

T-COFFEE, Version_7.38Thu Nov 20 17:26:38 WEST 2008
Cedric Notredame
CPU TIME:42 sec.
SCORE=94

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BAD AVG GOOD

Arabidopsis_tha	:	94
Cucumis_sativus	:	95
Hordeum_vulgare	:	93
Triticum_aestiv	:	93
Arabidopsis_tha	:	94
Hordeum_vulgare	:	94
Arabidopsis_tha	:	94
Avena_sativa	:	99
Chlamydomonas_r	:	90
Daucus_carota	:	94
Marchantia_pale	:	92
Pisum_sativum	:	94
Plectonema_bory	:	99
Synechocystis_s	:	98
cons	:	94

Arabidopsis_tha	1	MALQAA-S	7
Cucumis_sativus	1	MALQAA-S	7
Hordeum_vulgare	1	MALQ	4
Triticum_aestiv	1	MALQ	4
Arabidopsis_tha	1	MALQAA-S	7
Hordeum_vulgare	1	MALQAATS	8
Arabidopsis_tha	1	MALQAAYS	8
Avena_sativa	1		0
Chlamydomonas_r	1	MALTMSAK	8
Daucus_carota	1	MALQAA-S	7
Marchantia_pale	1	MPKRSNGSLVVRCAVSVVRFSKENVSCDLASENFTFSRDSFPVVS	45
Pisum_sativum	1	MALQTA-S	7
Plectonema_bory	1	MAQD	4
Synechocystis_s	1	MEQP	4
cons	1		45

Arabidopsis_tha	8	LVSSAFSVRKDGKL-NAS-ASSSFKESSL	34
Cucumis_sativus	8	LVSPALSIPKEGKS-S-VCLKDSSL	30
Hordeum_vulgare	5	LLPSTLSVPKKGSS-M-GAVAVKDTAA	29
Triticum_aestiv	5	LLPSTLSVPKKGSS-M-GAAAVKDTAA	29
Arabidopsis_tha	8	LVSSAFSVRKDAKL-NAS-SSSFKDSSL	33
Hordeum_vulgare	9	FLPSALSARKEGA-AKDSAF	27
Arabidopsis_tha	9	LLPSTISIQKEGKF-N-ASLKETT	31
Avena_sativa	1		0
Chlamydomonas_r	9	SVSARAQV-SSKAQAA	23
Daucus_carota	8	FLPSSFSINKEGKA-N-VSLKETSL	30
Marchantia_pale	46	TVLRVSEAVYRMAAVA SLGSALSVSSAALSQNVSVSNNATKESAF	90
Pisum_sativum	8	MLPASFSIPKEGKI-G-ASLKDSTL	30
Plectonema_bory	5		4
Synechocystis_s	5		4
cons	46		90

Arabidopsis_tha	35	F	GVS	LSEQSKADFV	SSSLRCKREQSLRNN	KAIIRAQAIATS	75
Cucumis_sativus	31	F	GIS	FSDHLKSEFS	SSTLRCKRE--LNQQ	IGAIRAQTTATE	69
Hordeum_vulgare	30	FL	GVS	S----	KAKKA-SL-----	AVR-TQVATAPSPVTT	57
Triticum_aestiv	30	FL	GVS	S----	KAKKA-SL-----	AVR-TQVATAPSSVTT	57
Arabidopsis_tha	34	F	GAS	ITDQIKSEHG	SSSLRFKREQS--LR	NLAIRAQTAATS	72
Hordeum_vulgare	28	F	GVR	LADGLKL DAT	SLGLRTRKR---VNTS	SVAIRAQAAAVS	65
Arabidopsis_tha	32	T	GSS	FSNHLRAEKI	STLLTIKEQRRQKPRF	STGIRAQTV-TA	72
Avena_sativa	1						0
Chlamydomonas_r	24	P	AVA	VSGRTSSRVMP	PAPALAARSSV-A-R	TPLVVCAATATAP	63
Daucus_carota	31	F	GVT	FSDSLRTDFS	SLRTRRGCR--QISQ	TGAIRSQAVA-T	68
Marchantia_pale	91	L	GLR	MGEVAKFGGA	LLSVSTVA-ANLKSK	PGVLSV--NAV T	128
Pisum_sativum	31	F	GVS	SLSDSLKGDFT	SSALRCK-E--LRQK	VGAVRAETAAPA	69
Plectonema_bory	5						4
Synechocystis_s	5						4
cons	91						135

