

Supplemental Table 1. Plant EST datasets provided by PlantGDB.

It includes 961 files representing plant species, subspecies, and cultivars. The PMID column represents identifiers of PubMed with direct links to the PubMed site for individual EST datasets. The “na” values (meaning “not available”) in the column represent that information on the papers is not provided by PubMed.

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Acacia_mangium.mRNA.EST.fasta	147	15574398	Tree Physiol. 25 (2), 167-178 (2005)
Acanthus_ebracteatus.mRNA.EST.fasta	521	na	Tree Genet. Genomes 2 (4), 196-201 (2006)
Acanthus_ebracteatus.mRNA.EST.fasta	107	na	Trees (Berl. West) 21, 515-520 (2007)
Acetabularia_acetabulum.mRNA.EST.fasta	941	15070428	BMC Plant Biol. 4 (1), 3 (2004)
Actaea_racemosa.mRNA.EST.fasta	2066	21188383	Plant Cell Rep. 30 (4), 613-629 (2010)
Actinidia_arguta.mRNA.EST.fasta	7257	18655731	BMC Genomics 9, 351 (2008)
Actinidia_chinensis.mRNA.EST.fasta	47379	18655731	BMC Genomics 9, 351 (2008)
Actinidia_deliciosa.mRNA.EST.fasta	57751	18655731	BMC Genomics 9, 351 (2008)
Actinidia_eriantha.mRNA.EST.fasta	12647	18655731	BMC Genomics 9, 351 (2008)
Actinidia_hemsleyana.mRNA.EST.fasta	5101	18655731	BMC Genomics 9, 351 (2008)
Actinidia_indochinensis.mRNA.EST.fasta	74	18655731	BMC Genomics 9, 351 (2008)
Actinidia_polygama.mRNA.EST.fasta	1348	18655731	BMC Genomics 9, 351 (2008)
Actinidia_setosa.mRNA.EST.fasta	1020	18655731	BMC Genomics 9, 351 (2008)
Adiantum_capillus_veneris.mRNA.EST.fasta	30540	15940394	J. Plant Res. 118 (3), 223-227 (2005)
Adonis_aestivalis.mRNA.EST.fasta	4189	18956021	Genome 51 (11), 888-896 (2008)
Aeluropus_lagopoides.mRNA.EST.fasta	10	na	Environ. Exp. Bot. 56 (3), 314-322 (2006)
Aeluropus_littoralis.mRNA.EST.fasta	492	17916418	Gene 404 (1-2), 61-69 (2007)
Ageratum_conyzoides.mRNA.EST.fasta	20	11535836	Proc. Natl. Acad. Sci. U.S.A. 98 (19), 10954-10959 (2001)
Agrostis_canina.mRNA.EST.fasta	4	17371611	Genet. Res. 88 (3), 165-175 (2006)
Agrostis_capillaris.mRNA.EST.fasta	7729	17497174	Mol. Genet. Genomics 278 (2), 197-209 (2007)
Agrostis_capillaris.mRNA.EST.fasta	4	17371611	Genet. Res. 88 (3), 165-175 (2006)
Agrostis_scabra.mRNA.EST.fasta	121	18950897	J. Plant Physiol. 166 (6), 588-601 (2009)
Agrostis_stolonifera.mRNA.EST.fasta	8472	17497174	Mol. Genet. Genomics 278 (2), 197-209 (2007)
Agrostis_stolonifera.mRNA.EST.fasta	7	17371611	Genet. Res. 88 (3), 165-175 (2006)
Allium_cepa.mRNA.EST.fasta	128	na	Theor. Appl. Genet. 103 (6-7), 979-991 (2001)
Allium_cepa.mRNA.EST.fasta	54	na	Theor. Appl. Genet. 96 (1), 52-62 (1998)
Allium_cepa.mRNA.EST.fasta	14	na	J. Food Biochem. 22 (2), 91-103 (1998)
Ananas_comosus.mRNA.EST.fasta	4102	17050644	J. Exp. Bot. 57 (14), 3921-3932 (2006)
