

182481_at	F35D2.1 /// status:Predicted	1.34
182489_at	F53B3.2 /// status:Partially_confirmed	1.17
182490_at	status:Partially_confirmed /// ZK470.6	2.52
182510_at	T07G12.3 /// status:Partially_confirmed	1.21
182522_at	status:Predicted /// Y17D7B.5	1.90
182561_at	status:Partially_confirmed /// Y11D7A.9	1.83
182600_at	F02D10.6 /// status:Partially_confirmed	1.16
182629_at	C50D2.1 /// status:Partially_confirmed	1.47
182639_at	C25F6.6 /// status:Predicted	2.15
182683_at	status:Partially_confirmed /// ZC250.3	1.80
182696_s_at	Chondroitin ProteoGlycan	1.23
182733_at	C31A11.5 /// status:Partially_confirmed	1.98
182743_at	status:Confirmed /// Y47D3B.6	1.34
182761_at	R04A9.1 /// status:Predicted	1.20
182776_at	status:Confirmed /// Y45F3A.8	2.59
182812_at	F22F4.1 /// status:Confirmed	1.26
182820_at	C54G10.1 /// status:Partially_confirmed	1.38
182923_at	F56D2.3 /// status:Partially_confirmed	1.25
182987_at	W03B1.2 /// status:Partially_confirmed	1.29
183049_at	B0432.7 /// status:Partially_confirmed	1.11
183092_at	F15H10.7 /// status:Partially_confirmed	1.77
183120_at	C06G1.1b	1.04
183136_at	R05G6.9 /// status:Confirmed	1.07
183144_at	B0348.5 /// status:Partially_confirmed	1.58
183147_at	C06G1.2 /// status:Confirmed	2.43
183159_at	F09F9.2 /// status:Confirmed	1.51
183176_at	T07C12.8 /// status:Predicted	1.61
183185_at	F15H10.8 /// status:Partially_confirmed	1.61
183191_at	F38B6.7 /// status:Partially_confirmed	1.55
183199_at	F59A1.12 /// status:Predicted	1.09
183222_at	status:Partially_confirmed /// ZC449.1	1.01
183237_s_at	ZK180.5c	2.46
183263_s_at	F59B1.8 /// status:Partially_confirmed	2.16
183271_at	---	1.14
183289_at	---	2.33
183300_at	F41A4.1 /// locus:cutl-28	1.38
183333_at	Kinesin-Like Protein	1.09
183363_at	status:Partially_confirmed /// ZC513.1	1.48
183379_at	K01D12.9 /// status:Confirmed	1.38
183407_at	F40H3.3 /// status:Partially_confirmed	1.61
183415_at	K01D12.3 /// status:Partially_confirmed	4.40
183454_at	F38B6.1 /// status:Partially_confirmed	1.07
183480_at	C32H11.6 /// status:Predicted	1.83
183537_at	C03G6.6 /// status:Partially_confirmed	1.28
183539_s_at	C06E7.4 /// status:Confirmed	1.77

183577_at	Prion-like-(Q/N-rich)-domain-bearing protein /// locus:pqn-95	2.53
183582_at	C47E12.6a	2.58
183601_at	C06E7.4 /// status:Confirmed	2.13
183673_at	F36H5.4 /// status:Confirmed	1.33
183702_at	K05B2.4 /// status:Partially_confirmed	1.51
183717_at	F20B10.3 /// status:Partially_confirmed	1.01
183724_at	status:Confirmed /// Y11D7A.5	1.44
183726_at	Serpentine Receptor, class Z /// locus:srz-4	1.97
183731_at	K01D12.1 /// status:Partially_confirmed	3.77
183733_at	F44C8.7 /// status:Confirmed	2.33
183755_s_at	F46C8.8 /// status:Confirmed	2.14
183765_s_at	F34D10.4 /// status:Partially_confirmed	1.23
183818_s_at	C05E7.1 /// status:Partially_confirmed	1.69
183826_at	F01G10.9 /// status:Partially_confirmed	1.48
183859_at	F13H10.1 /// status:Partially_confirmed	1.01
183911_at	T19A5.3b	1.41
183941_s_at	F56D3.1 /// status:Confirmed	1.39
183961_at	K05F1.5 /// status:Partially_confirmed	2.35
183967_at	B0432.8 /// status:Partially_confirmed	1.32
183975_s_at	Prion-like-(Q/N-rich)-domain-bearing protein /// locus:pqn-46	1.65
183995_s_at	MATH (meprin-associated Traf homology) domain containing /// MATH (meprin-associated Traf homology) domain containing /// locus:math-48	2.14
184008_at	D1022.5 /// status:Partially_confirmed	2.04
184023_at	D1022.3 /// status:Partially_confirmed	1.26
184047_at	UDP-GlucuronosylTransferase /// locus:ugt-36	2.62
184068_at	R05D8.7 /// status:Partially_confirmed	7.13
184081_at	Serpentine Receptor, class X /// locus:srx-16	1.05
184091_at	GRound-Like (grd related) /// locus:grl-10	1.65
184106_at	M03F8.1 /// status:Partially_confirmed	4.04
184125_at	C05D12.1 /// status:Partially_confirmed	1.16
184138_at	K02E7.1a	2.18
184140_at	F49E8.2 /// status:Partially_confirmed	1.88
184151_at	F26A1.9 /// status:Partially_confirmed	1.70
184199_s_at	C04F1.1 /// status:Partially_confirmed	2.04
184205_at	status:Partially_confirmed /// Y52B11B.1	2.04
184218_at	W06B11.1 /// status:Confirmed	1.11
184220_at	C56C10.4 /// status:Confirmed	1.65
184236_at	T24C12.4 /// status:Partially_confirmed	2.76
184247_at	F49D11.4 /// status:Predicted	1.31
184248_at	C.Elegans Y-box /// locus:cey-2	1.08
184254_at	C46H11.7 /// status:Partially_confirmed	2.13
184255_at	PHaryngeal gland Toxin-related /// locus:phat-1	1.17
184268_s_at	PHaryngeal gland Toxin-related /// locus:phat-2	1.71
184335_at	Dumpy : shorter than wild-type /// locus:dpy-1	1.03

