

**Table S1.** Identification of *Botrytis cinerea* proteins in coculture with Mp-30 cells in wounded apples. Protein names and accession numbers correspond to the UniProt database.

Entry	Gene names	Protein names	Mol. weight [kDa]
G2XN03	BofuT4_P013870.1	Proteasome subunit alpha type	31,99
G2XN09	BofuT4P9000002001	Proteasome subunit beta	24,77
G2XN77	BofuT4_P014610.1	5-aminoimidazole-4-carboxamide ribonucleotide formyltransferase (EC 2.1.2.3) (EC 3.5.4.10) (AICAR transformylase) (IMP synthase) (Inosinicase) (Phosphoribosylaminoimidazolecarboxamide formyltransferase)	65,056
G2XN83	BofuT4P9000017001	Similar to 60s ribosomal protein l11	19,963
G2XND8	BofuT4_P015220.1	Clathrin heavy chain	189,4
G2XNN8	BofuT4_P075770.1	Triosephosphate isomerase (EC 5.3.1.1)	27,012
G2XPB8	BofuT4_P073360.1	60S ribosomal protein L27	15,611
G2XPL9	BofuT4_P072420.1	TPR_MLP1_2 domain-containing protein	230,23
G2XPM1	BofuT4_P072440.1	NADH dehydrogenase [ubiquinone] flavoprotein 1, mitochondrial (EC 7.1.1.2)	55,722
G2XPW6	BofuT4_P071500.1	Similar to heat shock protein 90	79,536
G2XPY2	BofuT4_P071660.1	Similar to vacuolar sorting-associated protein	25,297
G2XQE0	BofuT4_P068850.1	Pyruvate kinase (EC 2.7.1.40)	57,905
G2XQE4	BofuT4_P068890.1	Similar to flavin-nucleotide-binding protein	29,434
G2XQE6	BofuT4_P068910.1	Similar to phosphatidylserine decarboxylase	46,447
G2XZU5	BofuT4_P050000.1	Phospho-2-dehydro-3-deoxyheptonate aldolase (EC 2.5.1.54)	39,371
G2XQL5	BofuT4_P069600.1	Uncharacterized protein	61,536
G2XQQ6	BofuT4_P068640.1	Pyruvate carboxylase (EC 6.4.1.1)	130,52
G2XQV1	BofuT4_P068540.1	Similar to stress response protein Rds1	55,355
G2XRG1	BofuT4_P067370.1	AP complex subunit beta	80,832
G2XS58	BofuT4P6000002001	Uncharacterized protein	55,382
G2XS65	BofuT4P6000004001	Similar to 40s ribosomal protein s5	23,746
G2XSE9	BofuT4_P012620.1	Uncharacterized protein	52,119
G2XSU2	BofuT4_P064120.1	Ribosomal_L18A domain-containing protein	27,397
G2XT31	BofuT4_P009930.1	Similar to transcription factor Zn, C2H2	67,994
G2XT46	BofuT4_P010080.1	Uncharacterized protein	46,866
G2XT89	BofuT4_P010510.1	Similar to U1biquitin-specific peptidase	80,062
G2XTL2	BofuT4_P062780.1	ADH_zinc_N domain-containing protein	17,121
G2XTR1	BofuT4_P060910.1	ADP/ATP translocase (ADP,ATP carrier protein)	33,957
G2XTR5	BofuT4_P060950.1	Similar to mitochondrial phosphate carrier protein	33,612
G2XU47	BofuT4_P062270.1	Catalase (EC 1.11.1.6)	57,482
G2XUD1	BofuT4_P060200.1	RRM domain-containing protein	10,351
G2XUE9	BofuT4P53000001001	Similar to alcohol dehydrogenase	36,015
G2XUF2	BofuT4_P057020.1	Uncharacterized protein	162,01
G2XUJ6	BofuT4_P057460.1	BcGBL1, heterotrimeric Gbeta-like protein	34,931
G2XUP1	BofuT4_P057910.1	40S ribosomal protein S24	15,439
G2XV82	BofuT4_P059820.1	BcNRPS9, nonribosomal peptide synthetase	485,2
G2XVE8	BofuT4_P053690.1	Similar to 60S ribosomal protein L44	12,045
G2XVH9	BofuT4_P054000.1	Similar to ATK1 (ARABIDOPSIS THALIANA KINESIN 1) microtubule motor/ minus-end-directed microtubule motor	78,262
G2XVU6	BofuT4_P055170.1	Similar to ATP-dependent RNA helicase sub2	49,626