Ananas_comosus.mRNA.EST.fasta	1547	16202174	BMC Plant Biol. 5 (1), 21 (2005)
Antirrhinum_majus.mRNA.EST.fasta	25310	15539471	Plant Cell 16 (12), 3197-3215 (2004)
Apium_graveolens.mRNA.EST.fasta	1218	15821978	Plant Mol. Biol. 57 (4), 517-540 (2005)
Apium_graveolens_Dulce_Group.mRNA.EST.fasta	996	12974812	Plant J. 36 (1), 67-81 (2003)
Arabidopsis_arenosa_x_Arabidopsis_thaliana.mRNA.EST.fasta	2122	17177805	Plant Biotechnol. J. 4 (4), 393-407 (2006)
Arabidopsis_halleri_subsp_halleri.mRNA.EST.fasta	307	16916885	J. Exp. Bot. 57 (12), 2967-2983 (2006)
Arabidopsis_lyrata.mRNA.EST.fasta	561	12618409	Genetics 163 (2), 723-733 (2003)
Arabidopsis_lyrata_subsp_petraea.mRNA.EST.fasta	11	na	Anim. Genet. 7 (4), 631-634 (2007)
Arabidopsis_thaliana.mRNA.EST.fasta	541852	17351049	Plant Physiol. 144 (1), 32-42 (2007)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Arabidopsis_thaliana.mRNA.EST.fasta	201750	16463100	Plant Mol. Biol. 60 (1), 69-85 (2006)
Arabidopsis_thaliana.mRNA.EST.fasta	111816	11910074	Science 296 (5565), 141-145 (2002)
Arabidopsis_thaliana.mRNA.EST.fasta	53236	10907847	DNA Res. 7 (3), 175-180 (2000)
Arabidopsis_thaliana.mRNA.EST.fasta	21937	na	Plant Sci. 124 (1), 68-78 (1997)
Arabidopsis_thaliana.mRNA.EST.fasta	11521	12799357	Genome Res. 13 (6), 1250-1257 (2003)
Arabidopsis_thaliana.mRNA.EST.fasta	11237	11115876	Plant Physiol. 124 (4), 1582-1594 (2000)
Arabidopsis_thaliana.mRNA.EST.fasta	9099	7846151	Plant Physiol. 106, 1241-1255 (1994)
Arabidopsis_thaliana.mRNA.EST.fasta	6164	na	Plant Cell Environ. 27 (5), 521-549 (2004)
Arabidopsis_thaliana.mRNA.EST.fasta	4504	16009969	Mol. Cell Proteomics 4 (10), 1558-1568 (2005)
Arabidopsis_thaliana.mRNA.EST.fasta	2256	21199568	Mol. Plant Pathol. 12 (2), 187-201 (2011)
Arabidopsis_thaliana.mRNA.EST.fasta	2105	na	Comp. Funct. Genomics 3 (2), 102-108 (2002)
Arabidopsis_thaliana.mRNA.EST.fasta	1587	15010618	Plant Mol. Biol. 53 (4), 545-563 (2003)
Arabidopsis_thaliana.mRNA.EST.fasta	722	11351099	Plant Physiol. 126 (1), 363-375 (2001)
Arabidopsis_thaliana.mRNA.EST.fasta	620	12620105	Genome Biol. 4 (3), R20 (2003)
Arabidopsis_thaliana.mRNA.EST.fasta	465	17956627	BMC Plant Biol. 7 (1), 56 (2007)
Arabidopsis_thaliana.mRNA.EST.fasta	239	16389298	PLoS Genet. 1 (6), e79 (2005)
Arabidopsis_thaliana.mRNA.EST.fasta	178	16504176	BMC Genomics 7, 38 (2006)
Arabidopsis_thaliana.mRNA.EST.fasta	113	10205900	Plant J. 17 (4), 433-444 (1999)
Arabidopsis_thaliana.mRNA.EST.fasta	84	14704363	Nucleic Acids Res. 32 (1), E6 (2004)
Arabidopsis_thaliana.mRNA.EST.fasta	44	10483124	Plant Cell Physiol. 40 (6), 624-639 (1999)
Arabidopsis_thaliana.mRNA.EST.fasta	35	9778851	Plant J. 15 (5), 707-720 (1998)
