i> NRRL 1	4e-37
FE526505	ref XP_747457.1	DUF895 domain membrane protein	<i>Aspergillus fumigatus</i> Af293	3e-11
FE526506	ref XP_970785.1	PREDICTED: similar to CG6957-PA, isoform A	<i>Tribolium castaneum</i>	3e-34
FE526507	ref XP_001261902.1	oligopeptide transporter	<i>Neosartorya fischeri</i> NRRL 181	2e-17
FE526508	ref XP_001271762.1	short chain dehydrogenase, putative	<i>Aspergillus clavatus</i> NRRL 1	1e-25
FE526509	ref XP_001261901.1	extracellular lipase, putative	<i>Neosartorya fischeri</i> NRRL 181	5e-19
FE526510	ref XP_001244025.1	hypothetical protein CIMG_03466	<i>Coccidioides immitis</i> RS	7e-08
FE526511	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i> <i>verrucosum</i>	7e-25
FE526512	ref XP_553256.2	ENSANGP00000027824	<i>Anopheles gambiae</i> str. PEST	6e-13
FE526513	emb CAK48124.1	unnamed protein product	<i>Aspergillus niger</i>	4e-12
FE526514	ref XP_001218693.1	cholinesterase	<i>Aspergillus terreus</i> NIH2624	2e-08
FE526515	gb EEQ29740.1	Aminoacid permease	<i>Microsporum canis</i>	1e-24
FE526516	ref XP_001244025.1	hypothetical protein CIMG_03466	<i>Coccidioides immitis</i> RS	1e-24
FE526517	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	6e-22
FE526518	-	No significant similarity	-	-
FE526519	ref XP_001245309.1	hypothetical protein CIMG_04750	<i>Coccidioides immitis</i> RS	1e-07
FE526520	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	1e-20
FE526521	ref XP_755153.1	flotillin domain protein	<i>Aspergillus fumigatus</i> Af293	4e-16
FE526522	ref XP_001246883.1	hypothetical protein CIMG_00654	<i>Coccidioides immitis</i> RS	3e-04
FE526523	-	No significant similarity	-	-
FE526524	ref XP_001246883.1	hypothetical protein CIMG_00654	<i>Coccidioides immitis</i> RS	3e-04
FE526525	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i> <i>verrucosum</i>	1e-45
FE526526	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i> <i>verrucosum</i>	1e-46

FE526527	gb AAR02424.1	subtilisin-like protease SUB5	<i>Trichophyton rubrum</i>	4e-47
FE526528	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	2e-47
FE526529	-	No significant similarity	-	-
FE526530	-	No significant similarity	-	-
FE526531	-	No significant similarity	-	-
FE526532	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	1e-40
FE526533	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	2e-48
FE526534	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	1e-45
FE526535	-	No significant similarity	-	-
FE526536	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	5e-46
FE526537	-	No significant similarity	-	-
FE526538	-	No significant similarity	-	-
FE526539	-	No significant similarity	-	-
FE526540	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	1e-34
FE526541	gb AAR08135.1	small GTPase RanA	<i>Emericella nidulans</i>	2e-40
FE526542	-	No significant similarity	-	-
FE526543	ref XP_001208665.1	vacuolar ATP synthase subunit B	<i>Aspergillus terreus</i> NIH2624	2e-17
FE526544	-	No significant similarity	-	-
FE526545	-	No significant similarity	-	-
FE526546	-	No significant similarity	-	-
FE526547	ref XP_001275026.1	oligopeptide transporter	<i>Aspergillus clavatus</i> NRRL 1	7e-37
FE526548	ref XP_001244970.1	glucosamine-6-phosphate deaminase	<i>Coccidioides immitis</i> RS	8e-61
FE526549	-	No significant similarity	-	-
FE526550	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton verrucosum</i>	1e-44
FE526551	-	No significant similarity	-	-
FE526552	ref XP_747457.1	DUF895 domain membrane protein	<i>Aspergillus fumigatus</i> Af293	4e-18
FE526553	gb ABL84992.1	metalloprotease Mep4	<i>Trichophyton tonsurans</i>	4e-23
FE526554	-	No significant similarity	-	-

FE526555	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i> NRRL 1	9e-09
FE526556	ref XP_754782.1	hypothetical protein Afu3g08320	<i>Aspergillus fumigatus</i> Af293	2e-05
FE526557	ref XP_001244184.1	hypothetical protein CIMG_03625	<i>Coccidioides immitis</i> RS	9e-49
FE526558	ref XP_001209001.1	predicted protein	<i>Aspergillus terreus</i> NIH2624	2e-05
FE526559	-	No significant similarity	-	-
FE526560	-	No significant similarity	-	-
FE526561	ref XP_001244184.1	hypothetical protein CIMG_03625	<i>Coccidioides immitis</i> RS	1e-51
FE526562	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE526563	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	4e-15
FE526564	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i> NRRL 1	2e-05
FE526565	ref YP_840033.1	Glyoxalase/bleomycin resistance protein/dioxygenase	<i>Burkholderia</i> <i>cenocepacia</i> HI2424	4e-11
FE526566	ref YP_840033.1	Glyoxalase/bleomycin resistance protein/dioxygenase	<i>Burkholderia</i> <i>cenocepacia</i> HI2424	3e-11
FE526567	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08
FE526568	gb EEQ33141.1	NIMA interactive protein	<i>Microsporum canis</i> CBS 113480	2e-23
FE526569	-	No significant similarity	-	-
FE526570	-	No significant similarity	-	4e-05
FE526571	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526572	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	8e-07
FE526573	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526574	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	3e-22
FE526575	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	9e-27
FE526576	-	No significant similarity	-	-
FE526577	ref YP_840033.1	Glyoxalase/bleomycin resistance protein/dioxygenase	<i>Burkholderia</i> <i>cenocepacia</i> HI2424	3e-11
FE526578	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526579	-	No significant similarity	-	-
FE526580	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-18
FE526581	ref ZP_01556758.1	aldehyde dehydrogenase	<i>Burkholderia ambifaria</i> MC40-6	5e-65
FE526582	ref YP_371113.1	Aldehyde dehydrogenase	<i>Burkholderia</i> sp. 383	7e-78
FE526583	ref XP_001242594.1	hypothetical protein CIMG_06490	<i>Coccidioides immitis</i> RS	6e-08

FE526584	-	No significant similarity	-	-
FE526585	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526586	-	No significant similarity	-	-
FE526587	-	No significant similarity	-	-
FE526588	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526589	-	No significant similarity	-	-
FE526590	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	3e-27
FE526591	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-18
FE526592	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	7e-10
FE526593	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	2e-28
FE526594	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-18
FE526595	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	4e-09
			<i>NRRL 1</i>	
FE526596	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	7e-27
FE526597	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	2e-16
FE526598	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	2e-14
FE526599	-	No significant similarity	-	-
FE526600	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	5e-42
FE526601	gb AAP78735.1	nonribosomal peptide synthase	<i>Alternaria brassicae</i>	5e-06
FE526602	ref XP_001244184.1	hypothetical protein CIMG_03625	<i>Coccidioides immitis</i> RS	7e-53
FE526603	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	2e-23
FE526604	-	No significant similarity	-	-
FE526605	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	2e-06
			<i>NRRL 1</i>	
FE526606	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	8e-26
FE526607	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	3e-11
FE526608	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	4e-38
FE526609	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-15
FE526610	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	6e-38
FE526611	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	4e-43
FE526612	emb CAK37677.1	unnamed protein product	<i>Aspergillus niger</i>	6e-46
FE526613	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	6e-06
FE526614	-	No significant similarity	-	2e-04
FE526615	ref XP_001244184.1	hypothetical protein CIMG_03625	<i>Coccidioides immitis</i> RS	8e-36
FE526616	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-26
FE526617	-	No significant similarity	-	-
FE526618	-	No significant similarity	-	-
FE526619	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	6e-20

FE526620	ref XP_001239812.1	hypothetical protein CIMG_09433	<i>Coccidioides immitis</i> RS	4e-08
FE526621	ref ZP_01420399.1	O-antigen polymerase	<i>Caulobacter</i> sp. K31	4e-17
FE526622	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	5e-43
FE526623	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526624	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	4e-20
FE526625	-	No significant similarity	-	-
FE526626	-	No significant similarity	-	-
FE526627	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-18
FE526628	gb AAP78735.1	nonribosomal peptide synthase	<i>Alternaria brassicae</i>	5e-06
FE526629	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	1e-07
			NRRL 1	
FE526630	gb AAP78735.1	nonribosomal peptide synthase	<i>Alternaria brassicae</i>	5e-06
FE526631	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	2e-24
FE526632	-	No significant similarity	-	-
FE526633	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	6e-25
FE526634	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-22
FE526635	ref YP_371113.1	Aldehyde dehydrogenase	<i>Burkholderia</i> sp. 383	3e-75
FE526636	-	No significant similarity	-	-
FE526637	-	No significant similarity	-	-
FE526638	ref XP_001244184.1	hypothetical protein CIMG_03625	<i>Coccidioides immitis</i> RS	3e-49
FE526639	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	7e-20
FE526640	-	No significant similarity	-	-
FE526641	ref XP_001244184.1	hypothetical protein CIMG_03625	<i>Coccidioides immitis</i> RS	2e-48
FE526642	ref XP_001264989.1	leucine rich repeat protein	<i>Neosartorya fischeri</i>	3e-09
			NRRL 181	
FE526643	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-17
FE526644	gb ABF22673.1	copper-sulfate regulated protein 1	<i>Ajellomyces capsulatus</i>	7e-10
FE526645	-	No significant similarity	-	-
FE526646	ref XP_001269482.1	Leucine Rich Repeat domain protein	<i>Aspergillus clavatus</i>	3e-04
			NRRL 1	
FE526647	gb AAL26311.2 AF362957_1	polyprotein	<i>Aspergillus flavus</i>	2e-31
FE526648	-	No significant similarity	-	-
FE526649	-	No significant similarity	-	-
FE526650	-	No significant similarity	-	-
FE526651	-	No significant similarity	-	-
FE526652	-	No significant similarity	-	-
FE526653	ref YP_371113.1	Aldehyde dehydrogenase	<i>Burkholderia</i> sp. 383	3e-77
FE526654	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	1e-22