184342_at	F56F11.5 /// status:Partially_confirmed	1.28
184352_at	C17H12.6 /// status:Partially_confirmed	2.03
184447_at	status:Partially_confirmed /// Y59A8B.22	1.27
184517_at	status:Predicted /// Y51H4A.21	1.31
184560_s_at	F16G10.9 /// status:Partially_confirmed	1.02
184602_at	UDP-GlucuronosylTransferase /// locus:ugt-25	2.10
184629_at	Ligand-Gated ion Channel	1.36
184662_at	status:Partially_confirmed /// Y46G5A.29	1.33
184667_at	H27M09.3 /// status:Partially_confirmed	1.85
184682_s_at	K09B11.10 /// status:Partially_confirmed	1.97
184684_at	Y53F4B.27a	1.51
184693_s_at	C32H11.11 /// C32H11.8 /// status:Predicted /// status:Predicted	1.78
184694_at	LYSozyme /// locus:lys-10	1.68
184746_at	F52F10.2 /// status:Partially_confirmed	1.25
184772_at	C01G10.14 /// C25D7.1	1.16
184814_at	Serpentine Receptor, class Z /// locus:srz-28	1.90
184828_s_at	MoLTing defective	1.44
184834_s_at	F26F12.4 /// status:Confirmed	1.39
184948_at	status:Predicted /// Y57G11B.2	1.01
184955_at	F35F10.7 /// status:Partially_confirmed	1.81
184968_at	F01G12.1 /// status:Predicted	1.37
185008_at	Serpentine Receptor, class BC (class B-like) /// locus:srbc-1	1.20
185020_at	C33E10.5 /// status:Predicted	1.19
185074_at	GRound-Like (grd related) /// locus:grl-12	1.18
185081_at	C-type LECTin /// locus:clec-210	3.62
185108_at	F36F12.1 /// status:Partially_confirmed	2.83
185172_s_at	CHitin Synthase /// locus:chs-2	1.22
185186_at	Serpentine Receptor, class V /// locus:srv-19	2.14
185231_at	F41E6.14 /// status:Predicted	1.28
185238_at	status:Partially_confirmed /// Y67A10A.8	1.12
185287_at	status:Partially_confirmed /// Y53F4B.25	2.11
185328_at	UDP-GlucuronosylTransferase /// locus:ugt-28	1.33
185334_s_at	R02F11.1 /// status:Confirmed	1.52
185343_at	T28C12.6 /// status:Predicted	1.00
185371_at	F38H12.5 /// status:Predicted	3.25
185390_at	F49D11.6 /// status:Predicted	2.45
185412_at	T22B11.2 /// status:Partially_confirmed	1.10
185424_s_at	F33D4.6b	1.65
185468_at	GRound-Like (grd related) /// locus:grl-15	1.75
185483_at	C16A11.4a	1.09
185536_at	F26D10.11 /// status:Partially_confirmed	1.52
185593_at	H23N18.5 /// status:Partially_confirmed	2.36
185606_at	F56A4.11 /// status:Predicted /// status:Predicted /// Y19D10A.10	2.60
185608_at	H03E18.1 /// status:Partially_confirmed	1.04
185646_at	GRoundhog (hedgehog-like family) /// locus:grd-14	1.43

185650_at	status:Partially_confirmed /// Y75B8A.34	1.85
185708_s_at	F56A4.10 /// status:Partially_confirmed	2.46
185764_s_at	F33D4.6b	1.67
185795_at	Prion-like-(Q/N-rich)-domain-bearing protein	2.39
185835_at	K08B12.1 /// status:Partially_confirmed	1.06
185862_at	LONg /// locus:lon-8	1.35
185863_at	Serpentine Receptor, class H /// locus:srh-24	1.02
185867_s_at	status:Partially_confirmed /// status:Partially_confirmed /// Y50E8A.11 /// Y75B12A.2	5.67
185916_at	R11G11.6 /// status:Partially_confirmed	1.68
185933_at	F49E10.2a	1.70
185944_at	LIPaSe related /// locus:lips-15	1.28
185949_at	C42D4.13 /// status:Partially_confirmed	1.27
185963_at	status:Confirmed /// Y46D2A.1	1.26
185970_at	status:Partially_confirmed /// ZK1055.4	1.14
185972_at	H42K12.3 /// status:Confirmed	1.51
185984_at	Nematode AStacin protease /// locus:nas-38	1.27
186051_at	C-type LECTin /// locus:clec-78	2.43
186106_at	T14G12.6 /// status:Predicted	2.54
186134_at	status:Partially_confirmed /// Y62H9A.12	1.25
186135_at	C24A11.5 /// status:Predicted	1.58
186180_at	E02C12.6 /// status:Partially_confirmed	2.85
186249_at	C03A7.12 /// status:Partially_confirmed	1.60
186362_s_at	GaLactoFuranose synthesis (UGM, UDP-galactopyranose mutase) /// locus:glf-1	1.21
186403_at	status:Partially_confirmed /// Y49F6C.2	3.37
186406_at	F26D11.4 /// status:Predicted	1.74
186467_at	VC27A7L.1 /// status:Predicted	1.42
186611_at	status:Partially_confirmed /// Y45G12B.3	1.19
186643_at	status:Partially_confirmed /// Y95B8A.11	1.00
186680_s_at	C27A7.2 /// status:Partially_confirmed	1.58
186693_at	C40H1.2 /// status:Partially_confirmed	1.13
186758_at	status:Partially_confirmed /// Y39A3CR.5	1.67
186764_s_at	F16G10.11 /// status:Predicted	3.02
186791_at	R07B1.7 /// status:Predicted	1.14
186814_at	R07B1.6 /// status:Predicted	1.15
186815_s_at	E02C12.10 /// E02C12.11 /// status:Predicted /// status:Partially_confirmed	2.98
186865_at	R07B1.5 /// status:Predicted	1.63
186892_at	C-type LECTin /// locus:clec-133	1.14
186927_s_at	GROund-Like (grd related)	1.98
186998_at	C35A5.5 /// status:Partially_confirmed	1.17
187063_at	status:Partially_confirmed /// Y53G8AR.9	1.89
187075_at	---	1.14
187162_s_at	status:Partially_confirmed /// Y38A10A.2	1.52
187176_s_at	D2096.6 /// status:Partially_confirmed	1.14

187228_at	R07G3.6 /// status:Confirmed	1.75
187251_at	B0353.1b	1.05
187278_at	C28H8.5a	2.80
187306_at	F37A4.3 /// status:Partially_confirmed	1.27
187329_at	T19C3.2 /// status:Partially_confirmed	1.51
187343_at	C01C4.2 /// status:Partially_confirmed	2.02
187347_at	Serpentine Receptor, class A (alpha) /// locus:sra-33	1.09
187353_s_at	PaTched family	1.52
187431_at	C45G9.15 /// status:Partially_confirmed	1.81
187460_at	status:Partially_confirmed /// ZK381.2	1.04
187502_s_at	D1022.3 /// K02A2.1 /// status:Partially_confirmed /// status:Predicted	1.16
187503_at	C14B9.3 /// status:Confirmed	1.38
187509_at	B0280.7 /// status:Confirmed	2.22
187533_at	status:Partially_confirmed /// Y55F3AM.14	1.70
187628_s_at	C30G12.6a	1.12
187630_at	C30G12.1 /// status:Predicted	1.66
187738_at	---	1.02
187753_at	C41A3.2a	1.30
187859_s_at	---	1.05
187885_at	C02D5.1 /// locus:acdh-6	1.79
187901_s_at	Male ABnormal /// locus:mab-7	1.85
187906_at	abnormal DAuer Formation /// locus:daf-6	1.17
187930_at	TWiK family of potassium channels /// locus:twk-6	1.86
188037_at	locus:wrt-6 /// WaRThog (hedgehog-like family)	1.13
188061_s_at	locus:wrt-2 /// WaRThog (hedgehog-like family)	1.13
188102_s_at	PaTched Related family /// locus:ptr-2	1.27
188116_at	Onchocerca Related Antigen family /// locus:ora-1	2.24
188144_s_at	PUF (Pumilio/FBF) domain-containing /// locus:puf-8	1.12
188188_s_at	PaTched Related family /// locus:ptr-18	1.95
188217_at	PaTched Related family /// locus:ptr-22	1.34
188224_s_at	C52A10.2 /// carboxylesterase /// carboxylesterase	2.03
188240_at	PaTched Related family /// locus:ptr-6	1.10
188306_at	Guanylyl CYclase /// locus:gy-25	1.42
188307_at	PaTched Related family /// locus:ptr-20	1.11
188310_at	Serpentine Receptor, class D (delta) /// locus:srd-2	2.48
188312_at	F57B7.3 /// locus:col-156	1.51
188331_s_at	C39E9.3 /// locus:col-131	2.03
188368_at	ZK836.1 /// locus:lon-3	1.30
188404_at	W06B11.4 /// locus:hog-1	1.13
188461_s_at	F29B9.9 /// locus:col-111	1.85
188524_at	M28.1 /// locus:cutl-9	1.23
188537_s_at	ZK20.6 /// locus:nep-1	1.59
188589_at	F33D11.3 /// locus:col-54	1.66
188610_at	Nuclear Hormone Receptor family /// locus:nhr-238	1.26
188611_s_at	Fem-3 mRNA Binding Factor /// locus:fbf-2	1.18