Arabidopsis_thaliana.mRNA.EST.fasta	33	11158529	Plant Cell 13 (1), 61-72 (2001)
Arabidopsis_thaliana.mRNA.EST.fasta	27	16455359	J. Plant Physiol. 163 (4), 452-462 (2006)
Arabidopsis_thaliana.mRNA.EST.fasta	27	10707356	Mol. Plant Microbe Interact. 13 (3), 309-315 (2000)
Arabidopsis_thaliana.mRNA.EST.fasta	10	15020432	Genetics 166 (1), 389-417 (2004)
Arabidopsis_thaliana.mRNA.EST.fasta	8	17339215	Genetics 176 (1), 151-160 (2007)
Arabidopsis_thaliana.mRNA.EST.fasta	5	10548732	Gene 239 (2), 309-316 (1999)
Arabidopsis_thaliana.mRNA.EST.fasta	3	7569999	Science 270, 467-470 (1995)
Arabidopsis_thaliana.mRNA.EST.fasta	2	10492172	DNA Res. 6, 247-253 (1999)
Arabidopsis_thaliana.mRNA.EST.fasta	2	9112769	Plant Physiol. 113 (4), 1081-1090 (1997)
Arabidopsis_thaliana.mRNA.EST.fasta	1	9358062	Gene 199 (1-2), 241-253 (1997)
Arachis_hypogaea.mRNA.EST.fasta	16931	19584933	Int J Plant Genomics 2009, doi:10.1155/2009/715605 (2009)
Arachis_hypogaea.mRNA.EST.fasta	575	19224247	Mol. Genet. Genomics 281 (6), 591-605 (2009)
Arachis_hypogaea.mRNA.EST.fasta	103	na	J. Plant Physiol. 168 (5), 481-492 (2011)
Arachis_hypogaea.mRNA.EST.fasta	1	12582692	Theor. Appl. Genet. 104 (2-3), 236-240 (2002)
Arachis_hypogaea_subsp_fastigiata.mRNA.EST.fasta	745	na	Plant Sci. 174 (1), 78-87 (2008)
Arachis_stenosperma.mRNA.EST.fasta	6264	17302987	BMC Plant Biol. 7 (1), 7 (2007)
Artemisia_annua.mRNA.EST.fasta	9322	16458889	FEBS Lett. 580 (5), 1411-1416 (2006)
Asarum_caudigerum.mRNA.EST.fasta	568	16675604	Ann. Bot. 98 (1), 157-163 (2006)
Athyrium_distentifolium.mRNA.EST.fasta	10	na	Mol. Ecol. Notes 3 (2), 287-290 (2003)
Aulacomnium_turgidum.mRNA.EST.fasta	439	na	Polar Biol. 33 (5), 617-626 (2010)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Avena_sativa.mRNA.EST.fasta	125	na	Theor. Appl. Genet. 97 (3), 356-369 (1998)
Avena_sativa.mRNA.EST.fasta	65	12669794	Genome 46 (1), 28-47 (2003)
Avena_sativa.mRNA.EST.fasta	30	7896104	Genetics 138 (4), 1251-1274 (1994)
Avena_sativa.mRNA.EST.fasta	19	na	Mol. Breed. 14 (4), 349-361 (2004)
Avena_sativa.mRNA.EST.fasta	9	10628998	Genetics 154, 397-412 (2000)
Avena_sativa.mRNA.EST.fasta	8	16137325	BMC Plant Biol. 5, 18 (2005)
Bambusa_balcooa.mRNA.EST.fasta	156	21272623	Gene j.gene.2011.01.004, doi:10.1016/j.gene.2011.01.004 (2011) In press
Beta_vulgaris.mRNA.EST.fasta	14697	12472698	Plant J. 32 (5), 845-857 (2002)
Beta_vulgaris.mRNA.EST.fasta	710	19121102	Physiol Plant 135 (1), 84-97 (2009)
Beta_vulgaris.mRNA.EST.fasta	53	na	Egypt J Genet Cytol 38, 187-206 (2009)
Beta_vulgaris.mRNA.EST.fasta	42	na	J. Gen. Plant Pathol. 73 (4), 260-265 (2007)
Beta_vulgaris.mRNA.EST.fasta	37	12669798	Genome 46 (1), 70-82 (2003)