FE526655	-	No significant similarity	-	-
FE526656	ref XP_001264989.1	leucine rich repeat protein	<i>Neosartorya fischeri</i> NRRL 181	1e-04
FE526657	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	7e-25
FE526658	ref XP_001240454.1	hypothetical protein CIMG_07617	<i>Coccidioides immitis</i> RS	1e-05
FE526659	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	2e-19
FE526660	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	6e-23
FE526661	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	2e-33
FE526662	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	5e-28
FE526663	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	4e-16
FE526664	gb AAR29046.2	gag-pol polyprotein	<i>Aspergillus flavus</i>	1e-28
FE526665	-	No significant similarity	-	-
FE526666	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	1e-33
FE526667	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	1e-22
FE526668	gb AAG24792.1 AF264028_2	pol protein	<i>Glomerella cingulata</i>	4e-42
FE526669	-	No significant similarity	-	-
FE526670	-	No significant similarity	-	-
FE526671	ref XP_001226392.1	hypothetical protein CHGG_08465	<i>Chaetomium globosum</i> CBS 148.51	4e-10
FE526672	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	2e-23
FE526673	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	2e-16
FE526674	gb AAG01549.3 AF291822_1	multidrug resistance protein MDR	<i>Trichophyton rubrum</i>	7e-20
FE526675	gb AAP23304.1	NIMA interactive protein	<i>Emericella nidulans</i>	5e-25
FE526676	-	No significant similarity	-	-
FE526677	ref YP_443455.1	secretion protein, putative	<i>Burkholderia</i> <i>thailandensis</i> E264	3e-04
FE526678	-	No significant similarity	-	-
FE526679	-	No significant similarity	-	-
FE526680	-	No significant similarity	-	-
FE526681	gb AAN62296.1 AF440524_83	transposase domain protein	<i>Pseudomonas</i> <i>aeruginosa</i>	8e-22
FE526682	-	No significant similarity	-	-
FE526683	ref XP_001262797.1	FYVE zinc finger protein	<i>Neosartorya fischeri</i> NRRL 181	2e-09
FE526684	ref XP_001262797.1	FYVE zinc finger protein	<i>Neosartorya fischeri</i> NRRL 181	4e-26
FE526685	ref XP_001246883.1	hypothetical protein CIMG_00654	<i>Coccidioides immitis</i> RS	3e-04
FE526686	-	No significant similarity	-	-

FE526687	-	No significant similarity	-	-
FE526688	-	No significant similarity	-	-
FE526689	-	No significant similarity	-	-
FE526690	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	5e-21
FE526691	gb AAN62296.1 AF440524_83	transposase domain protein	<i>Pseudomonas</i> <i>aeruginosa</i>	2e-22
FE526692	ref XP_001262797.1	FYVE zinc finger protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	4e-12
FE526693	-	No significant similarity	-	-
FE526694	-	No significant similarity	-	-
FE526695	-	No significant similarity	-	-
FE526696	-	No significant similarity	-	-
FE526697	ref XP_001262797.1	FYVE zinc finger protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-27
FE526698	ref XP_001262797.1	FYVE zinc finger protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	2e-27
FE526699	-	No significant similarity	-	-
FE526700	-	No significant similarity	-	-
FE526701	ref XP_001264529.1	GABA permease, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	3e-17
FE526702	ref XP_001264853.1	amino acid permease, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	3e-26
FE526703	ref XP_001262797.1	FYVE zinc finger protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	6e-26
FE526704	-	No significant similarity	-	-
FE526705	-	No significant similarity	-	-
FE526706	-	No significant similarity	-	-
FE526707	ref XP_001263899.1	60S ribosomal protein L20, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	5e-11
FE526708	-	No significant similarity	-	-
FE526709	ref XP_001262797.1	FYVE zinc finger protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	2e-22
FE526710	ref XP_001262797.1	FYVE zinc finger protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	8e-13
FE526711	-	No significant similarity	-	-
FE526712	ref YP_443455.1	secretion protein, putative	<i>Burkholderia</i> <i>thailandensis E264</i>	3e-04

FE526713	-	No significant similarity	-	-
FE526714	-	No significant similarity	-	-
FE526715	-	No significant similarity	-	-
FE526716	-	No significant similarity	-	-
FE526717	gb AAN62296.1 AF440524_83	transposase domain protein	<i>Pseudomonas aeruginosa</i>	2e-22
FE526718	gb AAF21678.1 AF051915_2	pol polyprotein	<i>Cladosporium fulvum</i>	1e-09
FE526719	ref XP_001263899.1	60S ribosomal protein L20, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	5e-11
FE526720	-	No significant similarity	-	-
FE526721	-	No significant similarity	-	-
FE526722	-	No significant similarity	-	-
FE526723	-	No significant similarity	-	-
FE526724	ref XP_001263899.1	60S ribosomal protein L20, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	5e-11
FE526725	-	No significant similarity	-	-
FE526726	ref XP_001260782.1	V-type ATPase, B subunit, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	5e-21
FE526727	-	No significant similarity	-	-
FE526728	-	No significant similarity	-	-
FE526729	-	No significant similarity	-	-
FE526730	ref YP_443455.1	secretion protein, putative	<i>Burkholderia thailandensis</i> E264	4e-04
FE526731	-	No significant similarity	-	-
FE526732	ref XP_001246883.1	hypothetical protein CIMG_00654	<i>Coccidioides immitis</i> RS	3e-04
FE526733	-	No significant similarity	-	-
FE526734	-	No significant similarity	-	-
FE526735	-	No significant similarity	-	-
FE526736	-	No significant similarity	-	-
FE526737	-	No significant similarity	-	-
FE526738	ref XP_001246883.1	hypothetical protein CIMG_00654	<i>Coccidioides immitis</i> RS	3e-04
FE526739	-	No significant similarity	-	-
FE526740	ref XP_001246883.1	hypothetical protein CIMG_00654	<i>Coccidioides immitis</i> RS	3e-04
FE526741	gb EEQ32880.1	FYVE zinc finger protein	<i>Microsporum canis</i> CBS 113480	1e-34
FE526742	-	No significant similarity	-	-
FE526743	-	No significant similarity	-	-
FE526744	-	No significant similarity	-	-

FE526745	-	No significant similarity	-	-
FE526746	-	No significant similarity	-	-
FE526747	-	No significant similarity	-	-
FE526748	ref XP_001262797.1	FYVE zinc finger protein	<i>Neosartorya fischeri</i> NRRL 181	9e-14
FE526749	ref XP_001272385.1	FYVE zinc finger protein	<i>Aspergillus clavatus</i> NRRL 1	3e-04
FE526750	-	No significant similarity	-	-
FE526751	gb AAN62296.1 AF440524_83	transposase domain protein	<i>Pseudomonas</i> <i>aeruginosa</i>	2e-22
FE526752	-	No significant similarity	-	-
FE526753	-	No significant similarity	-	-
FE526754	-	No significant similarity	-	-
FE526755	-	No significant similarity	-	-
FE526756	-	No significant similarity	-	-
FE526757	ref XP_001242709.1	hypothetical protein CIMG_06605	<i>Coccidioides immitis</i> RS	3e-10
FE526758	ref XP_753997.1	60s ribosomal protein y16	<i>Aspergillus fumigatus</i> Af293	1e-29
FE526759	-	No significant similarity	-	-
FE526760	ref XP_753233.1	ThiJ/PfpI family protein	<i>Aspergillus fumigatus</i> Af293	9e-07
FE526761	-	No significant similarity	-	-
FE526762	ref XP_001243030.1	hypothetical protein CIMG_06926	<i>Coccidioides immitis</i> RS	2e-06
FE526763	sp Q462Q7 COX19_PARBR	Cytochrome c oxidase assembly protein COX19 gb AAY64182.1 Cox19p	<i>Paracoccidioides</i> <i>brasiliensis</i>	1e-12
FE526764	ref XP_001259622.1	pentatricopeptide repeat protein	<i>Neosartorya fischeri</i> NRRL 181	1e-20
FE526765	-	No significant similarity	-	-
FE526766	-	No significant similarity	-	-
FE526767	-	No significant similarity	-	-
FE526768	-	No significant similarity	-	-
FE526769	ref XP_001259414.1	50S ribosomal protein L4	<i>Neosartorya fischeri</i> NRRL 181	7e-05
FE526770	-	No significant similarity	-	-
FE526771	gb AAG36933.1 AF262955_1	oleate delta-12 desaturase	<i>Emericella nidulans</i>	9e-21
FE526772	ref XP_001248270.1	hypothetical protein CIMG_02041	<i>Coccidioides immitis</i> RS	1e-04
FE526773	ref XP_001212165.1	HNRNP arginine N-methyltransferase	<i>Aspergillus terreus</i> NIH2624	3e-16