188618_at	Nuclear Hormone Receptor family /// locus:nhr-237	2.55
188659_at	Y38C1BA.3 /// locus:col-109	1.30
188705_s_at	T06D8.10 /// peroxidase	2.31
188747_at	Y54E10BL.2 /// locus:col-48	2.06
188758_at	F07H5.8 /// keratin like	1.13
188830_at	Seven TM Receptor /// locus:str-183	1.98
ã	R08H2.1 /// locus:dhs-23	10.31
188921_at	F25D1.5 /// dehydrogenase	7.72
188932_at	D1054.8 /// dehydrogenase	1.49
188960_s_at	C54G4.4 /// sushi repeats	1.12
188962_at	UDP-GlucuronosylTransferase /// locus:ugt-8	1.26
188987_at	F20G2.1 /// dehydrogenase	2.58
189000_at	T01G6.1 /// dehydrogenase	2.94
189031_at	Nuclear Hormone Receptor family /// locus:nhr-142	1.38
189082_at	C06E4.3 /// dehydrogenase	4.01
189142_at	CYtochrome P450 family /// locus:cyp-29A4	2.11
189174_at	K09C8.3 /// locus:nas-10	2.59
189192_at	C23H4.4 /// carboxyesterase	1.42
189219_s_at	CYtochrome P450 family /// CYtochrome P450 family /// locus:cyp-13A11 /// locus:cyp-13A12	1.87
189221_at	CYtochrome P450 family /// locus:cyp-37B1	1.56
189278_s_at	T21D11.1 /// zinc protease	5.81
189282_at	CYtochrome P450 family /// locus:cyp-35C1	3.11
189283_s_at	CYtochrome P450 family /// locus:cyp-35C1	2.51
189292_at	CYtochrome P450 family /// locus:cyp-25A6	1.73
189297_at	CYtochrome P450 family /// locus:cyp-33E2	1.39
189305_at	F55G1.5 /// carrier protein	1.53
189313_at	K03F8.2 /// locus:acr-5	2.71
189326_at	Serpentine Receptor, class J /// locus:srj-53	1.02
189336_at	CYtochrome P450 family /// locus:cyp-14A2	5.67
189378_at	CYtochrome P450 family /// locus:cyp-14A3	5.44
189405_at	CYtochrome P450 family /// locus:cyp-35D1	4.06
189443_at	T19B10.9 /// max protein like	1.08
189455_at	C27A2.4 /// acid phosphotase	1.74
189457_at	CYtochrome P450 family	3.16
189458_s_at	T07F10.1a	1.37
189481_at	CYtochrome P450 family /// locus:cyp-34A5	2.15
189512_at	CYtochrome P450 family /// locus:cyp-35A2	1.60
189521_s_at	T06A4.3a	1.18
189524_at	B0238.13 /// carboxylesterase	1.49
189549_at	T06A4.1b	1.06
189556_at	CYtochrome P450 family /// locus:cyp-33C2	3.28
189633_at	CYtochrome P450 family /// locus:cyp-34A10	8.81
189644_at	C03B1.5 /// protein kinase	1.02
189747_at	F14E5.1 /// glucose transporter	1.42

189752_s_at	R53.2 /// thymidylate kinase	1.32
189755_at	W01A11.1 /// epoxide hydrolase	1.94
189772_at	CaeNaCin (Caenorhabditis bacteriocin) /// R09B5.11 /// sugar transporter /// locus:cnc-11	1.01
189797_s_at	COLagen /// locus:col-104	1.70
189810_at	CYtochrome P450 family /// locus:cyp-13A10	2.33
189814_s_at	COLagen /// locus:col-173	1.58
189847_at	R12A1.3 /// protease inhibitor	1.83
189869_at	F56C11.6a	1.04
189891_at	M02G9.1 /// keratin like protein	2.28
189926_at	T07D10.3 /// zinc finger protein	1.33
189952_at	R08B4.1b	1.36
189964_at	Nuclear Hormone Receptor family /// locus:nhr-106	1.52
189987_s_at	Nematode AStacin protease /// locus:nas-36	1.09
190002_at	C33C12.8 /// glucosylceramidase	4.51
190020_at	T23F2.1 /// locus:bus-8	1.60
190032_s_at	Nematode AStacin protease	1.16
190048_s_at	CathePsin Z /// locus:cpz-1	1.02
190091_at	SQuaT /// locus:sqt-2 /// status:Confirmed	1.87
190104_at	CUTiclin /// locus:cut-4	1.52
190134_s_at	T10B10.1 /// locus:col-41	2.02
190188_s_at	Fem-3 mRNA Binding Factor /// Fem-3 mRNA Binding Factor /// locus:fbf-1 /// locus:fbf-2	1.34
190193_s_at	R11A5.7 /// zinc carboxypeptidase	1.50
190213_s_at	C16C8.2 /// peroxidase precursor	2.23
190254_at	SKN-1 Dependent Zygotic transcript /// locus:sdz-8	1.41
190298_s_at	DeHydrogenases, Short chain /// locus:dhs-14	1.89
190306_s_at	locus:wrt-8 /// WaRThog (hedgehog-like family)	1.08
190312_at	F20H11.5 /// D-amino acid oxidase	1.16
190318_at	R05D8.9 /// alcohol dehydrogenase	8.31
190326_at	CaDHerin family /// locus:cdh-9	1.35
190340_s_at	locus:wrt-4 /// WaRThog (hedgehog-like family)	2.35
190360_s_at	DumPY : shorter than wild-type /// locus:dpy-5	1.57
190386_at	T13H5.3 /// chitinase-like repeats	1.57
190388_at	T16G12.7 /// Protein phosphatase 1	1.23
190425_at	T05E11.6 /// haemoglobinase like	1.08
190461_at	F28H7.2 /// glucose-1-dehydrogenase	6.36
190479_s_at	QUAhog (hedgehog related) /// locus:qua-1	1.00
190563_at	F39D8.3 /// carbonic anhydrase like	1.10
190570_at	F52D1.1 /// acid alpha-glucosidase	1.05
190580_at	T09B4.6 /// SER/THR-protein kinase	1.22
190585_at	OV-17 antigen precursor /// ZK970.7	1.21
190601_at	C05C9.1 /// LBP / BPI / CETP family	1.44
190615_s_at	F13B12.4 /// cysteine synthase like	1.20
190643_at	H37N21.1 /// tyrosine-protein kinase	1.09
190711_at	Germ-Line Helicase /// locus:glh-1	1.08