Arabidopsis_tha	76	TPSV	TKSS	---	LDRKKT LRKGNVVVTGASSGLGLATAKALAE T	115
Cucumis_sativus	70	SPAV	NKAT	---	PDGKKT LRKGSVVITGASSGLGLATAKALAE T	109
Hordeum_vulgare	58	SPGS	TASS	---	PSGKKT LRQGVVVITGASSGLGLAAAKALAE T	97
Triticum_aestiv	58	SPGS	ATAK	---	PSGKKT LRQGVVVITGASSGLGLAAAKALAE T	97
Arabidopsis_tha	73	SPTV	TK-S	---	VDGKKT LRKGNVVVTGASSGLGLATAKALAE T	111
Hordeum_vulgare	66	APTA	TPAS	---	PAGKKT VRTGNAIITGASSGLGLATAKALAE S	105
Arabidopsis_tha	73	TPPA	NEAS	---	PEQKKT ERKGTAVITGASSGLGLATAKALAD T	112
Avena_sativa	1				---VVVITGASSGLGLAAAKALAE T	22
Chlamydomonas_r	64	SPSLA	DKFK	PNAI	ARVPATQQKQTAIITGASSGLGLNAAKALAA T	108
Daucus_carota	69	TPSV	NRAT	---	GEGKKT LRKGSVIITGASSGLGLATAKALAE T	108
Marchantia_pale	129	APAE	TMNK	---	PSSKKT ATKSTCIITGASSGLGLATAKALAD T	168
Pisum_sativum	70	TPAV	NKSS	---	SEGKKT LRKGNVVITGASSGLGLATAKALAE S	109
Plectonema_bory	5				---QKPTVVITGASSGVGLYAAKALVKR	29
Synechocystis_s	5				---MKPTVIITGASSGVGLYGAKALIDK	29
cons	136				: : * * * * * : * * * * *	180

Arabidopsis_tha	116	GKWHVIMACRDFLKAERAAQSAGMPKDSYTMHLDLASLDSVRQF	160
Cucumis_sativus	110	GKWHVIMACRDFLKAERAAKSAGITKENYTMHLDLASLDSVRQF	154
Hordeum_vulgare	98	GKWHVVIMACRDFLKAASKAAKAAGMADGSYTMHLDLASLDSVRQF	142
Triticum_aestiv	98	GKWHVVIMACRDFLKAASKAAKAAGMADGSYTMHLDLASLDSVRQF	142
Arabidopsis_tha	112	GKWNVIMACRDFLKAERAAKSVGMPKDSYTMHLDLASLDSVRQF	156
Hordeum_vulgare	106	GKWHVIMACRDYLLKTARAARAAGMPKGSYTI VHLDLASLDSVRQF	150
Arabidopsis_tha	113	GKWHVIMACRNFLKA EKAARSVGMSKEDYTMHLDLASLESVKQF	157
Avena_sativa	23	GKWHVVIMACRDFLKAASKAAKAAGMADGSYTMHLDLASLDSVRQF	67
Chlamydomonas_r	109	GEWHVVIMACRDFLKA EQAAKKVGMPAGSYSILHLDLSSLESVRQF	153
Daucus_carota	109	GKWHVIMACRDFLKAERAAKSAGMPKENYTIMHLDLASLDSVRQF	153
Marchantia_pale	169	GEWHVIMACRDFLKAERAA RSVGIPKDSYTVIHCDLASFDSVRAF	213
Pisum_sativum	110	GKWHVIMACRDYLLKAARA A K SAGLAKENYTIMHLDLASLDSVRQF	154
Plectonema_bory	30	G-WHVIMACRNLEKADSA A KSLGMSPDSYTLMHIDLGS LDSVRKF	73
Synechocystis_s	30	G-WHVIMACRNLDKTQKV ADELGF PKDSYTI IKL DLGYLDSVRRE	73
cons	181	* : * : * : * : * : * : . * * : . * : : : * * . : * : * : *	225