FE526774	-	No significant similarity	-	-
FE526775	ref XP_001269624.1	ADP,ATP carrier protein	<i>Aspergillus clavatus</i> NRRL 1	2e-23
FE526776	-	No significant similarity	-	-
FE526777	ref XP_001268371.1	CECR1 family adenosine deaminase, putative	<i>Aspergillus clavatus</i> NRRL 1	4e-14
FE526778	-	No significant similarity	-	-
FE526779	ref XP_001260012.1	conserved hypothetical protein	<i>Neosartorya fischeri</i> NRRL 181	3e-07
FE526780	ref XP_001257300.1	monocarboxylate permease homologue, mch4	<i>Neosartorya fischeri</i> NRRL 181	3e-08
FE526781	-	No significant similarity	-	-
FE526782	ref XP_001260419.1	60S ribosomal protein L10	<i>Neosartorya fischeri</i> NRRL 181	1e-34
FE526783	ref XP_001275190.1	amino acid transporter, putative	<i>Aspergillus clavatus</i> NRRL 1	4e-13
FE526784	-	No significant similarity	-	-
FE526785	-	No significant similarity	-	-
FE526786	ref XP_001245093.1	hypothetical protein CIMG_04534	<i>Coccidioides immitis</i> RS	3e-07
FE526787	ref XP_726427.1	RNA 3l-terminal phosphate cyclase protein	<i>Plasmodium yoelii yoelii</i> str. 17XNL	7e-04
FE526788	ref XP_001272674.1	taurine catabolism dioxygenase TauD, TfdA family protein	<i>Aspergillus clavatus</i> NRRL 1	1e-24
FE526789	-	No significant similarity	-	-
FE526790	ref XP_001257403.1	benzoate 4-monooxygenase cytochrome P450	<i>Neosartorya fischeri</i> NRRL 181	1e-12
FE526791	ref XP_001267237.1	60S ribosomal protein L38, putative	<i>Neosartorya fischeri</i> NRRL 181	5e-14
FE526792	ref XP_001273971.1	60S ribosomal protein L8, putative	<i>Aspergillus clavatus</i> NRRL 1	9e-23
FE526793	-	No significant similarity	-	-
FE526794	ref XP_001246583.1	succinate-semialdehyde dehydrogenase	<i>Coccidioides immitis</i> RS	3e-27
FE526795	ref XP_001267237.1	60S ribosomal protein L38, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-14
FE526796	ref XP_001272771.1	40S ribosomal protein S11	<i>Aspergillus clavatus</i> NRRL 1	2e-06
FE526797	-	No significant similarity	-	-
FE526798	ref XP_001267248.1	60S ribosomal protein L11	<i>Neosartorya fischeri</i>	2e-19

FE526799	-	No significant similarity	NRRL 181	-
FE526800	-	No significant similarity	-	-
FE526801	-	No significant similarity	-	-
FE526802	ref XP_001213404.1	predicted protein	<i>Aspergillus terreus</i> NIH2624	4e-15
FE526803	-	No significant similarity	-	-
FE526804	-	No significant similarity	-	-
FE526805	ref XP_001267237.1	60S ribosomal protein L38, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-13
FE526806	gb ABH10644.1	aconitase	<i>Coccidioides posadasii</i>	2e-42
FE526807	-	No significant similarity	-	-
FE526808	gb EAT82743.1	predicted protein	<i>Phaeosphaeria nodorum</i> SN15	6e-04
FE526809	-	No significant similarity	-	-
FE526810	ref XP_001267762.1	glutathione S-transferase, putative	<i>Aspergillus clavatus</i> NRRL 1	1e-23
FE526811	-	No significant similarity	-	-
FE526812	-	No significant similarity	-	-
FE526813	ref XP_001248510.1	alanyl-tRNA synthetase, cytoplasmic	<i>Coccidioides immitis</i> RS	1e-17
FE526814	-	No significant similarity	-	-
FE526815	ref XP_001243359.1	hypothetical protein CIMG_07255	<i>Coccidioides immitis</i> RS	7e-04
FE526816	gb EDJ95143.1	60S ribosomal protein L7	<i>Magnaporthe grisea</i> 70-15	1e-09
FE526817	ref XP_001216898.1	ADP-ribose pyrophosphatase	<i>Aspergillus terreus</i> NIH2624	2e-22
FE526818	ref XP_001260458.1	SNARE protein (Ufe1), putative	<i>Neosartorya fischeri</i> NRRL 181	5e-09
FE526819	-	No significant similarity	-	-
FE526820	ref XP_001212923.1	60S ribosomal protein L20	<i>Aspergillus terreus</i> NIH2624	2e-05
FE526821	ref XP_001213571.1	mitochondrial genome maintenance protein MGM101	<i>Aspergillus terreus</i> NIH2624	9e-48
FE526822	ref XP_001260097.1	cytochrome b5, putative	<i>Neosartorya fischeri</i> NRRL 181	6e-16
FE526823	ref XP_001215797.1	multicopy enhancer of UAS2	<i>Aspergillus terreus</i> NIH2624	5e-50
FE526824	sp P34763 NMT_AJECA	Glycylpeptide N-tetradecanoyltransferase (Peptide N-myristoyltransferase)	<i>Ajellomyces capsulatus</i>	6e-24

		(Myristoyl-CoA:protein N-myristoyltransferase) (NMT) gb AAA17549.1		
		N-myristoyltransferase		
FE526825	ref XP_001269377.1	DUF814 domain protein	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-08
FE526826	ref XP_001247536.1	lysyl-tRNA synthetase, cytoplasmic	<i>Coccidioides immitis</i> RS	4e-19
FE526827	gb EAL91735.2	integral membrane protein	<i>Aspergillus fumigatus</i> <i>Af293</i>	4e-23
FE526828	ref XP_001240306.1	cytochrome P450 51	<i>Coccidioides immitis</i> RS	1e-63
FE526829	-	No significant similarity	-	-
FE526830	ref XP_752041.1	Ribosomal L38e protein family	<i>Aspergillus fumigatus</i> <i>Af293</i>	5e-14
FE526831	ref XP_001244411.1	hypothetical protein CIMG_03852	<i>Coccidioides immitis</i> RS	6e-04
FE526832	-	No significant similarity	-	-
FE526833	-	No significant similarity	-	-
FE526834	ref XP_001259852.1	WD repeat protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	2e-22
FE526835	gb EAL90446.2	eukaryotic translation elongation factor 1 subunit Eef1-beta, putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	2e-13
FE526836	-	No significant similarity	-	-
FE526837	ref XP_001273261.1	tropomyosin, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	6e-35
FE526838	-	No significant similarity	-	-
FE526839	-	No significant similarity	-	-
FE526840	-	No significant similarity	-	-
FE526841	ref XP_001260419.1	60S ribosomal protein L10	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	7e-37
FE526842	-	No significant similarity	-	-
FE526843	-	No significant similarity	-	-
FE526844	ref XP_751502.1	HAD superfamily hydrolase, putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	3e-39
FE526845	emb CAK97343.1	unnamed protein product	<i>Aspergillus niger</i>	4e-13
FE526846	-	No significant similarity	-	-
FE526847	ref XP_001270525.1	conserved hypothetical protein	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	9e-07
FE526848	gb EAL90446.2	eukaryotic translation elongation factor 1 subunit Eef1-beta, putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	4e-13
FE526849	ref XP_753997.1	60s ribosomal protein y16	<i>Aspergillus fumigatus</i> <i>Af293</i>	5e-30

