

Table S2 Potential *N. crassa* IME-2 targets for phosphorylation, predicted by Scansite^{a, b}

NCU #	Annotation
NCU00101	PBN-1
NCU00258	40S ribosomal protein S7
NCU00269	Histone-lysine N-methyltransferase, H3 lysine-36 specific
NCU00276	DNA polymerase gamma, mitochondrial
NCU00311	Protein VTS1
NCU00552	Phytoene dehydrogenase; albino 1 protein
NCU00675	Protein EFR-3
NCU01157	Glutamate-cysteine ligase
NCU01206	Histone-lysine N-methyltransferase, H3 lysine-4 specific
NCU01312	Myb-like DNA-binding protein MYB-1
NCU01797	Serine/threonine-protein kinase NRC-2 (Non-repressible conidiation protein 2)
NCU01954	Pre-mRNA-splicing factor CWC-24
NCU02466	SVP1-like protein 2
NCU03678	Protein SSH-4
NCU03725	Transcription factor VIB-1
NCU03819	COPII coat assembly protein SEC-16
NCU03894	Serine/threonine-protein kinase STE-20
NCU04190	Probable exocyst complex component SEC-8
NCU04191	Protein RAI-1
NCU05589	Protein SIP-5
NCU05876	pH-response regulator protein PAL-A/RIM-20
NCU06276	Increased rDNA silencing protein 4
NCU06498	Pre-mRNA-splicing factor CWC-21
NCU07693	Protein STU-1
NCU07863	Chromatin modification-related protein VID-21
NCU08566	Nuclear distribution protein nudE homolog 1 (RO-11)
NCU08699	Putative oxidoreductase BLI-4, mitochondrial
NCU09068	Nitrogen catabolic enzyme regulatory protein
NCU09730	Kinesin heavy chain
NCU09993	Helicase SWR-1

^a <http://scansite.mit.edu/>; ^b one mismatch was allowed at the final consensus position