190713_at	Fungus-Induced Protein /// locus:fip-1	1.94
190747_s_at	F53B2.2 /// Y45F10B.1 /// locus:tsp-4 /// locus:tsp-5	1.69
190760_at	B0395.2 /// Sterol O-acyltransferase	1.16
190766_s_at	Nematode AStacin protease /// locus:nas-15	2.07
190841_at	Serpentine Receptor, class D (delta) /// locus:srd-8	1.68
190849_at	UDP-GlucuronosylTransferase /// locus:ugt-41	1.49
190861_at	Nematode AStacin protease /// locus:nas-4	1.08
190877_at	Nematode AStacin protease /// locus:nas-7	1.29
190879_at	AC3.7 /// locus:ugt-1	2.56
190895_at	AC3.8 /// locus:ugt-2	1.03
190901_s_at	K04H4.2a	1.61
190907_at	serine protease inhibitor /// ZK287.4	2.13
190916_at	C23G10.6 /// UDP-glucosyltransferase	1.98
190929_at	Glutathione S-Transferase /// locus:gst-14	1.65
190954_at	SOrbitol DeHydrogenase family /// locus:sodh-2	1.37
190973_at	---	1.40
190979_at	Glutathione S-Transferase /// locus:gst-12	1.59
190981_at	TransThyretin-Related family domain /// locus:ttr-12	2.05
191001_s_at	H10E21.4 /// calcium-binding protein	2.16
191036_at	LIPaSe related /// locus:lips-3	2.35
191042_at	Nuclear Hormone Receptor family /// locus:nhr-181	1.09
191066_s_at	UDP-GlucuronosylTransferase	1.74
191137_at	Serpentine Receptor, class X	1.22
191180_at	locus:wrt-1 /// WaRThog (hedgehog-like family)	1.70
191217_s_at	F30H5.3 /// Serine protease inhibitor	1.70
191255_s_at	INneXin	1.25
191307_at	Glutathione S-Transferase /// locus:gst-31	1.43
191319_at	Serpentine Receptor, class H /// locus:srh-204	1.39
191337_at	Glutathione S-Transferase /// locus:gst-39	1.30
191414_s_at	Glutathione S-Transferase /// locus:gst-33	4.49
191418_at	ZC443.6 /// locus:ugt-16	2.21
191466_at	C35A5.2 /// locus:ugt-33	3.05
191500_at	Serpentine Receptor, class H /// locus:srh-270	4.13
191522_s_at	Osmotic Stress Resistant /// locus:osr-1	1.24
191532_at	C26C6.8 /// tyrosine-protein kinase like	1.28
191599_at	T25D1.3 /// cleavage stimulation factor	1.74
191611_at	CaDmium Responsive /// locus:cdr-1	1.45
191612_at	F28D1.9 /// fatty acid transport protein	1.38
191688_at	F15B9.8 /// Thrombospondin type 1 domain	1.62
191698_at	fatty acid ELongation /// locus:elo-4	1.17
191720_at	MAP kinase kinase or Erk Kinase	1.09
191776_at	UDP-GlucuronosylTransferase	3.40
191797_at	status:Partially_confirmed /// ZC449.2	1.03
191811_at	ImmunoGlobulin-like Cell adhesion Molecule family /// locus:igcm-2	1.19
191829_at	F18A11.2 /// retinal-binding like protein	1.08

191837_at	K04A8.10 /// UDP-glucuronosyltransferase	1.70
191846_s_at	SQuashed Vulva /// locus:sqv-4	1.01
191871_s_at	CaDmium Responsive /// locus:cdr-4	3.46
191963_at	SuRFace antigenicity abnormal /// locus:srf-3	1.23
192026_at	COLlagen	1.97
192036_at	SPiNdle orientation defective /// locus:spn-4	1.01
192038_at	C06H2.3 /// Acetyltransferase (GNAT) family	1.41
192089_s_at	F13H8.5 /// Phospholipase ADRAB-B (Rabbit)	1.74
192153_at	M153.1 /// pyrroline-5-carboxylate reductase	1.37
192252_at	F18F11.4 /// serine/threonine-protein kinase	1.21
192391_at	Nuclear Hormone Receptor family /// locus:nhr-248	1.05
192447_at	Nematode AStacin protease	1.32
192448_at	F53F8.1 /// locus:klf-2	1.07
192497_at	Nuclear Hormone Receptor family /// locus:nhr-205	2.13
192605_at	STeroid DeHydrogenase family /// locus:stdh-2	3.73
192607_at	Bacterially Un-Swollen (M. nematophilum resistant)	1.16
192610_at	Nematode AStacin protease /// locus:nas-12	2.34
192671_at	Stress Induced Protein /// locus:sip-1	1.32
192672_s_at	Stress Induced Protein /// locus:sip-1	1.10
192674_s_at	SQuaT /// locus:sqt-1	1.09
192682_at	ROlLer: helically twisted, animals roll when moving /// locus:rol-6	1.53
192820_at	Glutathione S-Transferase /// locus:gst-5	1.90
192948_at	Glutathione S-Transferase /// locus:gst-7	1.11
192967_at	ROlLer: helically twisted, animals roll when moving	1.24
192968_s_at	ROlLer: helically twisted, animals roll when moving	1.33
192970_at	T14E8.4 /// status:Partially_confirmed	1.38
193007_s_at	P-GlycoProtein related /// locus:pgp-1	1.30
193147_at	F01G10.10 /// lipopolysaccharide-binding protein like	1.26
193258_at	PHarynx-associated GAS (growth arrest protein) related /// locus:phg-1	1.71
193270_at	Fatty Acid/Retinol binding protein /// locus:far-4	1.74
193272_s_at	Fatty Acid/Retinol binding protein /// locus:far-3	3.07
193327_at	F53B1.4 /// thymidine diphosphoglucose 4,6-dehydratase	2.21
193330_at	Serpentine Receptor, class SX /// locus:srsx-19	1.01
193338_at	Homolog Of Presenilin /// locus:hop-1	1.21
193409_at	LRP X(cross)-hybridizing /// locus:lrx-1	1.99
193460_at	Zinc finger present in dystrophin, CBP/p300 /// Y32B12A.1	1.35
193466_s_at	CHiTinase /// locus:cht-1	1.12
193477_s_at	E01G6.1 /// pancreatic trypsin inhibitor (kunitz) protein	1.44
193543_at	DeHydrogenases, Short chain	1.46
193573_at	F26D2.15 /// Alcohol/other dehydrogenases, short chain type	3.69
193595_at	THaumatIN family /// locus:thn-1	1.03
193601_at	K10H10.6 /// Alcohol/other dehydrogenases, short chain type	2.41
193677_at	F32D8.7b	1.14
193686_at	See oma /// locus:moe-3	1.91

193712_s_at	C14F11.6 /// DTDP-4-dehydrorhamnose 3,5-epimerase (RFBC gene)	1.16
193735_at	B0334.6 /// Gonadotropin-releasing hormone receptor like protein	1.03
193864_at	Cell Death abnormality /// locus:ced-9	1.33
193885_at	B0393.5 /// EGF-like aspartic acid and asparagine hydroxylation sites	1.13
193903_at	P-GlycoProtein related /// locus:pgp-9	2.09
193991_at	F41G3.3 /// calcium channel protein alpha-1 chain isoform (PIR:A41098) (weak)	2.03
194146_x_at	F49E2.1b	1.33
194213_x_at	H31G24.4 /// locus:cyb-2.2	1.02
194238_x_at	MoLTing defective	1.39