Arabidopsis_tha	161	VDNFRRRAEMPLDVLVCNAAVYQPTANQPTFTAEGFELSVGINHLG	205
Cucumis_sativus	155	VDNFRQSGRPLDVLVCNAAVYLPTAKEPTFTAEGFELSVGTNHLG	199
Hordeum_vulgare	143	VDAFRRRAEMPLDVLVCNAAIYRPTARTPTFTADGHEMSVGVNHLG	187
Triticum_aestiv	143	VDAFRRRAEMPLDVLVCNAAIYRPTARTPTFTADGHEMSVGVNHLG	187
Arabidopsis_tha	157	VDNFRRTETPLDVLVCNAAVYFPTAKEPTYSAEGFELSVATNHLG	201
Hordeum_vulgare	151	VKNVRQLDMPIDVVVCNAAVYQPTAKEPSFTADGFEMSVGVNHLG	195
Arabidopsis_tha	158	VENFRRTEQPLDVLVCNAAVYQPTAKEPSFTAEGFEISVGTNHLG	202
Avena_sativa	68	VDAFRRRAEMPLDVLVCNAAIYRPTARKPTFTAEGVEMSVGVNHLG	112
Chlamydomonas_r	154	VQNFKASGRRLDALVCNAAVYLPTAKEPRFTADGFELSVGTNHLG	198
Daucus_carota	154	VETFRRSERPLDVLVCNAAVYFPTAKEPTYTADGFELSVGTNHLG	198
Marchantia_pale	214	VDNFRRTERQLDVLVCNAAVYFPTDKEPKFSAEGFELSVGTNHMG	258
Pisum_sativum	155	VDNFRRSEMPDLVLINNAAVYFPTAKEPSFTADGFEISVGTNHLG	199
Plectonema_bory	74	VTQFRESGKSLDALVCNAAVHMPLLKEPMRSPEGYELSVATNHFG	118
Synechocystis_s	74	VAQFRELGRPLKALVCNAAVYFPLLDEPLWSADDYELSVATNHLG	118
cons	226	* .: :.:: ***** * *:.* **:*	270

Arabidopsis_tha	206	HFLLSRLLLIDDLKNSDYPSKRLLIIVGSITGNTNTLAGNVP--PKA	248
Cucumis_sativus	200	HFLLSRLLLLEDLNKSSYPSKRLLIIVGSITGNTNTLAGNVP--PKA	242
Hordeum_vulgare	188	HFLLARLLMEDLQKSDYPSRRMVIVGSITGNSNTLAGNVP--PKA	230
Triticum_aestiv	188	HFLLARLLMEDLQKSDYPSRRMVIVGSITGNSNTLAGNVP--PKA	230
Arabidopsis_tha	202	HFLLARLLLLDDLKKSDYPSKRLLIIVGSITGNTNTLAGNVP--PKA	244
Hordeum_vulgare	196	HFLLARELLEDLKASDYPSKRLLIIVGSITGNTNTLAGNVP--PKA	238
Arabidopsis_tha	203	HFLLSRLLLDDLKKSDYPSKRMIIVGSITGNTNTLAGNVP--PKA	245
Avena_sativa	113	HFLLARLLLEDLQKSDYPSRRLVIVGSITGNDNTLAGNVP--PKA	155
Chlamydomonas_r	199	HFLLTNLLLLDDLKNAPNKQPRCIIIVGSITGNTNTLAGNVP--PKA	241
Daucus_carota	199	HFLLSRLLLDDLKNKSDYPSKRLLIIVGSITGNTNTLAGNVP--PKA	241
Marchantia_pale	259	HFLLARLLMEDLQKAKDSLKRMIIIVGSITGNSNTVAGNVP--PKA	301
Pisum_sativum	200	HFLLSRLLLLEDLKSDYPSKRLLIIVGSITGNTNTLAGNVP--PKA	242
Plectonema_bory	119	HFLLCNLLLEDLKHSTHNDPRLIILGTVTANSKELGGKIPIPAPA	163
Synechocystis_s	119	HFLLCNLLLEDLKACPDADKRLIILGTVTANSKELGGKIPIPAPP	163
cons	271	***** . *::*: . * ::*:*. * : :.::* . .	315