FE526850	-	No significant similarity	-	-
FE526851	ref XP_001274066.1	60S ribosome biogenesis protein Sqt1, putative	<i>Aspergillus clavatus</i> NRRL 1	7e-21
FE526852	ref XP_001214720.1	outer mitochondrial membrane protein porin	<i>Aspergillus terreus</i> NIH2624	4e-19
FE526853	ref XP_001257403.1	benzoate 4-monooxygenase cytochrome P450	<i>Neosartorya fischeri</i> NRRL 181	1e-13
FE526854	-	No significant similarity	-	-
FE526855	sp P07509 SODC_NEUCR	Superoxide dismutase	<i>Neurospora crassa</i>	2e-13
FE526856	gb ABN50029.1	putative senescence-associated protein	<i>Trichosanthes dioica</i>	2e-05
FE526857	ref XP_001268698.1	succinate dehydrogenase subunit CybS, putative	<i>Aspergillus clavatus</i> NRRL 1	4e-07
FE526858	ref XP_001211720.1	60S ribosomal protein L12	<i>Aspergillus terreus</i> NIH2624	9e-10
FE526859	-	No significant similarity	-	-
FE526860	-	No significant similarity	-	-
FE526861	-	No significant similarity	-	-
FE526862	-	No significant similarity	-	-
FE526863	-	No significant similarity	-	-
FE526864	ref XP_001271235.1	DUF1713 domain protein	<i>Aspergillus clavatus</i> NRRL 1	3e-07
FE526865	ref XP_001246620.1	40S ribosomal protein S4	<i>Coccidioides immitis</i> RS	1e-26
FE526866	ref XP_001262838.1	indoleamine 2,3-dioxygenase family protein	<i>Neosartorya fischeri</i> NRRL 181	1e-47
FE526867	ref XP_001387342.1	60S large subunit ribosomal protein	<i>Pichia stipitis</i> CBS 6054	4e-04
FE526868	emb CAK45164.1	protein O-mannosyl transferase pmtA- <i>Aspergillus niger</i>	<i>Aspergillus niger</i>	1e-27
FE526869	ref XP_753997.1	60s ribosomal protein y16	<i>Aspergillus fumigatus</i> Af293	2e-17
FE526870	gb EAL85168.2	U5 snRNP component Snu114, putative	<i>Aspergillus fumigatus</i> Af293	1e-18
FE526871	ref XP_001246620.1	40S ribosomal protein S4	<i>Coccidioides immitis</i> RS	1e-26
FE526872	ref XP_001269346.1	60S ribosomal protein L9, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-20
FE526873	ref XP_001243698.1	fibrillarin	<i>Coccidioides immitis</i> RS	7e-50
FE526874	ref XP_001264284.1	kelch repeat protein	<i>Neosartorya fischeri</i> NRRL 181	6e-15
FE526875	-	No significant similarity	-	-
FE526876	-	No significant similarity	-	-

FE526877	-	No significant similarity	-	-
FE526878	ref XP_001246583.1	succinate-semialdehyde dehydrogenase	<i>Coccidioides immitis</i> RS	3e-27
FE526879	ref XP_001265211.1	Ribosomal L18ae protein family	<i>Neosartorya fischeri</i> NRRL 181	6e-16
FE526880	-	No significant similarity	-	-
FE526881	-	No significant similarity	-	-
FE526882	ref XP_001260103.1	WD repeat protein	<i>Neosartorya fischeri</i> NRRL 181	3e-05
FE526883	gb EDJ98154.1	GTP cyclohydrolase II, putative	<i>Magnaporthe grisea</i> 70-15	3e-50
FE526884	gb EEQ30276.1	mitochondrial 2-methylisocitrate lyase	<i>Microsporum canis</i> CBS113480	9e-31
FE526885	-	No significant similarity	-	-
FE526886	ref XP_661668.1	ADP/ATP carrier protein	<i>Aspergillus nidulans</i> FGSC A4	2e-39
FE526887	ref XP_001178408.1	PREDICTED: similar to translation initiation factor 2 gamma subunit	<i>Strongylocentrotus purpuratus</i>	8e-17
FE526888	ref XP_001270945.1	PDCD2_C domain protein, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-08
FE526889	ref XP_001244571.1	40S ribosomal protein S26E	<i>Coccidioides immitis</i> RS	4e-10
FE526890	-	No significant similarity	-	-
FE526891	-	No significant similarity	-	-
FE526892	-	No significant similarity	-	-
FE526893	-	No significant similarity	-	-
FE526894	-	No significant similarity	-	-
FE526895	-	No significant similarity	-	-
FE526896	-	No significant similarity	-	-
FE526897	emb CAD29478.1	glutathione transferase F5	<i>Triticum aestivum</i>	7e-07
FE526898	-	No significant similarity	-	-
FE526899	gb AAM54368.1	elongation factor 1-alpha	<i>Trichophyton rubrum</i>	7e-20
FE526900	-	No significant similarity	-	-
FE526901	ref XP_661668.1	ADP/ATP carrier protein	<i>Aspergillus nidulans</i> FGSC A4	1e-40
FE526902	sp Q5BGF9 PAM17_EMENI	Presequence translocated-associated motor subunit pam17, mitochondrial precursor	<i>Emericella nidulans</i>	8e-22
FE526903	-	No significant similarity	-	-
FE526904	-	No significant similarity	-	-
FE526905	gb AAS76666.1	carboxypeptidase S1	<i>Trichophyton rubrum</i>	5e-48

FE526906	sp Q5AZJ7 DPH1_EMENI	Diphthamide biosynthesis protein	<i>Emericella nidulans</i>	3e-36
FE526907	ref XP_001273971.1	60S ribosomal protein L8, putative	<i>Aspergillus clavatus</i>	2e-12
			NRRL 1	
FE526908	ref XP_001246620.1	40S ribosomal protein S4	<i>Coccidioides immitis</i> RS	6e-28
FE526909	ref XP_001266080.1	seryl-tRNA synthetase	<i>Neosartorya fischeri</i>	4e-26
			NRRL 181	
FE526910	ref XP_001240451.1	aspartyl aminopeptidase, putative	<i>Coccidioides immitis</i> RS	5e-47
FE526911	-	No significant similarity	-	-
FE526912	-	No significant similarity	-	-
FE526913	-	No significant similarity	-	-
FE526914	-	No significant similarity	-	-
FE526915	-	No significant similarity	-	-
FE526916	ref XP_001244035.1	hypothetical protein CIMG_03476	<i>Coccidioides immitis</i> RS	2e-12
FE526917	ref XP_001246620.1	40S ribosomal protein S4	<i>Coccidioides immitis</i> RS	6e-28
FE526918	-	No significant similarity	-	-
FE526919	gb EAL90446.2	eukaryotic translation elongation factor 1 subunit Eef1-beta, putative	<i>Aspergillus fumigatus</i>	6e-19
			Af293	
FE526920	emb CAC18218.1	probable TRANSKETOLASE	<i>Neurospora crassa</i>	6e-22
FE526921	-	No significant similarity	-	-
FE526922	gb ABH11414.1	peroxin 3	<i>Penicillium</i>	1e-07
			<i>chrysogenum</i>	
FE526923	-	No significant similarity	-	-
FE526924	ref XP_567138.1	siderochrome-iron (ferrioxamine) uptake transporter	<i>Cryptococcus</i>	5e-21
			<i>neoformans</i> var.	
			<i>neoformans</i> JEC21	
FE526925	ref XP_001258439.1	60S ribosomal protein L7a	<i>Neosartorya fischeri</i>	1e-08
			NRRL 181	
FE526926	gb ABH10638.1	enolase	<i>Coccidioides posadasii</i>	5e-29
FE526927	ref XP_001261926.1	ELMO/CED-12 family protein	<i>Neosartorya fischeri</i>	1e-28
			NRRL 181	
FE526928	-	No significant similarity	-	-
FE526929	-	No significant similarity	-	-
FE526930	-	No significant similarity	-	-
FE526931	ref XP_001241489.1	hypothetical protein CIMG_08652	<i>Coccidioides immitis</i> RS	3e-23
FE526932	ref XP_001248308.1	60S ribosomal protein L28	<i>Coccidioides immitis</i> RS	4e-22
FE526933	ref XP_001241872.1	hypothetical protein CIMG_05768	<i>Coccidioides immitis</i> RS	2e-05
FE526934	ref XP_753437.1	integral membrane protein, putative	<i>Aspergillus fumigatus</i>	5e-05
			Af293	

FE526935	-	No significant similarity	-	-
FE526936	ref XP_001248510.1	alanyl-tRNA synthetase, cytoplasmic	<i>Coccidioides immitis</i> RS	6e-17
FE526937	ref XP_001267248.1	60S ribosomal protein L11	<i>Neosartorya fischeri</i>	4e-21
			NRRL 181	
FE526938	ref XP_001266988.1	phosphatase-like protein (PTPLA), putative	<i>Neosartorya fischeri</i>	7e-13
			NRRL 181	
FE526939	ref XP_001220066.1	40S ribosomal protein S11	<i>Chaetomium globosum</i>	3e-04
			CBS 148.51	
FE526940	-	No significant similarity	-	-
FE526941	-	No significant similarity	-	-
FE526942	-	No significant similarity	-	-
FE526943	ref XP_001270945.1	PDCD2_C domain protein, putative	<i>Aspergillus clavatus</i>	3e-07
			NRRL 1	
FE526944	gb EAL91735.2	integral membrane protein	<i>Aspergillus fumigatus</i>	2e-24
			Af293	
FE526945	ref XP_001247826.1	2-oxoglutarate dehydrogenase E1 component	<i>Coccidioides immitis</i> RS	3e-15
FE526946	-	No significant similarity	-	-
FE526947	-	No significant similarity	-	-
FE526948	gb EDJ95143.1	60S ribosomal protein L7	<i>Magnaporthe grisea</i> 70-15	1e-12
FE526949	-	No significant similarity	-	-
FE526950	-	No significant similarity	-	-
FE526951	-	No significant similarity	-	-
FE526952	ref XP_001269007.1	oleate delta-12 desaturase	<i>Aspergillus clavatus</i>	7e-13
			NRRL 1	
FE526953	ref XP_001273282.1	mitochondrial ribosomal protein DAP3, putative	<i>Aspergillus clavatus</i>	2e-12
			NRRL 1	
FE526954	ref XP_001246620.1	40S ribosomal protein S4	<i>Coccidioides immitis</i> RS	6e-28
FE526955	ref XP_001265779.1	fatty acid desaturase, putative	<i>Neosartorya fischeri</i>	2e-28
			NRRL 181	
FE526956	ref XP_001247300.1	hypothetical protein CIMG_01071	<i>Coccidioides immitis</i> RS	1e-05
FE526957	ref XP_001267424.1	phosphoethanolamine	<i>Neosartorya fischeri</i>	8e-22
			NRRL 181	
FE526958	ref XP_001242955.1	hypothetical protein CIMG_06851	<i>Coccidioides immitis</i> RS	1e-08
FE526959	-	No significant similarity	-	-
FE526960	gb EDJ95143.1	60S ribosomal protein L7	<i>Magnaporthe grisea</i> 70-15	8e-14
FE526961	-	No significant similarity	-	-