表 2. ケルセチンの 24 時間暴露により発現が誘導した線虫の遺伝子群

Probe Set ID	Gene Title	Log2 Ratio
171743_x_at	Chondroitin ProteoGlycan	1.04
171745_x_at	PaTched Related family /// locus:ptr-11	1.03
171795_x_at	R12E2.14 /// status:Confirmed	1.90
171796_x_at	C02E7.6 /// status:Confirmed	1.56
171815_x_at	OligoPeptide Transporter /// locus:opt-1	1.21
171970_x_at	H39E23.3 /// status:Partially_confirmed	1.39
172054_at	---	4.28
172055_x_at	HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// HIStone /// locus:his-3 /// locus:his-4 /// locus:his-7 /// locus:his-8 /// locus:his-12 /// locus:his-16 /// locus:his-19 /// locus:his-20 /// locus:his-21 /// locus:his-22 /// locus:his-29 /// locus:his-30 /// locus:his-33 /// locus:his-34 /// locus:his-43 /// locus:his-51 /// locus:his-52 /// locus:his-53 /// locus:his-54 /// locus:his-57 /// locus:his-65	1.00
172191_x_at	C27C7.2 /// status:Predicted /// ZK1025.4a	1.36
172196_x_at	COLlagen /// locus:col-65	1.59
172293_x_at	status:Partially_confirmed /// status:Predicted /// Y26D4A.17 /// Y26D4A.5	1.01
172346_x_at	R12E2.7 /// status:Confirmed	1.20
172353_x_at	C02E7.6 /// status:Confirmed	1.46
172387_x_at	---	1.01
172390_x_at	SKN-1 Dependent Zygotic transcript /// locus:sdz-37	1.14
172448_x_at	C29F9.4 /// status:Partially_confirmed	2.33
172497_x_at	R12E2.15 /// status:Confirmed	1.98
172547_x_at	R12E2.14 /// status:Confirmed	1.53
172791_x_at	K08C9.7 /// Ubiquitin family	1.59
172799_x_at	F53F1.5 /// cuticlin	1.62
172824_x_at	HIStone /// locus:his-4	1.10
172851_x_at	HIStone /// HIStone /// locus:his-29 /// locus:his-34	1.23
172976_s_at	UNCoordinated	1.12
172989_s_at	R11A5.7 /// zinc carboxypeptidase	1.42
173004_s_at	GROund-Like (grd related)	1.77
173014_at	ZK20.6 /// locus:nep-1	1.12
173030_at	BE10.2 /// status:Partially_confirmed	1.44
173083_s_at	H10E21.4 /// calcium-binding protein	2.69
173123_at	Cell Division Cycle related	2.53
173132_s_at	T06A1.1 /// status:Partially_confirmed	1.39
173235_at	Y38C1BA.3 /// locus:col-109	2.01
173261_s_at	B0507.1 /// status:Partially_confirmed	2.12
173273_at	SYnthetic lethal with Mec /// locus:sym-1	1.32
173494_s_at	T20F5.4 /// status:Confirmed	1.13
173567_s_at	GROund-Like (grd related)	1.92

173576_at	ZK180.5c	1.63
173584_s_at	Prion-like-(Q/N-rich)-domain-bearing protein /// locus:pqn-73	1.40
173654_at	ACE(angiotensin converting enzyme)-like Non-peptidase	1.02
173661_s_at	Lin-12 and Glp-1 X-hybridizing	1.19
173664_at	Male ABnormal /// locus:mab-7	2.02
173900_s_at	B0331.2 /// phosphate permease like	1.01
173947_at	F13H8.5 /// Phospholipase ADRAB-B (Rabbit)	2.61
174011_at	ForKHead transcription factor family	1.04
174044_at	K04H4.2a	1.07
174071_at	Prion-like-(Q/N-rich)-domain-bearing protein /// locus:pqn-95	1.42
174075_at	---	1.42
174178_at	Prion-like-(Q/N-rich)-domain-bearing protein	1.71
174256_at	Lin-12 and Glp-1 X-hybridizing	1.29
174340_at	status:Partially_confirmed /// Y18H1A.8	1.22
174378_at	T06A4.1b	1.34
174448_s_at	F28D1.9 /// fatty acid transport protein	1.06
174508_s_at	K08E7.5c	1.55
174650_at	F49E10.2a	1.32
174657_s_at	abnormal CHEmotaxis /// locus:che-14	1.03
174660_at	T19A5.3b	1.16
174661_s_at	T19A5.3b	1.28
174960_at	ZK1010.7 /// locus:col-97	1.43
174995_s_at	PaTched Related family /// locus:ptr-6	1.05
175008_at	F10D11.6 /// status:Partially_confirmed	2.24
175093_at	C26B9.7 /// status:Confirmed	1.18
175204_at	status:Partially_confirmed /// ZK829.3	1.52
175294_s_at	T17H7.7 /// status:Partially_confirmed	1.55
175393_at	status:Partially_confirmed /// Y47D7A.13	1.71
175576_s_at	status:Confirmed /// Y75B7AR.1	1.15
175642_s_at	AntiBacterial Factor related /// locus:abf-2	1.09
176074_s_at	status:Partially_confirmed /// Y32G9B.1	1.49
176188_at	CaDHerin family /// locus:cdh-12	1.05
176274_at	status:Partially_confirmed /// Y52E8A.4	1.07
176293_at	status:Partially_confirmed /// Y71G12B.17	1.31
176327_s_at	GRound-Like (grd related) /// locus:grl-5	1.75
176330_at	status:Confirmed /// Y22D7AL.14	1.44
176497_s_at	Y54F10AL.1a	1.22
176672_at	W08G11.1 /// status:Partially_confirmed	1.50
176674_at	Serpentine Receptor, class XA /// locus:srxa-5	1.28
176813_at	Serpentine Receptor, class W /// locus:srw-122	3.09
176857_s_at	SaPosin-like Protein family /// locus:spp-7	1.30
177001_at	UBX-containing protein in Nematode	1.89
177024_at	COLlagen /// locus:col-90	1.24
177049_s_at	status:Partially_confirmed /// status:Partially_confirmed /// Y73B6BL.23 /// ZK616.5	1.28

177085_at	status:Partially_confirmed /// Y55F3BR.2	1.25
177106_at	status:Partially_confirmed /// Y61A9LA.7	1.14
177113_at	Prion-like-(Q/N-rich)-domain-bearing protein /// locus:pqn-26	2.19
177135_at	status:Partially_confirmed /// Y48G1BR.1	1.22
177247_at	BE10.2 /// status:Partially_confirmed	1.08
177270_at	JC8.12a	1.28
177353_s_at	C-type LECTin /// locus:clcc-190	2.26
177359_at	status:Partially_confirmed /// Y97E10AM.1	1.36
177505_at	T06G6.6b	1.40
177581_at	BListered cuticle /// locus:bli-5	1.00
177586_at	C15C6.1 /// status:Confirmed	1.29
177687_at	K09G1.2 /// status:Partially_confirmed	1.20
177696_at	C30H6.5 /// status:Partially_confirmed	1.45
177801_at	C-type LECTin /// locus:clcc-230	1.14
177832_at	K02E2.8b	2.06
177870_at	C02F4.4 /// status:Confirmed	1.61
177929_s_at	T22H2.2	1.10
177942_at	F33A8.7 /// status:Confirmed	1.77
178012_at	Seven TM Receptor /// locus:str-97	1.08
178038_at	T23F6.1 /// status:Confirmed	1.11
178111_at	F59F5.7 /// status:Partially_confirmed	1.32
178165_at	F46F3.3 /// status:Confirmed	1.44
178168_at	T01G9.1 /// status:Partially_confirmed	1.72
178189_at	F28C6.5 /// status:Partially_confirmed	1.11
178194_at	R07E3.6 /// status:Partially_confirmed	1.01
178197_at	Serpentine Receptor, class E (epsilon) /// locus:sre-32	1.48
178204_at	T23F6.2 /// status:Predicted	1.04
178244_at	F01D5.6 /// status:Partially_confirmed	1.47
178285_at	---	2.68
178330_at	C54C8.2 /// status:Partially_confirmed	1.96
178333_at	K03H4.2 /// status:Partially_confirmed	3.84
178343_s_at	LIPaSe related /// locus:lips-9	1.99
178372_at	status:Partially_confirmed /// ZK829.3	1.56
178476_at	status:Partially_confirmed /// ZC434.3	1.39
178508_at	C14A4.9 /// status:Confirmed	1.06
178693_at	M151.3 /// status:Predicted	1.10
178715_at	Ligand-Gated ion Channel /// locus:lge-21	1.18
178716_at	AuTophagy (yeast Atg homolog) /// locus:atg-10	1.08
178728_at	R01E6.2 /// status:Partially_confirmed	1.25
178861_at	K02D3.2 /// status:Partially_confirmed	1.06
178912_at	F35G2.5a	1.11
178929_at	F47G9.3 /// locus:cutl-18	1.21
178979_at	K01C8.2 /// status:Partially_confirmed	1.21
178985_at	C04B4.1 /// status:Partially_confirmed	1.87
179020_at	C08E8.3 /// status:Partially_confirmed	1.22