Arabidopsis_tha	249	NLGDLRGLAGGLNGLN--SSAMIDGGD-FVGAKAYKDSKVCNMLTMQ	292
Cucumis_sativus	243	NLGDLRGLAGGLNGLK--SSMIDGGE-FDGAKAYKDSKVCNMLTMQ	285
Hordeum_vulgare	231	SLGDLRGLAGGLSGASGSAMIDGDESFDGAKAYKDSKVCNMLTMQ	275
Triticum_aestiv	231	SLGDLRGLAGGLSGASGSAMIDGDESFDGAKAYKDSKVCNMLTMQ	275
Arabidopsis_tha	245	NLGDLRGLAGGLNGLN--SSAMIDGGD-FDGAKAYKDSKVCNMLTMQ	288
Hordeum_vulgare	239	NLGDLRGLAAGLNGVGSAAMIDGAE-FDGAKAYKDSKVCNMLTMQ	282
Arabidopsis_tha	246	NLGDLRGLASGLNGQN--SSMIDGGE-FDGAKAYKDSKVCNMLTMQ	288
Avena_sativa	156	NLGDLRGLAGGLTGASGSAMIDGDESFDGAKAYKDSKVCNMLTMQ	200
Chlamydomonas_r	242	NLGDLSGLAAGVPAA--NPMMDGQE-FNGAKAYKDSKVACMMTVR	283
Daucus_carota	242	NLGDLRGLAGGLNGMNSSAMIDGAE-FDGAKAYKDSKVCNMLTMQ	285
Marchantia_pale	302	NLGHLRGLAGGLNGVNSSSMIDGGE-FDGAKAYKDSKVCNMFTMQ	345
Pisum_sativum	243	NLGDLRGLAGGLTGLNSSAMIDGGD-FDGAKAYKDSKVCNMLTMQ	286
Plectonema_bory	164	DLGDLSGLEAGFKK--PIAMIDGKP-FKAGKAYKDSKLCNMITSR	205
Synechocystis_s	164	DLGNFEGFEAGFKK--PIAMINNKK-FKSGKAYKDSKLCNMLTTR	205
cons	316	.**.: *: .* . .::* * .*****: **:	360

Arabidopsis_tha	293	EFHRRFHEDTGITFASLYPGCIATTGLFREHIPLFRTLFPFPFQKY	337
Cucumis_sativus	286	EFHKRYHEETGITFASLYPGCIATTGLFREHIPLFRILFPFPFQKF	330
Hordeum_vulgare	276	EFHRRYHEETGITFSSLYPGCIATTGLFREHIPLFRTLFPFPFQKF	320
Triticum_aestiv	276	EFHRRYHEETGITFSSLYPGCIATTGLFREHIPLFRTLFPFPFQKF	320
Arabidopsis_tha	289	EFHRRFHHEETGVTFASLYPGCIASTGLFREHIPLFRALFPFPFQKY	333
Hordeum_vulgare	283	EFHRRYHEETGVTFASLYPGCIATTGLFREHIPLFRLLFPFPFQKY	327
Arabidopsis_tha	289	ELHRRYHEETGVTFASLYPGCIATTGLFREHIPLFRLLFPFPFQKY	333
Avena_sativa	201	EFHRRYHEDTGITFSSLYPGCIATTGLFREHIPLFRTLFPFPFQKF	245
Chlamydomonas_r	284	QMHQRFHDATGITFASLYPGCIAETGLFREHVPLFKTLFPFPFQKY	328
Daucus_carota	286	EFHRRYHEETGITFASLYPGCIATTGLFREHIPLFRTLFPFPFQKY	330
Marchantia_pale	346	EFHRRYHAETGITFSSLYPGCIAETGLFRNHVTLFRTLFPFPFQKY	390
Pisum_sativum	287	EFHRRYHEETGITFASLYPGCIATTGLFREHIPLFRTLFPFPFQKY	331
Plectonema_bory	206	ELHRRYHDSTGIVFNTLYPGCVADTPLFRNSLPVFQKVFPWFQKN	250
Synechocystis_s	206	ELHRRFHOETGIVFNSLYPGCVADTPLFRNHYSLFRTIFPWFQKN	250