Beta_vulgaris.mRNA.EST.fasta	35	na	Mol. Breed. 8 (1), 63-74 (2001)
Beta_vulgaris.mRNA.EST.fasta	14	15159641	Plant Mol. Biol. 54 (1), 147-156 (2004)
Beta_vulgaris.mRNA.EST.fasta	5	na	Mol. Plant Pathol. 7, 325-339 (2006)
Beta_vulgaris_subsp__vulgaris.mRNA.EST.fasta	2784	19046420	BMC Plant Biol. 8 (1), 122 (2008)
Betula_pendula.mRNA.EST.fasta	2532	14964534	Mol. Plant Microbe Interact. 17 (2), 202-215 (2004)
Betula_pendula.mRNA.EST.fasta	3	16640599	Plant J. 46 (4), 628-640 (2006)
Betula_pendula.mRNA.EST.fasta	1	18467468	Plant Physiol. 147 (3), 1199-1211 (2008)
Betula_platyphylla.mRNA.EST.fasta	2878	na	Plant Mol. Biol. Rep. 28 (3), 438-449 (2010)
Bienertia_sinuspersici.mRNA.EST.fasta	1921	na	Funct. Plant Biol. 37 (1), 1-13 (2010)
Bixa_orellana.mRNA.EST.fasta	954	na	Planta 163 (1), 141-145 (2002)
Boechera_cf__holboellii_SvA_2008.mRNA.EST.fasta	8	19220792	Plant J. 58 (5), 870-882 (2009)
Brachypodium_distachyon.mRNA.EST.fasta	20440	16791686	Theor. Appl. Genet. 113 (2), 186-195 (2006)
Brassica_junceae.mRNA.EST.fasta	4982	19821065	Theor. Appl. Genet. 120 (1), 71-83 (2009)
Brassica_junceae.mRNA.EST.fasta	90	na	Plant Sci. 167 (6), 1191-1201 (2004)
Brassica_junceae.mRNA.EST.fasta	4	18761493	Plant Biol (Stuttg) 10 (5), 539-547 (2008)
Brassica_napus.mRNA.EST.fasta	24716	na	Plant Sci. 168 (2), 439-447 (2005)
Brassica_napus.mRNA.EST.fasta	23521	19490642	BMC Genomics 10 (1), 256 (2009)
Brassica_napus.mRNA.EST.fasta	11618	19821065	Theor. Appl. Genet. 120 (1), 71-83 (2009)
Brassica_napus.mRNA.EST.fasta	8462	na	Mol Plant 2009, doi:10.1093/mp/ssp042 (2009)
Brassica_napus.mRNA.EST.fasta	1317	16024910	Plant Cell Physiol. 46 (9), 1525-1539 (2005)
Brassica_napus.mRNA.EST.fasta	1000	17384168	Plant Physiol. 144 (1), 134-154 (2007)
Brassica_napus.mRNA.EST.fasta	672	8029332	Plant Physiol. 103, 359-370 (1993)
Brassica_napus.mRNA.EST.fasta	405	na	Plant Cell Rep. 17 (12), 930-936 (1998)
Brassica_napus.mRNA.EST.fasta	280	na	(er) Mol. Plant Pathol. none, 0301FRISTENSKY (1999)
Brassica_napus.mRNA.EST.fasta	136	16977456	Planta 225 (3), 641-652 (2007)
Brassica_napus.mRNA.EST.fasta	49	19184461	Plant Mol. Biol. 70 (1-2), 31-45 (2009)
Brassica_napus.mRNA.EST.fasta	40	17401749	Plant Mol. Biol. 64 (4), 425-438 (2007)
Brassica_napus.mRNA.EST.fasta	20	18754099	Theor. Appl. Genet. 117 (7), 1031-1040 (2008)
Brassica_napus.mRNA.EST.fasta	12	16909288	Planta 225 (2), 515-522 (2007)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Brassica_napus.mRNA.EST.fasta	9	18761493	Plant Biol (Stuttg) 10 (5), 539-547 (2008)
Brassica_napus_var__napus.mRNA.EST.fasta	63	16020789	Genetics 171 (2), 765-781 (2005)
Brassica_nigra.mRNA.EST.fasta	1809	19821065	Theor. Appl. Genet. 120 (1), 71-83 (2009)