179027_at	W02D9.4 /// status:Predicted	1.12
179034_s_at	M03B6.3 /// status:Confirmed	1.07
179131_at	T08G5.3 /// status:Confirmed	1.76
179159_at	C02B4.3 /// C02B4.4 /// status:Predicted /// status:Confirmed	2.16
179176_at	C52G5.2 /// status:Confirmed	1.20
179246_s_at	F54B8.1 /// M01B2.8 /// status:Partially_confirmed /// status:Partially_confirmed	1.50
179249_at	LIpaSe related /// locus:lips-6	2.60
179254_at	locus:zbed-6 /// Zinc finger protein (BED class)	1.91
179277_at	F15D4.3 /// status:Confirmed	1.39
179333_at	GRoundhog (hedgehog-like family) /// locus:grd-12	1.14
179335_at	K08E7.5c	1.72
179401_at	status:Confirmed /// ZK662.2	1.89
179600_at	C38C6.3 /// status:Confirmed	1.33
179741_at	T11F8.2 /// status:Partially_confirmed	1.14
179749_at	Serpentine Receptor, class H /// locus:srh-32	1.56
179830_at	F01D5.10 /// status:Partially_confirmed	1.05
179868_at	status:Partially_confirmed /// Y81G3A.4	1.13
180041_at	status:Partially_confirmed /// Y49A3A.4	1.52
180056_at	F31A3.3 /// status:Predicted	1.39
180136_at	status:Predicted /// ZK488.6	2.18
180189_s_at	T06A1.1 /// status:Partially_confirmed	1.07
180200_at	T13C2.2 /// status:Confirmed	1.20
180227_at	F21H7.10 /// status:Partially_confirmed	1.87
180277_at	T22D1.1 /// status:Predicted	1.41
180286_at	W03D8.11 /// status:Partially_confirmed	1.20
180418_at	T19D7.3 /// status:Confirmed	1.27
180462_at	C07E3.10 /// status:Confirmed	1.43
180498_at	F22E12.3 /// status:Partially_confirmed	1.22
180505_at	Prion-like-(Q/N-rich)-domain-bearing protein	1.66
180506_at	Seven TM Receptor /// locus:str-158	3.82
180524_at	T22B2.6 /// status:Confirmed	2.04
180605_s_at	C39E9.11 /// status:Confirmed	1.25
180614_at	T17H7.7 /// status:Partially_confirmed	1.40
180624_s_at	Prion-like-(Q/N-rich)-domain-bearing protein /// locus:pqn-40	1.59
180647_at	F53B1.6 /// status:Partially_confirmed	1.19
180664_s_at	status:Partially_confirmed /// ZK180.6	2.11
180671_at	CYtochrome P450 family /// locus:cyp-33E3	1.28
180704_at	F56C3.9 /// status:Confirmed	1.02
180741_at	F36H9.5 /// status:Partially_confirmed	1.12
180761_at	F42A9.3 /// status:Partially_confirmed	2.82
180841_at	C32H11.5 /// status:Confirmed	1.24
180846_at	C27D9.2 /// status:Confirmed	1.19
180848_at	C09D4.6 /// status:Partially_confirmed	1.91
180882_s_at	F18C5.5 /// status:Confirmed	1.26

180895_at	Delta and OSM-11-like /// locus:dos-2	1.15
180922_at	C52B9.2b	1.74
180930_s_at	F25E2.2 /// status:Partially_confirmed	1.05
180948_at	F52E1.5 /// status:Partially_confirmed	1.16
181113_at	F40E10.5 /// status:Confirmed	1.54
181133_at	F-box B protein /// locus:fbxb-78	1.04
181151_at	C26B9.2 /// status:Predicted	1.16
181182_at	R12E2.8 /// status:Partially_confirmed	1.13
181185_at	Serpentine Receptor, class H /// locus:srh-283	1.24
181253_at	T09B4.3 /// status:Partially_confirmed	1.07
181302_s_at	BTB and MATH domain containing /// BTB and MATH domain containing /// locus:bath-30	1.05
181317_at	C17F4.2 /// status:Partially_confirmed	1.35
181326_at	Ligand-Gated ion Channel /// locus:lgc-22	1.90
181343_at	C17F4.3 /// status:Confirmed	1.77
181410_s_at	F26G1.5 /// status:Partially_confirmed	1.54
181415_at	B0507.1 /// status:Partially_confirmed	1.14
181440_at	D1014.5 /// status:Confirmed	1.24
181523_s_at	F18A1.7 /// status:Confirmed	1.23
181544_at	status:Partially_confirmed /// Y53F4B.6	1.10
181586_at	F42C5.3 /// status:Partially_confirmed	1.02
181631_at	---	2.66
181638_at	Serpentine Receptor, class X /// locus:srx-108	1.38
181649_at	F48E3.9 /// status:Partially_confirmed	2.03
181681_at	SaPosin-like Protein family /// locus:spp-13	1.19
181707_at	F10D11.6 /// status:Partially_confirmed	1.65
181722_s_at	T01D1.4 /// status:Confirmed	1.54
181793_at	MoLTing defective /// locus:mlt-10	1.32
181872_at	Suppressor/Enhancer of Lin-12 /// locus:sel-7	1.06
181887_at	C01G12.7 /// status:Partially_confirmed	1.21
181890_at	C55B6.4 /// status:Predicted	1.37
181956_at	LIPaSe related	1.20
181963_at	C23H3.9d	1.01
182085_at	C23H3.9d	1.21
182104_at	status:Confirmed /// ZK1025.3	1.24
182111_at	D2024.4a	1.54
182305_at	F47F2.3 /// status:Partially_confirmed	3.27
182308_at	C11G10.2 /// status:Partially_confirmed	1.30
182346_at	status:Partially_confirmed /// ZC123.1	1.55
182356_at	GLoBin /// locus:glb-25	3.34
182419_at	F-box A protein /// locus:fbxa-203	1.56
182428_s_at	LYSozyme /// locus:lys-3	2.57
182431_at	T06C12.8 /// status:Partially_confirmed	2.97
182436_at	H19J13.1 /// locus:otpl-8	1.01
182455_at	F39G3.6 /// status:Predicted	3.36