FE526962	ref XP_001259336.1	cell cycle control protein (Cwf8), putative	<i>Neosartorya fischeri</i> NRRL 181	1e-27
FE526963	ref XP_001210588.1	U3 small nucleolar ribonucleoprotein protein IMP4	<i>Aspergillus terreus</i> NIH2624	5e-43
FE526964	ref XP_001261302.1	C2 domain protein	<i>Neosartorya fischeri</i> NRRL 181	5e-29
FE526965	ref XP_001212323.1	predicted protein	<i>Aspergillus terreus</i> NIH2624	8e-09
FE526966	ref XP_001267850.1	antigenic mitochondrial protein HSP60, putative	<i>Aspergillus clavatus</i> NRRL 1	5e-13
FE526967	ref XP_001244490.1	ADP,ATP carrier protein	<i>Coccidioides immitis</i> RS	8e-19
FE526968	ref XP_001248397.1	hypothetical protein CIMG_02168	<i>Coccidioides immitis</i> RS	2e-05
FE526969	gb AAL50803.1 AF452883_1	Y20 protein	<i>Paracoccidioides</i> <i>brasiliensis</i>	6e-30
FE526970	-	No significant similarity	-	-
FE526971	-	No significant similarity	-	-
FE526972	pir S63701	mannosyl-oligosaccharide 1,2-alpha-mannosidase (EC 3.2.1.113) precursor	<i>Aspergillus phoenicis</i>	2e-13
FE526973	ref XP_001242456.1	hypothetical protein CIMG_06352	<i>Coccidioides immitis</i> RS	1e-04
FE526974	ref XP_001242392.1	plasma membrane ATPase	<i>Coccidioides immitis</i> RS	3e-43
FE526975	gb AAT40563.1	small G-protein GPA3	<i>Paracoccidioides</i> <i>brasiliensis</i>	2e-29
FE526976	ref XP_001271126.1	ubiquitin	<i>Aspergillus clavatus</i> NRRL 1	3e-23
FE526977	ref XP_001266080.1	seryl-tRNA synthetase	<i>Neosartorya fischeri</i> NRRL 181	3e-25
FE526978	-	No significant similarity	-	-
FE526979	emb CAM37639.1	proteophosphoglycan ppg1	<i>Leishmania braziliensis</i>	1e-23
FE526980	ref XP_001269939.1	eukaryotic translation initiation factor 3 subunit EifCb, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-36
FE526981	ref XP_001248308.1	60S ribosomal protein L28	<i>Coccidioides immitis</i> RS	8e-24
FE526982	-	No significant similarity	-	-
FE526983	ref XP_001248096.1	acetamidase	<i>Coccidioides immitis</i> RS	6e-33
FE526984	-	No significant similarity	-	-
FE526985	ref XP_001241489.1	hypothetical protein CIMG_08652	<i>Coccidioides immitis</i> RS	7e-26
FE526986	-	No significant similarity	-	-
FE526987	-	No significant similarity	-	-
FE526988	-	No significant similarity	-	-
FE526989	ref XP_001257403.1	benzoate 4-monooxygenase cytochrome P450	<i>Neosartorya fischeri</i>	1e-13

			<i>NRRL 181</i>	
FE526990	-	No significant similarity	-	-
FE526991	-	No significant similarity	-	-
FE526992	-	No significant similarity	-	-
FE526993	-	No significant similarity	-	-
FE526994	ref XP_001386580.1	membrane protein involved in vacuolar protein sorting	<i>Pichia stipitis</i> CBS 6054	3e-07
FE526995	-	No significant similarity	-	-
FE526996	-	No significant similarity	-	-
FE526997	ref XP_001268779.1	BAP31 domain protein, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	8e-14
FE526998	ref XP_001263178.1	ubiquitin conjugating enzyme (UbcB), putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	3e-08
FE526999	ref XP_001271404.1	heat shock protein Hsp30/Hsp42, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-08
FE527000	ref XP_001248200.1	predicted protein	<i>Coccidioides immitis</i> RS	1e-09
FE527001	-	No significant similarity	-	-
FE527002	-	No significant similarity	-	-
FE527003	-	No significant similarity	-	-
FE527004	-	No significant similarity	-	-
FE527005	-	No significant similarity	-	-
FE527006	ref XP_001275968.1	60S ribosomal protein L37a	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	7e-21
FE527007	-	No significant similarity	-	-
FE527008	-	No significant similarity	-	-
FE527009	-	No significant similarity	-	-
FE527010	ref XP_001273971.1	60S ribosomal protein L8, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	9e-26
FE527011	-	No significant similarity	-	-
FE527012	-	No significant similarity	-	-
FE527013	ref XP_001240365.1	hypothetical protein CIMG_07528	<i>Coccidioides immitis</i> RS	3e-05
FE527014	-	No significant similarity	-	-
FE527015	-	No significant similarity	-	-
FE527016	ref XP_001259429.1	dihydroxyacetone kinase (DakA), putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	5e-10
FE527017	-	No significant similarity	-	-
FE527018	-	No significant similarity	-	-
FE527019	-	No significant similarity	-	-
FE527020	-	No significant similarity	-	-

FE527021	ref XP_001266736.1	MFS monosaccharide transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	3e-20
FE527022	-	No significant similarity	-	-
FE527023	-	No significant similarity	-	-
FE527024	-	No significant similarity	-	-
FE527025	ref XP_001248308.1	60S ribosomal protein L28	<i>Coccidioides immitis</i> RS	2e-21
FE527026	ref XP_001242709.1	hypothetical protein CIMG_06605	<i>Coccidioides immitis</i> RS	4e-10
FE527027	ref XP_001245288.1	heat shock protein hsp1	<i>Coccidioides immitis</i> RS	2e-11
FE527028	ref XP_001248308.1	60S ribosomal protein L28	<i>Coccidioides immitis</i> RS	2e-21
FE527029	-	No significant similarity	-	-
FE527030	-	No significant similarity	-	-
FE527031	emb CAD70291.1	related to mitochondrial serine--tRNA ligase	<i>Neurospora crassa</i>	7e-05
FE527032	-	No significant similarity	-	-
FE527033	ref XP_001387342.1	60S large subunit ribosomal protein	<i>Pichia stipitis</i> CBS 6054	3e-04
FE527034	gb AAK52822.1 AF365926_1	calmodulin-binding coil-coil protein	<i>Emericella nidulans</i>	4e-07
FE527035	ref XP_001387342.1	60S large subunit ribosomal protein	<i>Pichia stipitis</i> CBS 6054	3e-04
FE527036	ref XP_001268653.1	TIM barrel metal-dependent hydrolase, putative	<i>Aspergillus clavatus</i> NRRL 1	3e-07
FE527037	-	No significant similarity	-	-
FE527038	ref XP_001265779.1	fatty acid desaturase, putative	<i>Neosartorya fischeri</i> NRRL 181	9e-29
FE527039	ref XP_753997.1	60s ribosomal protein y16	<i>Aspergillus fumigatus</i> Af293	1e-29
FE527040	emb CAA49847.1	GMP synthase	<i>Saccharomyces cerevisiae</i>	7e-12
FE527041	ref XP_001262785.1	MFS peptide transporter Ptr2, putative	<i>Neosartorya fischeri</i> NRRL 181	1e-15
FE527042	-	No significant similarity	-	-
FE527043	-	No significant similarity	-	-
FE527044	ref XP_001387342.1	60S large subunit ribosomal protein	<i>Pichia stipitis</i> CBS 6054	3e-04
FE527045	-	No significant similarity	-	-
FE527046	-	No significant similarity	-	-
FE527047	-	No significant similarity	-	-
FE527048	ref XP_754922.1	cytosolic small ribosomal subunit S4, putative	<i>Aspergillus fumigatus</i> Af293	2e-19
FE527049	-	No significant similarity	-	-
FE527050	-	No significant similarity	-	-
FE527051	-	No significant similarity	-	-