G2XVW1	BofuT4_P055320.1	Similar to ribosomal protein L14	15,936
G2XW60	BofuT4_P056310.1	Phosphotransferase (EC 2.7.1.-)	60,598
G2XWG2	BofuT4_P051810.1	Diadenosine tetraphosphate synthetase (EC 6.1.1.14)	74,643
G2XW19	BofuT4P48000005001	Alpha-galactosidase (EC 3.2.1.22) (Melibiase)	54,183
G2XWL5	BofuT4_P052340.1	Similar to 60S ribosomal protein L8	27,405
G2XXA6	BofuT4_P009010.1	5-methyltetrahydropteroyltriglutamate--homocysteine S-methyltransferase (EC 2.1.1.14)	86,237
G2XXC6	BofuT4_P009210.1	BRO1 domain-containing protein	97,212
G2XXJ4	BofuT4_P007500.1	Similar to chromatin remodelling complex ATPase chain ISW1	128,94
G2XYH0	BofuT4_P045250.1	Similar to carbamoyl-phosphate synthase arginine-specific large chain	127,94
G2XYH5	BofuT4_P045300.1	Similar to eukaryotic initiation factor 4a-i	44,874
G2XYV3	BofuT4_P046580.1	2-phosphoglycerate dehydratase (EC 4.2.1.11) (Enolase)	47,154
G2XZH4	BofuT4_P048790.1	Lcc1, laccase	60,788
G2XZL6	BofuT4_P049210.1	Similar to oxidoreductase	37,524
G2XZP3	BofuT4_P049480.1	Similar to nuclear movement protein nudC	21,212
G2XZT8	BofuT4_P049930.1	Similar to ubiquitin-conjugating enzyme	15,5
G2Y0C8	BofuT4_P116430.1	Eukaryotic translation initiation factor 5A (eIF-5A)	17,824
G2Y0I2	BofuT4_P116970.1	Elongation factor 2	93,459
G2Y0V2	BofuT4_P118170.1	Similar to ThiJ/Pfpl family protein	31,615
G2Y0Z5	BofuT4_P118600.1	6-phosphogluconate dehydrogenase, decarboxylating (EC 1.1.1.44)	54,71
G2Y167	BofuT4P32000003001	Similar to mitochondrial-processing peptidase subunit beta	52,811

G2Y187	BofuT4_P040380.1	Saccharopine dehydrogenase [NAD(+), L-lysine-forming] (SDH) (EC 1.5.1.7) (Lysine--2-oxoglutarate reductase)	40,86
G2Y1A2	BofuT4P32000012001	Similar to ribosome assembly protein Noc2	88,03
G2Y1C6	BofuT4P32000015001	Snoal-like domain-containing protein	20,925
G2Y1Z2	BofuT4_P042930.1	Similar to 60S ribosomal protein l23	14,65
G2Y2E1	BofuT4_P115370.1	Similar to 60S ribosomal protein L18	20,735
G2Y2G1	BofuT4_P115570.1	Fatty acid hydroxylase domain-containing protein	40,969
G2Y3C1	BofuT4_P003200.1	Mitogen-activated protein kinase (EC 2.7.11.24)	40,499
G2Y3F9	BofuT4P2000014001	Similar to monothiol glutaredoxin-3	28,432
G2Y3T0	BofuT4P2000030001	Similar to amidophosphoribosyltransferase	236,28
G2Y3Y5	BofuT4_P005340.1	Glyceraldehyde-3-phosphate dehydrogenase (EC 1.2.1.12)	44,817
G2Y3Z0	BofuT4P2000050001	ATP synthase subunit beta (EC 7.1.2.2)	54,546
G2Y465	BofuT4_P006140.1	SET domain-containing protein	108,65
G2Y4A4	BofuT4_P006530.1	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial (EC 1.3.5.1)	54,065
G2Y4C1	BofuT4P2000074001	ATP synthase subunit alpha	59,85
G2Y4V3	BofuT4_P036730.1	Uncharacterized protein	257,86
G2Y528	BofuT4_P037480.1	Similar to phosphoglucomutase	60,028
G2Y5A0	BofuT4_P113440.1	Elongation factor 1-alpha	49,983
G2Y5Q7	BofuT4_P113070.1	Uncharacterized protein	152,49
G2Y5X8	BofuT4_P110430.1	40S ribosomal protein S27	8,8514
G2Y6F3	BofuT4_P034980.1	Uncharacterized protein	26,358
G2Y6Z6	BofuT4_P107820.1	Similar to replication factor C subunit 3	42,801