182490_at	status:Partially_confirmed /// ZK470.6	2.35
182561_at	status:Partially_confirmed /// Y11D7A.9	1.53
182629_at	C50D2.1 /// status:Partially_confirmed	1.34
182639_at	C25F6.6 /// status:Predicted	1.67
182683_at	status:Partially_confirmed /// ZC250.3	1.23
182696_s_at	Chondroitin ProteoGlycan	1.36
182776_at	status:Confirmed /// Y45F3A.8	2.12
182987_at	W03B1.2 /// status:Partially_confirmed	1.15
182988_at	C29G2.3 /// status:Partially_confirmed	1.19
183147_at	C06G1.2 /// status:Confirmed	1.81
183159_at	F09F9.2 /// status:Confirmed	1.24
183176_at	T07C12.8 /// status:Predicted	1.56
183185_at	F15H10.8 /// status:Partially_confirmed	1.29
183191_at	F38B6.7 /// status:Partially_confirmed	1.12
183230_at	R02E4.1 /// status:Predicted	1.27
183237_s_at	ZK180.5c	2.05
183258_at	F38B6.2 /// status:Partially_confirmed	1.07
183328_at	status:Confirmed /// Y38F1A.1	1.43
183379_at	K01D12.9 /// status:Confirmed	2.24
183407_at	F40H3.3 /// status:Partially_confirmed	1.33
183415_at	K01D12.3 /// status:Partially_confirmed	3.23
183454_at	F38B6.1 /// status:Partially_confirmed	1.09
183480_at	C32H11.6 /// status:Predicted	1.33
183496_at	C26B2.8 /// status:Partially_confirmed	2.85
183537_at	C03G6.6 /// status:Partially_confirmed	1.41
183539_s_at	C06E7.4 /// status:Confirmed	1.37
183561_s_at	F40F12.7 /// status:Confirmed	1.01
183577_at	Prion-like-(Q/N-rich)-domain-bearing protein /// locus:pqn-95	1.99
183582_at	C47E12.6a	2.01
183601_at	C06E7.4 /// status:Confirmed	1.88
183676_s_at	NUclear localized Metal Responsive /// NUclear localized Metal Responsive /// locus:numr-2 /// locus:numr-1	1.57
183717_at	F20B10.3 /// status:Partially_confirmed	1.07
183724_at	status:Confirmed /// Y11D7A.5	1.21
183731_at	K01D12.1 /// status:Partially_confirmed	2.31
183733_at	F44C8.7 /// status:Confirmed	1.44
183735_at	K03H9.3 /// status:Predicted	1.25
183755_s_at	F46C8.8 /// status:Confirmed	1.94
183794_at	status:Partially_confirmed /// ZK1225.1	1.00
183818_s_at	C05E7.1 /// status:Partially_confirmed	1.49
183907_at	status:Partially_confirmed /// ZK1025.7	1.69
183911_at	T19A5.3b	1.17
183941_s_at	F56D3.1 /// status:Confirmed	1.01
183975_s_at	Prion-like-(Q/N-rich)-domain-bearing protein /// locus:pqn-46	1.42
184091_at	GRound-Like (grd related) /// locus:grl-10	1.33

184102_s_at	status:Predicted /// status:Predicted /// ZK1025.2 /// ZK1025.8	1.23
184106_at	M03F8.1 /// status:Partially_confirmed	3.50
184125_at	C05D12.1 /// status:Partially_confirmed	1.19
184138_at	K02E7.1a	1.75
184140_at	F49E8.2 /// status:Partially_confirmed	1.13
184151_at	F26A1.9 /// status:Partially_confirmed	1.85
184194_s_at	status:Partially_confirmed /// ZK1025.7	1.40
184199_s_at	C04F1.1 /// status:Partially_confirmed	1.81
184205_at	status:Partially_confirmed /// Y52B11B.1	1.90
184220_at	C56C10.4 /// status:Confirmed	1.20
184236_at	T24C12.4 /// status:Partially_confirmed	1.75
184254_at	C46H11.7 /// status:Partially_confirmed	1.37
184268_s_at	PHaryngeal gland Toxin-related /// locus:phat-2	1.30
184335_at	DumPY : shorter than wild-type /// locus:dpy-1	1.06
184352_at	C17H12.6 /// status:Partially_confirmed	1.09
184601_at	DNaJ domain (prokaryotic heat shock protein) /// locus:dnj-26	1.07
184662_at	status:Partially_confirmed /// Y46G5A.29	1.53
184682_s_at	K09B11.10 /// status:Partially_confirmed	1.98
184684_at	Y53F4B.27a	1.31
184693_s_at	C32H11.11 /// C32H11.8 /// status:Predicted /// status:Predicted	1.67
184810_at	C01B10.7 /// status:Predicted	2.55
184828_s_at	MoLTing defective	1.39
184957_at	Seven TM Receptor /// locus:str-159	1.40
185108_at	F36F12.1 /// status:Partially_confirmed	1.37
185172_s_at	CHitin Synthase /// locus:chs-2	1.22
185231_at	F41E6.14 /// status:Predicted	1.22
185248_s_at	T09A12.1 /// status:Partially_confirmed	1.35
185287_at	status:Partially_confirmed /// Y53F4B.25	1.74
185334_s_at	R02F11.1 /// status:Confirmed	1.14
185337_at	C32B5.12 /// locus:fbxc-57	1.01
185390_at	F49D11.6 /// status:Predicted	1.24
185424_s_at	F33D4.6b	1.45
185468_at	GRound-Like (grd related) /// locus:grl-15	1.25
185593_at	H23N18.5 /// status:Partially_confirmed	1.98
185764_s_at	F33D4.6b	1.44
185771_at	Serpentine Receptor, class U /// locus:sru-1	1.16
185795_at	Prion-like-(Q/N-rich)-domain-bearing protein	1.97
185933_at	F49E10.2a	1.35
185972_at	H42K12.3 /// status:Confirmed	1.13
185985_s_at	F09E10.6 /// status:Confirmed	1.06
186106_at	T14G12.6 /// status:Predicted	2.12
186118_at	F35H12.1 /// status:Predicted	1.03
186134_at	status:Partially_confirmed /// Y62H9A.12	1.40
186171_at	Y44E3A.1a	1.23
186249_at	C03A7.12 /// status:Partially_confirmed	1.44