FE527052	ref XP_001273971.1	60S ribosomal protein L8, putative	<i>Aspergillus clavatus</i> NRRL 1	9e-26
FE527053	gb EDK03984.1	mitochondrial NADH-ubiquinone oxidoreductase 20 kD subunit	<i>Magnaporthe grisea</i> 70-15	8e-27
FE527054	-	No significant similarity	-	-
FE527055	ref XP_001218101.1	seryl-tRNA synthetase	<i>Aspergillus terreus</i> NIH2624	2e-08
FE527056	-	No significant similarity	-	-
FE527057	ref XP_001273971.1	60S ribosomal protein L8, putative	<i>Aspergillus clavatus</i> NRRL 1	3e-23
FE527058	ref XP_001265211.1	Ribosomal L18ae protein family	<i>Neosartorya fischeri</i> NRRL 181	2e-15
FE527059	ref XP_001261939.1	Hsp70 chaperone (HscA), putative	<i>Neosartorya fischeri</i> NRRL 181	1e-16
FE527060	ref XP_001241456.1	hypothetical protein CIMG_08619	<i>Coccidioides immitis</i> RS	2e-05
FE527061	-	No significant similarity	-	-
FE527062	-	No significant similarity	-	-
FE527063	-	No significant similarity	-	-
FE527064	-	No significant similarity	-	-
FE527065	-	No significant similarity	-	-
FE527066	-	No significant similarity	-	-
FE527067	ref XP_001240912.1	hypothetical protein CIMG_08075	<i>Coccidioides immitis</i> RS	2e-06
FE527068	ref XP_001268219.1	3-oxoacyl-(acyl-carrier-protein) reductase	<i>Aspergillus clavatus</i> NRRL 1	1e-21
FE527069	ref XP_001268984.1	PfkB family carbohydrate kinase (Mak32), putative	<i>Aspergillus clavatus</i> NRRL 1	1e-14
FE527070	ref XP_001215828.1	tRNA (uridine-2l-O-)-methyltransferase TRM7	<i>Aspergillus terreus</i> NIH2624	5e-13
FE527071	-	No significant similarity	-	-
FE527072	ref XP_001266337.1	phosphate transporter (Pho88), putative	<i>Neosartorya fischeri</i> NRRL 181	6e-07
FE527073	-	No significant similarity	-	-
FE527074	gb EAT88174.1	hypothetical protein SNOG_04414	<i>Phaeosphaeria nodorum</i> SN15	3e-04
FE527075	-	No significant similarity	-	-
FE527076	ref XP_001260419.1	60S ribosomal protein L10	<i>Neosartorya fischeri</i> NRRL 181	2e-28
FE527077	-	No significant similarity	-	-

FE527078	gb AAP22960.1	14-3-3-like protein	<i>Paracoccidioides brasiliensis</i>	9e-07
FE527079	sp Q8J1M3 DPP5_ARTBE	Dipeptidyl-peptidase 5 precursor (Dipeptidyl-peptidase V) (DPP V) (DppV) (Allergen Tri m 4) emb CAD23611.1 tri m 4 allergen	<i>Arthroderma benhamiae</i>	5e-21
FE527080	-	No significant similarity	-	-
FE527081	-	No significant similarity	-	-
FE527082	dbj BAB33421.1	putative senescence-associated protein	<i>Pisum sativum</i>	1e-26
FE527083	-	No significant similarity	-	-
FE527084	ref XP_001273971.1	60S ribosomal protein L8, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-16
FE527085	-	No significant similarity	-	-
FE527086	ref XP_001274406.1	DNA-directed RNA polymerases i, ii, and iii 145 kDa polypeptide	<i>Aspergillus clavatus</i> NRRL 1	2e-16
FE527087	ref XP_747704.1	prpd protein	<i>Aspergillus fumigatus</i> Af293	1e-24
FE527088	-	No significant similarity	-	-
FE527089	-	No significant similarity	-	-
FE527090	-	No significant similarity	-	-
FE527091	ref XP_001267123.1	60S ribosomal protein L25, putative	<i>Neosartorya fischeri</i> NRRL 181	4e-18
FE527092	-	No significant similarity	-	-
FE527093	-	No significant similarity	-	-
FE527094	sp A1CSR1 MRH4_ASPCL	ATP-dependent RNA helicase mrh4, mitochondrial precursor	<i>Aspergillus clavatus</i>	2e-14
FE527095	ref XP_001266736.1	MFS monosaccharide transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	7e-21
FE527096	ref XP_753520.1	D-amino acid oxidase	<i>Aspergillus fumigatus</i> Af293	4e-11
FE527097	gb ABE01845.1	beta-tubulin	<i>Microsporum canis</i>	7e-15
FE527098	-	No significant similarity	-	-
FE527099	-	No significant similarity	-	-
FE527100	gb EAT76933.1	hypothetical protein SNOG_15558	<i>Phaeosphaeria nodorum</i> SN15	1e-05
FE527101	-	No significant similarity	-	-
FE527102	ref XP_001268336.1	protein transport protein Sec24, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-20
FE527103	ref XP_001209399.1	succinate/fumarate mitochondrial transporter	<i>Aspergillus terreus</i> NIH2624	1e-32
FE527104	ref XP_001273981.1	vacuolar protein sorting protein DigA	<i>Aspergillus clavatus</i>	4e-24

FE527105	-	No significant similarity	NRRL 1	-
FE527106	ref XP_001270945.1	PDCD2_C domain protein, putative	<i>Aspergillus clavatus</i>	1e-08
FE527107	ref XP_001257579.1	2-methylcitrate dehydratase, putative	NRRL 1	
FE527108	ref XP_001264551.1	RING finger ubiquitin ligase (Tul1), putative	<i>Neosartorya fischeri</i>	3e-12
FE527109	dbj BAC01275.1	cytochrome P450nor	NRRL 181	
FE527110	ref XP_001275968.1	60S ribosomal protein L37a	<i>Neosartorya fischeri</i>	2e-09
FE527111	-	No significant similarity	NRRL 181	
FE527112	ref XP_001273487.1	short-chain dehydrogenase	<i>Aspergillus oryzae</i>	1e-19
FE527113	ref XP_001264756.1	adenosylhomocysteinase	<i>Aspergillus clavatus</i>	7e-21
FE527114	-	No significant similarity	NRRL 1	
FE527115	-	No significant similarity	<i>Neosartorya fischeri</i>	5e-23
FE527116	-	No significant similarity	NRRL 181	
FE527117	-	No significant similarity	-	-
FE527118	ref XP_001267248.1	60S ribosomal protein L11	<i>Aspergillus clavatus</i>	1e-05
FE527119	ref XP_001273971.1	60S ribosomal protein L8, putative	NRRL 1	
FE527120	-	No significant similarity	<i>Neosartorya fischeri</i>	6e-22
FE527121	-	No significant similarity	NRRL 181	
FE527122	-	No significant similarity	<i>Aspergillus clavatus</i>	3e-27
FE527123	-	No significant similarity	NRRL 1	
FE527124	ref XP_001387342.1	60S large subunit ribosomal protein	-	-
FE527125	-	No significant similarity	<i>Pichia stipitis</i> CBS 6054	3e-04
FE527126	ref XP_001271844.1	WD repeat protein	-	-
FE527127	ref XP_001268989.1	multidrug resistance protein 1, 2, 3 (p glycoprotein 1, 2, 3)	<i>Aspergillus clavatus</i>	1e-07
FE527128	ref XP_001240756.1	hypothetical protein CIMG_07919	NRRL 1	
FE527129	-	No significant similarity	<i>Aspergillus clavatus</i>	2e-09
FE527130	-	No significant similarity	NRRL 1	
FE527131	ref XP_752593.1	mitochondrial hypoxia responsive domain protein	<i>Coccidioides immitis</i> RS	2e-06
			-	-
			<i>Aspergillus fumigatus</i>	2e-33

FE527132	ref XP_001265914.1	sugar transporter, putative	<i>Af293</i> <i>Neosartorya fischeri</i>	1e-06
FE527133	ref XP_001269812.1	adenylate kinase 2	<i>NRRL 181</i> <i>Aspergillus clavatus</i>	9e-07
FE527134	-	No significant similarity	<i>NRRL 1</i>	-
FE527135	ref XP_001544887.1	conserved hypothetical protein	<i>Ajellomyces capsulatus</i>	4e-11
FE527136	ref XP_001541343.1	conserved hypothetical protein	<i>NAm1</i> <i>Ajellomyces capsulatus</i>	2e-05
FE527137	ref XP_001271103.1	Bromodomain protein	<i>NAm1</i> <i>Aspergillus clavatus</i>	3e-15
FE527138	ref XP_001248747.1	predicted protein	<i>NRRL 1</i> <i>Coccidioides immitis</i> RS	4e-08
FE527139	ref XP_001536346.1	40S ribosomal protein S18	<i>Ajellomyces capsulatus</i>	1e-31
FE527140	-	No significant similarity	<i>NAm1</i>	-
FE527141	ref XP_001217407.1	conserved hypothetical protein	<i>Aspergillus terreus</i>	1e-06
FE527142	-	No significant similarity	<i>NIH2624</i>	-
FE527143	-	No significant similarity	-	-
FE527144	ref XP_001543575.1	amino acid permease Dip5	<i>Ajellomyces capsulatus</i>	3e-26
FE527145	ref XP_001542093.1	sodium transport ATPase 5	<i>NAm1</i> <i>Ajellomyces capsulatus</i>	1e-12
FE527146	-	No significant similarity	<i>NAm1</i>	-
FE527147	-	No significant similarity	-	-
FE527148	-	No significant similarity	-	-
FE527149	-	No significant similarity	-	-
FE527150	-	No significant similarity	-	-
FE527151	ref XP_752484.2	eukaryotic translation elongation factor 1 subunit Eef1-beta, putative	<i>Aspergillus fumigatus</i>	5e-13
FE527152	ref XP_001257858.1	isocitrate dehydrogenase LysB	<i>Af293</i> <i>Neosartorya fischeri</i>	2e-47
FE527153	-	No significant similarity	<i>NRRL 181</i>	-
FE527154	-	No significant similarity	-	-
FE527155	-	No significant similarity	-	-
FE527156	ref XP_001268212.1	PF02656 domain protein	<i>Aspergillus clavatus</i>	2e-13
			<i>NRRL 1</i>	