G2Y729	BofuT4_P108150.1	Uncharacterized protein	40,327
G2Y7A0	BofuT4_P108860.1	BcPKS2, polyketide synthase	349,78
G2Y7A5	BofuT4_P108910.1	PKS_ER domain-containing protein	34,991
G2Y7D4	BofuT4_P109200.1	Similar to 26S protease regulatory subunit 7	47,492
G2Y7M1	BofuT4_P110070.1	Similar to mitochondrial carrier protein	32,945
G2Y7P9	BofuT4_P110350.1	Citrate synthase	51,928
G2Y7U9	BofuT4_P033190.1	Uncharacterized protein	163,63
G2Y819	BofuT4P25000008001	ActA, actin	47,927
G2Y8V5	BofuT4P22000016001	Similar to isopropylmalate dehydrogenase	38,214
G2Y937	BofuT4_P029890.1	Uncharacterized protein	20,214
G2Y9J7	BofuT4_P031490.1	Uncharacterized protein	40,277
G2Y9N5	BofuT4_P031870.1	Similar to 40S ribosomal protein S11	18,395
G2Y9Q7	BofuT4_P032090.1	Uncharacterized protein	21,247
G2Y9Z0	BofuT4_P027370.1	Similar to ubiquitin-conjugating enzyme	23,468
G2YAL4	BofuT4_P103830.1	Uncharacterized protein	16,715
G2YB79	BofuT4_P102650.1	Similar to 60S ribosomal protein L27a	16,867
G2YBG2	BofuT4_P102540.1	Ribosomal_S10 domain-containing protein	11,238
G2YBK9	BofuT4_P101990.1	Adenosylhomocysteinase (EC 3.3.1.1)	48,849
G2YBN8	BofuT4P173000001001	Glycoside hydrolase family 17 protein	44,123
G2YBZ3	BofuT4_P101380.1	Similar to mRNA export factor elf1	122,83
G2YC48	BofuT4_P099990.1	Uncharacterized protein	48,063
G2YCC8	BofuT4_P099250.1	Similar to AhpC/TSA family protein	16,306
G2YCE4	BofuT4_P099410.1	Similar to type 11 methyltransferase	26,05
G2YCG2	BofuT4_P098500.1	Ribosomal protein L15	33,944
G2YCI1	BofuT4_P098790.1	Similar to zinc-containing alcohol dehydrogenase	37,477
G2YCV9	BofuT4_P097150.1	Similar to Mis12 domain-containing protein	38,225
G2YD26	BofuT4_P093820.1	40S ribosomal protein S8	28,811
G2YD44	BofuT4_P094000.1	40S ribosomal protein S6	27,201
G2YD46	BofuT4_P094020.1	Ribose-phosphate diphosphokinase (EC 2.7.6.1)	34,771
G2YD81	BofuT4P153000002001	Vacuolar proton pump subunit B (V-ATPase subunit B) (Vacuolar proton pump subunit B)	57,599
G2YDD2	BofuT4_P094880.1	Similar to Mitochondrial porin (Voltage-dependent anion channel)	34,416
G2YDQ1	BofuT4_P096070.1	Uncharacterized protein	48,529
G2YE11	BofuT4_P092810.1	Similar to 60S ribosomal protein L5	34,022
G2YEA8	BofuT4_P025050.1	Similar to 60S ribosomal protein L24	17,478
G2YEC3	BofuT4_P025200.1	Similar to NACHT domain protein	112,17
G2YEW4	BofuT4P143000024001	Similar to 26S protease regulatory subunit 6A	51,348
G2YEY4	BofuT4P143000032001	60S ribosomal protein L6	22,137
G2YF95	BofuT4_P089890.1	Signal peptidase complex catalytic subunit SEC11 (EC 3.4.21.89)	19,047
G2YFT6	BofuT4_P024430.1	RNA polymerase II subunit A C-terminal domain phosphatase (EC 3.1.3.16)	93,336
G2YFU1	BofuT4_P024480.1	EkdA	13,141
G2YG89	BofuT4_P088500.1	Similar to 26S protease regulatory subunit 6b	47,174
G2YGC2	BofuT4_P087240.1	Similar to 60S ribosomal protein L10	25,372
G2YGC3	BofuT4_P087250.1	Uncharacterized protein	34,243
G2YGL9	BofuT4_P086960.1	Similar to 40S ribosomal protein S3	28,512
G2YGR0	BofuT4_P022120.1	Translationally-controlled tumor protein homolog	16,771
G2YGR9	BofuT4_P022210.1	Similar to 60S ribosomal protein L17	19,293