186357_at	T15B7.17 /// status:Partially_confirmed	3.43
186375_at	F43C11.4 /// status:Predicted	1.21
186505_at	F49H12.3 /// status:Confirmed	2.60
186758_at	status:Partially_confirmed /// Y39A3CR.5	1.22
186764_s_at	F16G10.11 /// status:Predicted	2.48
186814_at	R07B1.6 /// status:Predicted	1.32
186821_at	Serpentine Receptor, class W /// locus:srw-38	2.11
186865_at	R07B1.5 /// status:Predicted	1.21
186886_s_at	status:Partially_confirmed /// Y18D10A.23	1.28
186912_at	status:Confirmed /// Y45F10D.10	1.03
186927_s_at	GRound-Like (grd related)	1.76
186966_at	Helix Loop Helix /// locus:hlh-33	2.15
186998_at	C35A5.5 /// status:Partially_confirmed	1.06
187021_at	C-type LECTin /// locus:clec-171	2.37
187162_s_at	status:Partially_confirmed /// Y38A10A.2	1.22
187174_at	C33H5.2 /// status:Partially_confirmed	1.50
187216_s_at	D1044.2c	1.37
187228_at	R07G3.6 /// status:Confirmed	1.60
187278_at	C28H8.5a	2.44
187329_at	T19C3.2 /// status:Partially_confirmed	1.58
187440_at	MeDiaTor /// R12B2.6 /// status:Predicted	1.01
187509_at	B0280.7 /// status:Confirmed	1.90
187514_at	F52C9.5 /// status:Partially_confirmed	1.30
187570_at	B0310.6 /// status:Partially_confirmed	2.88
187630_at	C30G12.1 /// status:Predicted	1.17
187708_s_at	Suppressor with Morphological effect on Genitalia	3.69
187885_at	C02D5.1 /// locus:acdh-6	1.67
187901_s_at	Male ABnormal /// locus:mab-7	1.34
187911_at	COHesin family /// locus:coh-3	1.32
188102_s_at	PaTched Related family /// locus:ptr-2	1.01
188116_at	Onchocerca Related Antigen family /// locus:ora-1	1.78
188188_s_at	PaTched Related family /// locus:ptr-18	1.54
188257_at	T15H9.1	1.45
188270_at	PaTched Related family /// locus:ptr-16	1.66
188312_at	F57B7.3 /// locus:col-156	1.49
188331_s_at	C39E9.3 /// locus:col-131	2.18
188368_at	ZK836.1 /// locus:lon-3	1.03
188461_s_at	F29B9.9 /// locus:col-111	1.81
188512_at	R09D1.3 /// chitinase	1.13
188527_s_at	ACE(angiotensin converting enzyme)-like Non-peptidase	1.08
188537_s_at	ZK20.6 /// locus:nep-1	1.37
188589_at	F33D11.3 /// locus:col-54	1.27
188611_s_at	Fem-3 mRNA Binding Factor /// locus:fbf-2	1.42
188659_at	Y38C1BA.3 /// locus:col-109	1.09
188679_s_at	F52E1.10 /// locus:vha-18	1.12

188705_s_at	T06D8.10 /// peroxidase	2.36
188747_at	Y54E10BL.2 /// locus:col-48	1.79
189213_at	C56G2.5 /// Protein kinase	1.33
189221_at	Cytochrome P450 family /// locus:cyp-37B1	1.15
189278_s_at	T21D11.1 /// zinc protease	5.20
189443_at	T19B10.9 /// max protein like	1.07
189455_at	C27A2.4 /// acid phosphatase	1.37
189458_s_at	T07F10.1a	1.36
189587_s_at	W03F9.4 /// acetyltransferase	1.12
189600_at	COLlagen /// locus:col-89	2.07
189747_at	F14E5.1 /// glucose transporter	1.29
189797_s_at	COLlagen /// locus:col-104	1.38
189814_s_at	COLlagen /// locus:col-173	1.43
189847_at	R12A1.3 /// protease inhibitor	1.29
189891_at	M02G9.1 /// keratin like protein	1.65
189899_at	Nematode AStacin protease /// locus:nas-14	3.66
189952_at	R08B4.1b	1.05
189987_s_at	Nematode AStacin protease /// locus:nas-36	1.38
190020_at	T23F2.1 /// locus:bus-8	1.31
190091_at	SQuaT /// locus:sqt-2 /// status:Confirmed	1.27
190134_s_at	T10B10.1 /// locus:col-41	1.81
190193_s_at	R11A5.7 /// zinc carboxypeptidase	1.22
190213_s_at	C16C8.2 /// peroxidase precursor	2.26
190306_s_at	locus:wrt-8 /// WaRThog (hedgehog-like family)	1.26
190326_at	CaDHerin family /// locus:cdh-9	1.12
190340_s_at	locus:wrt-4 /// WaRThog (hedgehog-like family)	1.97
190360_s_at	DumPY : shorter than wild-type /// locus:dpy-5	1.42
190386_at	T13H5.3 /// chitinase-like repeats	1.02
190431_at	Pachytene CHeckpoint protein (yeast PCH homolog) /// locus:pch-2	1.11
190479_s_at	QUAhog (hedgehog related) /// locus:qua-1	1.34
190601_at	C05C9.1 /// LBP / BPI / CETP family	1.01
190615_s_at	F13B12.4 /// cysteine synthase like	1.48
190713_at	Fungus-Induced Protein /// locus:fip-1	2.18
190766_s_at	Nematode AStacin protease /// locus:nas-15	1.65
190877_at	Nematode AStacin protease /// locus:nas-7	1.31
190901_s_at	K04H4.2a	1.11
191001_s_at	H10E21.4 /// calcium-binding protein	2.30
191036_at	LIPaSe related /// locus:lips-3	2.14
191042_at	Nuclear Hormone Receptor family /// locus:nhr-181	1.03
191066_s_at	UDP-GlucuronosylTransferase	1.02
191085_at	Serpentine Receptor, class I /// locus:sri-45	1.07
191180_at	locus:wrt-1 /// WaRThog (hedgehog-like family)	1.06
191217_s_at	F30H5.3 /// Serine protease inhibitor	1.15
191311_at	Serpentine Receptor, class H /// locus:srh-60	1.95

191450_at	Seven TM Receptor /// locus:str-28	2.80
191522_s_at	Osmotic Stress Resistant /// locus:osr-1	1.07
191612_at	F28D1.9 /// fatty acid transport protein	1.52
191680_at	Serpentine Receptor, class H /// locus:srh-40	2.59
191688_at	F15B9.8 /// Thrombospondin type 1 domain	1.37
191845_at	C46A5.1 /// protein-tyrosine phosphatase	3.93
191913_s_at	C17H12.3 /// protein-tyrosine phosphatase	1.01
192026_at	COLlagen	1.85
192038_at	C06H2.3 /// Acetyltransferase (GNAT) family	1.08
192089_s_at	F13H8.5 /// Phospholipase ADRAB-B (Rabbit)	1.46
192153_at	M153.1 /// pyrroline-5-carboxylate reductase	1.05
192252_at	F18F11.4 /// serine/threonine-protein kinase	1.85
192385_at	INeXin /// locus:inx-20	1.11
192445_at	Motor AXon guidance	2.16
192610_at	Nematode ASTacin protease /// locus:nas-12	1.97
192682_at	ROlLer: helically twisted, animals roll when moving /// locus:rol-6	1.21
192711_at	IMportin Alpha family /// locus:ima-2	1.03
192967_at	ROlLer: helically twisted, animals roll when moving	1.08
192968_s_at	ROlLer: helically twisted, animals roll when moving	1.10
193147_at	F01G10.10 /// lipopolysaccharide-binding protein like	1.36
193258_at	PHarynx-associated GAS (growth arrest protein) related /// locus:phg-1	1.17
193327_at	F53B1.4 /// thymidine diphosphoglucose 4,6-dehydratase	1.77
193409_at	LRP X(cross)-hybridizing /// locus:lrx-1	1.35
193477_s_at	E01G6.1 /// pancreatic trypsin inhibitor (kunitz) protein	1.22
193601_at	K10H10.6 /// Alcohol/other dehydrogenases, short chain type	1.74
193677_at	F32D8.7b	1.20
193686_at	See oma /// locus:moe-3	1.59
193991_at	F41G3.3 /// calcium channel protein alpha-1 chain isoform (PIR:A41098) (weak)	1.37
194146_x_at	F49E2.1b	1.08
194238_x_at	MoLTing defective	1.36