FE527157	ref XP_001248747.1	predicted protein	<i>Coccidioides immitis</i> RS	1e-07
FE527158	ref XP_001260492.1	SET and MYND domain protein, putative	<i>Neosartorya fischeri</i> NRRL 181	6e-06
FE527159	ref XP_001273282.1	mitochondrial ribosomal protein DAP3, putative	<i>Aspergillus clavatus</i> NRRL 1	2e-12
FE527160	ref XP_001273062.1	non-classical export protein Nce2, putative	<i>Aspergillus clavatus</i> NRRL 1	5e-10
FE527161	ref XP_001273744.1	arginine permease	<i>Aspergillus clavatus</i> NRRL 1	1e-20
FE527162	ref XP_001536307.1	predicted protein	<i>Ajellomyces capsulatus</i> NA _m 1	1e-10
FE527163	ref XP_001259189.1	F-box domain protein	<i>Neosartorya fischeri</i> NRRL 181	7e-25
FE527164	ref XP_001537488.1	ER-derived vesicles protein ERV14	<i>Ajellomyces capsulatus</i> NA _m 1	2e-07
FE527165	-	No significant similarity	-	-
FE527166	-	No significant similarity	-	-
FE527167	ref XP_750288.1	mitochondrial ADP,ATP carrier protein (Ant), putative	<i>Aspergillus fumigatus</i> Af293	4e-22
FE527168	-	No significant similarity	-	-
FE527169	ref XP_001248747.1	predicted protein	<i>Coccidioides immitis</i> RS	3e-06
FE527170	ref XP_001265779.1	fatty acid desaturase, putative	<i>Neosartorya fischeri</i> NRRL 181	4e-29
FE527171	gb ABH10645.1	ATP synthase beta chain	<i>Coccidioides posadasii</i>	4e-21
FE527172	ref XP_001536346.1	40S ribosomal protein S18	<i>Ajellomyces capsulatus</i> NA _m 1	1e-26
FE527173	ref XP_001211936.1	26S protease regulatory subunit 8	<i>Aspergillus terreus</i> NIH2624	9e-12
FE527174	ref XP_001537882.1	thiamine-phosphate pyrophosphorylase	<i>Ajellomyces capsulatus</i> NA _m 1	2e-17
FE527175	ref XP_001260814.1	translation elongation factor EF-2 subunit, putative	<i>Neosartorya fischeri</i> NRRL 181	4e-11
FE527176	-	No significant similarity	-	-
FE527177	-	No significant similarity	-	-
FE527178	ref XP_001213274.1	conserved hypothetical protein	<i>Aspergillus terreus</i> NIH2624	1e-04
FE527179	ref XP_001242914.1	hypothetical protein CIMG_06810	<i>Coccidioides immitis</i> RS	3e-08
FE527180	-	No significant similarity	-	-

FE527181	-	No significant similarity	-	-
FE527182	-	No significant similarity	-	-
FE527183	-	No significant similarity	-	-
FE527184	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	2e-23
FE527185	ref XP_001266126.1	oxidosqualene:lanosterol cyclase	<i>Neosartorya fischeri</i> NRRL 181	2e-10
FE527186	ref YP_704246.1	possible dipeptidase	<i>Rhodococcus</i> sp. RHA1	2e-13
FE527187	ref XP_753972.1	glutathione synthetase, putative	<i>Aspergillus fumigatus</i> Af293	5e-12
FE527188	gb AAM54368.1	elongation factor 1-alpha	<i>Trichophyton rubrum</i>	2e-64
FE527189	ref XP_001240587.1	hypothetical protein CIMG_07750	<i>Coccidioides immitis</i> RS	1e-04
FE527190	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	3e-19
FE527191	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	3e-16
FE527192	ref XP_001544403.1	Cox19p	<i>Ajellomyces capsulatus</i> NAm1	1e-11
FE527193	-	No significant similarity	-	-
FE527194	ref XP_001540148.1	conserved hypothetical protein	<i>Ajellomyces capsulatus</i> NAm1	1e-18
FE527195	-	No significant similarity	-	-
FE527196	ref XP_001241535.1	hypothetical protein CIMG_08698	<i>Coccidioides immitis</i> RS	4e-06
FE527197	-	No significant similarity	-	-
FE527198	-	No significant similarity	-	-
FE527199	gb ABF82266.1	heat shock protein 30	<i>Penicillium marneffei</i>	2e-06
FE527200	-	No significant similarity	-	-
FE527201	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	4e-23
FE527202	-	No significant similarity	-	-
FE527203	ref XP_750370.1	N-acetylglucosamine-phosphate mutase	<i>Aspergillus fumigatus</i> Af293	1e-15
FE527204	-	No significant similarity	-	-
FE527205	ref XP_001257807.1	Rho GTPase Rho1	<i>Neosartorya fischeri</i> NRRL 181	3e-81
FE527206	ref XP_750999.1	DUF domain protein	<i>Aspergillus fumigatus</i> Af293	4e-05
FE527207	ref XP_001246171.1	hypothetical protein CIMG_05612	<i>Coccidioides immitis</i> RS	2e-08
FE527208	-	No significant similarity	-	-
FE527209	ref XP_001542733.1	ATP-binding cassette sub-family F member 2	<i>Ajellomyces capsulatus</i> NAm1	7e-43
FE527210	-	No significant similarity	-	-

FE527211	-	No significant similarity	-	-
FE527212	-	No significant similarity	-	-
FE527213	-	No significant similarity	-	-
FE527214	-	No significant similarity	-	-
FE527215	-	No significant similarity	-	-
FE527216	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	6e-24
FE527217	ref ZP_01504263.1	transcriptional regulator, AraC family	<i>Burkholderia phymatum</i> STM815	6e-08
FE527218	ref XP_001264609.1	CORD and CS domain protein	<i>Neosartorya fischeri</i> NRRL 181	5e-05
FE527219	ref XP_362903.2	c-14 sterol reductase	<i>Magnaporthe grisea</i> 70-15	9e-41
FE527220	ref XP_001276316.1	actin interacting protein 2	<i>Aspergillus clavatus</i> NRRL 1	2e-53
FE527221	-	No significant similarity	-	-
FE527222	-	No significant similarity	-	-
FE527223	dbj BAE65103.1	unnamed protein product	<i>Aspergillus oryzae</i>	1e-08
FE527224	-	No significant similarity	-	-
FE527225	ref XP_001258414.1	40S ribosomal protein S29, putative	<i>Neosartorya fischeri</i> NRRL 181	6e-10
FE527226	-	No significant similarity	-	-
FE527227	gb EEQ28484.1	iron sulfur cluster assembly protein 1, mitochondrial precursor	<i>Microsporum canis</i> CBS113480	1e-44
FE527228	ref XP_001238954.1	cytochrome c1,	<i>Coccidioides immitis</i> RS	1e-44
FE527229	-	No significant similarity	-	-
FE527230	ref XP_001543311.1	mitochondrial import inner membrane translocase subunit tim22	<i>Ajellomyces capsulatus</i> NAm1	9e-25
FE527231	ref XP_749294.1	non-classical export protein Nce102, putative	<i>Aspergillus fumigatus</i> Af293	3e-11
FE527232	-	No significant similarity	-	-
FE527233	gb ABG67901.1	putative phospholipase	<i>Trichophyton rubrum</i>	9e-12
FE527234	ref XP_001217348.1	lipoic acid synthetase, mitochondrial precursor	<i>Aspergillus terreus</i> NIH2624	1e-46
FE527235	-	No significant similarity	-	-
FE527236	ref XP_001255645.1	PREDICTED: similar to Keratin associated protein 10-4	<i>Bos taurus</i>	1e-07
FE527237	-	No significant similarity	-	-
FE527238	-	No significant similarity	-	-
FE527239	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	1e-41