G2YGS0	BofuT4_P022220.1	Similar to 40S ribosomal protein S18	16,858

G2YGW4	BofuT4_P022660.1	Similar to 60S ribosomal protein l12	20,653
G2YGW7	BofuT4_P022690.1	40S ribosomal protein S7	22,526
G2YHA6	BofuT4_P021590.1	Uncharacterized protein	332,83
G2YHB1	BofuT4_P021640.1	Transaldolase (EC 2.2.1.2)	35,402
G2YHI5	BofuT4_P085340.1	BcPKS20, polyketide synthase	294,72
G2YHK1	BofuT4_P085500.1	Plasma membrane ATPase (EC 7.1.2.1)	96,33
G2YHK3	BofuT4_P085520.1	Similar to O-acetylhomoserine/O-acetylserine sulfhydrylase	48,927
G2YHN9	BofuT4_P015320.1	Similar to 60s ribosomal protein l7	28,62
G2YHW8	BofuT4_P016110.1	Similar to 40S ribosomal protein S22	14,644
G2YHZ3	BofuT4_P016360.1	Similar to 40S ribosomal protein S16	15,9
G2YHZ5	BofuT4P10000009001	Similar to 40S ribosomal protein S14	16,013
G2YI43	BofuT4_P016860.1	Similar to heat shock protein 70	68,669
G2YI83	BofuT4P10000021001	GDP-mannose pyrophosphorylase (EC 2.7.7.13) (GTP-mannose-1-phosphate guanylyltransferase) (Mannose-1-phosphate guanylyltransferase)	39,906
G2YIA7	BofuT4P10000025001	Seryl-tRNA synthetase (EC 6.1.1.11) (Seryl-tRNA(Ser/Sec) synthetase)	54,535
	RPS1		
G2YIB7	BofuT4_P017600.1	40S ribosomal protein S1	29,246
G2YIE3	BofuT4_P017860.1	Similar to elongation factor 1-gamma	46,935
G2YJN1	BofuT4_P084720.1	Similar to ras-related protein rab-7a	22,998
G2YJR1	BofuT4_P083810.1	Uncharacterized protein	70,692
G2YKK1	BofuT4P103000006001	Similar to ADP-ribosylation factor	20,994
G2YKM7	BofuT4_P079700.1	Similar to ATP-dependent Clp protease	88,366
G2YL11	BofuT4_P081040.1	S-adenosylmethionine synthase (EC 2.5.1.6)	43,121
G2YL85	BofuT4_P078980.1	Isoleucyl-tRNA synthetase (EC 6.1.1.5)	123,13
G2YLR6	BofuT4_P000110.1	Similar to transcription factor Zn, C2H2	30,771
G2YLZ3	BofuT4P0000017001	Serine/threonine-protein phosphatase (EC 3.1.3.16)	37,227
G2YM46	BofuT4P0000042001	Similar to elongation factor 3	118,52
G2YMA9	BofuT4_P002040.1	Carbamoyl-phosphate synthase (glutamine-hydrolyzing) (EC 6.3.5.5)	184,19
G2YMB4	BofuT4P0000076001	Similar to transcription factor Zn, C2H2	78,143
G2YMH6	BofuT4_P137480.1	Uncharacterized protein	52,804
G2YND3	BofuT4_P121500.1	Similar to 40s ribosomal protein S15	17,571
G2YNI1	BofuT4_P121980.1	Similar to eukaryotic translation initiation factor eIF-4A subunit	45,484
G2YNN0	BofuT4_P122470.1	Similar to 60S ribosomal protein L4	40,04
G2YNN4	BofuT4_P122510.1	Similar to cytochrome c oxidase subunit 5A	18,6
G2YNV9	BofuT4_P123260.1	Ricin B-type lectin domain-containing protein	16,626
G2YNY7	BofuT4_P123540.1	Similar to ubiquitin-conjugating enzyme E2	16,527
G2YP99	BofuT4P79000021001	Peptidyl-prolyl cis-trans isomerase (PPIase) (EC 5.2.1.8)	24,225
	RPS0		
G2YPH0	BofuT4P79000014001	40S ribosomal protein S0	31,721
G2YPH3	BofuT4P79000016001	Nucleolar protein 58	61,264
G2YQ72	BofuT4_P133240.1	Similar to 40s ribosomal protein S17	16,881
G2YQ78	BofuT4_P133300.1	Similar to glutathione S-transferase	42,801
G2YQS9	BofuT4_P131570.1	Similar to subtilisin-like serine protease (Secreted protein)	55,253
G2YR90	BofuT4_P128380.1	E3 ubiquitin protein ligase (EC 2.3.2.27)	80,449