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cons      361  ::*:~*  :*****:*  *  ****:  .:::  :~~  ***  405
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Arabidopsis_tha	338	ITKGYVSESEAGKRLAQVVADPSLTKSGVYWSWNKT	----	SASFE	378
Cucumis_sativus	331	ITQGYVSEDEAGKRLAQVVSEPSLTKSGVYWSWNKN	----	SASFE	371
Hordeum_vulgare	321	VTKGfVSEAESGKRLAQVVAEPVLTKSGVYWSWNKD	----	SASFE	361
Triticum_aestiv	321	VTKGfVSEAESGKRLAQVVAEPSLTKSGVYWSWNKD	----	SASFE	361
Arabidopsis_tha	334	ITKGYVSETESGKRLAQVVS DPSLTKSGVYWSWNNA	----	SASFE	374
Hordeum_vulgare	328	ITKGYVSEEEAGKRLAQVVSEPSLTKSGVYWSWNKN	----	SASFE	368
Arabidopsis_tha	334	ITKGYVSEEEAGKRLAQVVS DPSLGLKSGVYWSWNNN	----	SSSFE	374
Avena_sativa	246	VTKGfVSEAESGKRLAQVVGEP SLTKSGVYWSWNKD	----	SASFE	286
Chlamydomonas_r	329	ITKGYVSEEEAGRRLLAAVISDPKLNKSGAYWSWSST	----	TGSFD	369
Daucus_carota	331	ITKGYVSEAESGKRLAQVVSEPSLTKSGVYWSWNKD	----	SASFE	371
Marchantia_pale	391	ITKGYVSEEEAGKRMAQVVS DPKLKSGVYWSWNKD	----	SGSFE	431
Pisum_sativum	332	ITKGYVSEEEESGKRLAQVVS DPSLTKSGVYWSWNNA	----	SASFE	372
Plectonema_bory	251	ITGGYVSQELAGERTAQVVADPEFKQSGVHWSWGN	NRQKEG	RESFV	295
Synechocystis_s	251	VTKGYVSQELAGERVAMVVADDKFKDSGVHWSWGN	NRQQAG	REAFV	295

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cons      406  :*  * : ** :   : * . * * * : . :   :   . ** . : ** * . .   : *      450
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Arabidopsis_tha	379	NQLSQEASDVEKARRVWEVSEKLVGL	-A	405
Cucumis_sativus	372	NQLSQEASDAEKARKVWELSEKLVGL	-A	398
Hordeum_vulgare	362	NQLSQEASDPEKARKVWELSEKLVGL	-A	388
Triticum_aestiv	362	NQLSQEASDPEKARKVWELSEKLVGL	-A	388
Arabidopsis_tha	375	NQLSEEASDVEKARKVWEISEKLVGL	-A	401
Hordeum_vulgare	369	NQLSEEASDTEKARKVWELSEKLVGL	-A	395
Arabidopsis_tha	375	NQLSKEASDAEKAKKLWEVSEKLVGL	-A	401
Avena_sativa	287	NQLSQEASDPEKARKVWELSEKLVGL	-A	313
Chlamydomonas_r	370	NQVSEEVADDSKASKLWDISAKLVGL	SA	397
Daucus_carota	372	NQLSEEASDVEKARKVWEVSEKLVGL	-A	398
Marchantia_pale	432	NELSEEASNPEKAKRLWELSERLSGL	-V	458
Pisum_sativum	373	NQLSQEASDAEKARKVWEVSEKLVGL	-A	399
Plectonema_bory	296	QELSEKVTDDAKAKRMWELSEKLVGL	A-	322
Synechocystis_s	296	QELSEOGSDAQKAQRMWDLSEKLVGL	V-	322

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cons      451  :::*::  ::  **  ::*::*  :*  **  478
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