FE527240	ref XP_001542349.1	nuclear protein SNF4	<i>Ajellomyces capsulatus</i> <i>NAm1</i>	1e-20
FE527241	sp Q0CA25 FYV10_ASPTN	Protein fyv10	<i>Aspergillus terreus</i> <i>NIH2624</i>	6e-31
FE527242	-	No significant similarity	-	-
FE527243	-	No significant similarity	-	-
FE527244	gb AAS45677.1	subtilisin-like protease SUB5	<i>Trichophyton</i> <i>verrucosum</i>	1e-43
FE527245	ref XP_001544767.1	coatomer beta subunit	<i>Ajellomyces capsulatus</i> <i>NAm1</i>	1e-16
FE527246	ref XP_001544767.1	coatomer beta subunit	<i>Ajellomyces capsulatus</i> <i>NAm1</i>	2e-31
FE527247	ref NP_487763.1	cation-transporting P-type ATPase	<i>Nostoc sp. PCC 7120</i>	7e-04
FE527248	ref XP_001268985.1	conserved hypothetical protein	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	9e-12
FE527249	gb AAO47089.1	mannitol-1-phosphate dehydrogenase	<i>Paracoccidioides</i> <i>brasiliensis</i>	1e-42
FE527250	-	No significant similarity	-	-
FE527251	dbj BAE65103.1	unnamed protein product	<i>Aspergillus oryzae</i>	5e-08
FE527252	ref XP_001273062.1	non-classical export protein Nce2, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	4e-10
FE527253	ref XP_001248046.1	casein kinase I homolog	<i>Coccidioides immitis RS</i>	6e-35
FE527254	-	No significant similarity	-	-
FE527255	-	No significant similarity	-	-
FE527256	ref XP_001242709.1	hypothetical protein CIMG_06605	<i>Coccidioides immitis RS</i>	5e-04
FE527257	ref XP_001542533.1	40S ribosomal protein S15	<i>Ajellomyces capsulatus</i> <i>NAm1</i>	4e-25
FE527258	ref XP_001538877.1	conserved hypothetical protein	<i>Ajellomyces capsulatus</i> <i>NAm1</i>	1e-17
FE527259	ref XP_001212080.1	enolase	<i>Aspergillus terreus</i> <i>NIH2624</i>	4e-28
FE527260	-	No significant similarity	-	-
FE527261	ref XP_001543567.1	lanosterol synthase	<i>Ajellomyces capsulatus</i> <i>NAm1</i>	4e-34
FE527262	-	No significant similarity	-	-
FE527263	ref XP_001266272.1	histone acetylase complex subunit Paf400, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-43
FE527264	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis RS</i>	3e-24

FE527265	-	No significant similarity	-	-
FE527266	ref XP_001261041.1	calcium dependent mitochondrial carrier protein, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-25
FE527267	ref XP_001274394.1	nucleoside diphosphatase (Ynd1), putative	<i>Aspergillus clavatus</i> NRRL 1	6e-10
FE527268	ref NP_830817.1	Oxalate decarboxylase	<i>Bacillus cereus</i> ATCC 14579	3e-20
FE527269	gb EAA29158.2	predicted protein	<i>Neurospora crassa</i> OR74A	1e-14
FE527270	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	1e-22
FE527271	gb ABB96277.1	hesp-767	<i>Melampsora lini</i>	2e-06
FE527272	ref XP_001263556.1	Coatomer subunit alpha, putative	<i>Neosartorya fischeri</i> NRRL 181	2e-07
FE527273	emb CAC28076.1	glucoamylase	<i>Talaromyces emersonii</i>	2e-29
FE527274	-	No significant similarity	-	-
FE527275	ref XP_001275305.1	actin cytoskeleton protein (VIP1), putative	<i>Aspergillus clavatus</i> NRRL 1	8e-21
FE527276	-	No significant similarity	-	-
FE527277	-	No significant similarity	-	-
FE527278	-	No significant similarity	-	-
FE527279	-	No significant similarity	-	-
FE527280	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	9e-22
FE527281	-	No significant similarity	-	-
FE527282	dbj BAF48663.1	leucine-rich repeat/extensin	<i>Nicotiana</i> <i>plumbaginifolia</i>	1e-04
FE527283	ref ZP_01617781.1	glutathione-dependent formaldehyde-activating, GFA	<i>marine gamma</i> <i>proteobacterium</i> HTCC2143	8e-05
FE527284	ref XP_001263728.1	phosphopantothenate-cysteine ligase, putative	<i>Neosartorya fischeri</i> NRRL 181	3e-19
FE527285	-	No significant similarity	-	-
FE527286	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	5e-24
FE527287	-	No significant similarity	-	-
FE527288	-	No significant similarity	-	-
FE527289	ref XP_001262431.1	phthalate transporter, putative	<i>Neosartorya fischeri</i> NRRL 181	3e-22
FE527290	gb ABY21304.1	thioredoxin TrxA	<i>Trichophyton</i> <i>mentagrophytes</i>	9e-41

FE527291	ref XP_752593.1	mitochondrial hypoxia responsive domain protein	<i>Aspergillus fumigatus</i> <i>Af293</i>	1e-24
FE527292	ref XP_001263032.1	HORMA domain protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	3e-18
FE527293	-	No significant similarity	-	-
FE527294	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	4e-23
FE527295	-	No significant similarity	-	-
FE527296	-	No significant similarity	-	-
FE527297	-	No significant similarity	-	-
FE527298	ref XP_754442.2	galactose-proton symport, putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	7e-04
FE527299	ref XP_001262893.1	DNA-directed RNA polymerases N/8 kDa subunit superfamily	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	4e-35
FE527300	-	No significant similarity	-	-
FE527301	ref XP_750553.1	endoplasmic reticulum DnaJ domain protein Erj5, putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	9e-21
FE527302	ref NP_730262.2	CG13731-PA	<i>Drosophila melanogaster</i>	5e-05
FE527303	-	No significant similarity	-	-
FE527304	ref XP_001227059.1	hypothetical protein CHGG_09132	<i>Chaetomium globosum</i> <i>CBS 148.51</i>	4e-34
FE527305	gb AAN87885.1	NADH-ubiquinone oxidoreductase	<i>Paracoccidioides brasiliensis</i>	3e-25
FE527306	ref XP_001264534.1	mitochondrial hypoxia responsive domain protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	1e-16
FE527307	-	No significant similarity	-	-
FE527308	-	No significant similarity	-	-
FE527309	ref XP_001214203.1	conserved hypothetical protein	<i>Aspergillus terreus</i> <i>NIH2624</i>	6e-25
FE527310	-	No significant similarity	-	-
FE527311	ref XP_001213878.1	homoserine dehydrogenase	<i>Aspergillus terreus</i> <i>NIH2624</i>	8e-52
FE527312	ref XP_748143.1	BYS1 domain protein, putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	2e-35
FE527313	dbj BAD90767.1	histone 3	<i>Conocephalum supradecompositum</i>	2e-22
FE527314	ref XP_001247458.1	78 kDa glucose-regulated protein homolog precursor	<i>Coccidioides immitis</i> RS	3e-73
FE527315	ref XP_001541705.1	hydroxymethylglutaryl-CoA lyase, mitochondrial precursor	<i>Ajellomyces capsulatus</i>	5e-50

FE527316	-	No significant similarity	<i>NAm1</i>	-
FE527317	-	No significant similarity	-	-
FE527318	ref XP_001261075.1	C-x8-C-x5-C-x3-H type zinc finger protein	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	7e-41
FE527319	ref XP_001268941.1	methyltransferase small domain protein	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	2e-23
FE527320	-	No significant similarity	-	-
FE527321	-	No significant similarity	-	-
FE527322	ref XP_001247341.1	hypothetical protein CIMG_01112	<i>Coccidioides immitis</i> RS	3e-24
FE527323	sp Q0CMM5 DBP4_ASPTN	ATP-dependent RNA helicase dbp4	<i>Aspergillus terreus</i> <i>NIH2624</i>	4e-31
FE527324	dbj BAD90801.1	histone 3	<i>Conocephalum conicum</i>	6e-38
FE527325	-	No significant similarity	-	-
FE527326	ref XP_001266180.1	thioredoxin reductase, putative	<i>Neosartorya fischeri</i> <i>NRRL 181</i>	4e-24
FE527327	-	No significant similarity	-	-
FE527328	ref XP_748143.1	BYS1 domain protein, putative	<i>Aspergillus fumigatus</i> <i>Af293</i>	8e-15
FE527329	ref XP_001672380.1	Hypothetical protein CBG01536	<i>Caenorhabditis briggsae</i>	5e-21
FE527330	-	No significant similarity	-	-
FE527331	-	No significant similarity	-	-
FE527332	dbj BAD90801.1	histone 3	<i>Conocephalum conicum</i>	2e-42
FE527333	ref XP_001244171.1	predicted protein	<i>Coccidioides immitis</i> RS	2e-19
FE527334	ref XP_001244171.1	predicted protein	<i>Coccidioides immitis</i> RS	4e-05
FE527335	ref XP_001270181.1	sodium P-type ATPase, putative	<i>Aspergillus clavatus</i> <i>NRRL 1</i>	9e-20
FE527336	ref XP_001543624.1	NADH-ubiquinone oxidoreductase 9.5 kDa subunit	<i>Ajellomyces capsulatus</i> <i>NAm1</i>	2e-13

Sequences with similarity in the nonredundant NCBI database (1e-3) using BLASTx.