G2YRD8	BofuT4_P128860.1	Similar to aspartyl aminopeptidase	55,924
G2YRE7	BofuT4_P128950.1	Small COPII coat GTPase SAR1 (Small COPII coat GTPase sar1)	21,519
G2YRP9	BofuT4_P129970.1	Succinate--CoA ligase [ADP-forming] subunit alpha, mitochondrial (EC 6.2.1.5) (Succinyl-CoA synthetase subunit alpha) (SCS-alpha)	34,398
G2YRU0	BofuT4_P130380.1	Similar to 60S ribosomal protein L16	23,031
G2YRV9	BofuT4_P124050.1	Similar to pre-mRNA splicing factor ATP-dependent RNA helicase PRP43	85,722
G2YSM3	BofuT4_P126690.1	Endoplasmic reticulum chaperone BiP (EC 3.6.4.10) (Immunoglobulin heavy chain-binding protein homolog)	73,12
G2YSN7	BofuT4_P126830.1	Similar to P-type ATPase	114,3
G2YT44	BofuT4_P162370.1	UTP--glucose-1-phosphate uridylyltransferase (EC 2.7.7.9)	58,609
	MET3		
G2YTD1	BofuT4_P163240.1	Sulfate adenyltransferase (EC 2.7.7.4) (ATP-sulfurylase) (Sulfate adenylate transferase) (SAT)	54,9
G2YTJ0	BofuT4_P160390.1	Similar to 60s ribosomal protein l21	18,296
G2YTP4	BofuT4_P160930.1	Similar to 14-3-3 protein	29,623
G2YTV0	BofuT4_P161490.1	Inorganic diphosphatase (EC 3.6.1.1)	32,172
G2YU25	BofuT4_P159440.1	Actin-related protein 2/3 complex subunit 4	19,574
G2YUG2	BofuT4_P156560.1	Similar to HET domain protein	75,58
G2YUP2	BofuT4_P157360.1	Similar to mitochondrial heat shock protein	73,462
G2YV39	BofuT4_P158830.1	Similar to heat shock protein 60	61,162
G2YV41	BofuT4_P158850.1	Actin-depolymerizing factor 1 (Cofilin)	15,509
		3-oxoacyl-[acyl-carrier-protein] reductase (EC 1.1.1.100) (EC 2.3.1.41) (EC 2.3.1.86) (3-oxoacyl-[acyl-carrier-protein] synthase) (Acyl carrier) (Beta-ketoacyl reductase) (Beta-ketoacyl synthase) (Fatty acid synthase subunit alpha)	
G2YVH4	BofuT4_P154340.1		204,22
G2YVK4	BofuT4_P154640.1	Similar to eukaryotic translation initiation factor 2 subunit alpha	35,217
G2YVU9	BofuT4_P153090.1	40S ribosomal protein S26	13,472
G2YVZ6	BofuT4_P153560.1	Dihydrolipoyl dehydrogenase (EC 1.8.1.4)	54,55
G2YW49	BofuT4_P154090.1	Histone H2B	14,941

G2YWD7	BofuT4P145000004001	Similar to aldo/keto reductase	31,594
G2YWP7	BofuT4_P151770.1	Malate dehydrogenase (EC 1.1.1.37)	34,473
G2YWP8	BofuT4_P151780.1	Similar to 60S ribosomal protein L28	16,008
G2YWQ9	BofuT4_P151890.1	Uncharacterized protein	22,194
G2YWX7	BofuT4_P148250.1	Catalase (EC 1.11.1.6)	63,852
G2YX82	BofuT4_P149300.1	Fructose-bisphosphate aldolase (FBP aldolase) (EC 4.1.2.13)	39,241
G2YXB1	BofuT4_P149590.1	Uncharacterized protein	22,075
G2YXS7	BofuT4_P145940.1	Uncharacterized protein	59,922
G2YXT2	BofuT4_P145990.1	t-SNARE coiled-coil homology domain-containing protein	37,879
G2YY24	BofuT4_P146910.1	40S ribosomal protein S4	29,332
G2YY59	BofuT4P106000003001	Ketol-acid reductoisomerase, mitochondrial (EC 1.1.1.86) (Acetohydroxy-acid reductoisomerase) (Alpha-keto-beta-hydroxylacyl reductoisomerase)	44,169
G2YY64	BofuT4P106000005001	Tryptophan synthase (EC 4.2.1.20)	78,038
G2YYA1	BofuT4P106000015001	Similar to actin-like protein (Contractin)	42,223
G2YYT2	BofuT4_P140420.1	Similar to GTP-binding protein rho3	21,646
R7R399	BofuT4P114000012001	Uncharacterized protein	66,856