Brassica_nigra.mRNA.EST.fasta	1	18761493	Plant Biol (Stuttg) 10 (5), 539-547 (2008)
Brassica_oleracea.mRNA.EST.fasta	351	16545472	J. Microbiol. Methods 66 (3), 466-478 (2006)
Brassica_rapa.mRNA.EST.fasta	966	10417710	Plant J. 18 (6), 589-600 (1999)
Brassica_rapa_subsp__chinensis.mRNA.EST.fasta	26	na	Acta Physiol. Plant. 32 (4), 781-787 (2010)
Brassica_rapa_subsp__pekinensis.mRNA.EST.fasta	127144	18797175	Mol. Cells 26 (6), 595-605 (2008)
Brassica_rapa_subsp__pekinensis.mRNA.EST.fasta	2166	na	Plant Biotechnol. 23, 503-508 (2007)
Brassica_rapa_subsp__pekinensis.mRNA.EST.fasta	536	9177047	Planta 202 (1), 9-17 (1997)
Brassica_rapa_subsp__pekinensis.mRNA.EST.fasta	1	8787028	Plant Physiol. 111 (2), 577-588 (1996)
Bruguiera_cylindrica.mRNA.EST.fasta	126	na	Mol. Genet. Genomics 86 (2), 117-122 (2007)
Bruguiera_gymnorhiza.mRNA.EST.fasta	14842	na	Plant Sci. 171 (2), 234-241 (2006)
Bruguiera_gymnorhiza.mRNA.EST.fasta	5456	18568404	Plant Mol. Biol. 68 (1-2), 119-129 (2008)
Bupleurum_kaoi.mRNA.EST.fasta	41	na	Plant Sci. 173 (1), 12-24 (2007)
Cajanus_cajan.mRNA.EST.fasta	9888	20222972	BMC Plant Biol. 10 (1), 45 (2010)
Camellia_sinensis.mRNA.EST.fasta	350	na	Plant Sci. 166, 953-961 (2004)
Camellia_sinensis.mRNA.EST.fasta	129	18454353	Mol. Biol. Rep. 36 (5), 921-927 (2009)
Camellia_sinensis.mRNA.EST.fasta	51	17587678	Biosci. Biotechnol. Biochem. 71 (6), 1476-1486 (2007)
Camellia_sinensis.mRNA.EST.fasta	10	na	Plant Mol. Biol. Rep. 27 (3), 257-265 (2009)
Camellia_sinensis.mRNA.EST.fasta	1	16235114	Plant Mol. Biol. 59 (3), 515-532 (2005)
Cannabis_sativa.mRNA.EST.fasta	1575	na	Plant Sci. 174 (2), 205-220 (2008)
Cannabis_sativa.mRNA.EST.fasta	1565	19581347	J. Exp. Bot. 60 (13), 3715-3726 (2009)
Capsicum_annuum.mRNA.EST.fasta	25	10656596	Mol. Plant Microbe Interact. 13 (1), 136-142 (2000)
Capsicum_chinense.mRNA.EST.fasta	12	11355703	Mol. Cells 11 (2), 213-219 (2001)
Carica_papaya.mRNA.EST.fasta	75846	18432245	Nature 452 (7190), 991-996 (2008)
Carica_papaya.mRNA.EST.fasta	1216	na	Plant Sci. 170, 356-363 (2006)
Carica_papaya.mRNA.EST.fasta	30	na	Physiol. Mol. Plant Pathol. 65 (1), 21-30 (2004)
Carica_papaya.mRNA.EST.fasta	29	na	Sci. Hortic. 121 (2), 159-165 (2009)
Carica_papaya.mRNA.EST.fasta	23	na	Plant Sci. 174 (1), 38-50 (2008)
Carica_papaya.mRNA.EST.fasta	14	na	Postharvest Biol. Technol. 44 (2), 179-183 (2007)
Castanopsis_sieboldii.mRNA.EST.fasta	3354	na	Ann. For. Sci. 66, 509 (2009)
Casuarina_glauca.mRNA.EST.fasta	2028	16441749	New Phytol. 169 (4), 681-688 (2006)
Casuarina_glauca.mRNA.EST.fasta	1	na	Plant Physiol. 154 (3), 1372-1380 (2010)
Catharanthus_roseus.mRNA.EST.fasta	10282	16870181	FEBS Lett. 580 (18), 4501-4507 (2006)
Catharanthus_roseus.mRNA.EST.fasta	575	18326827	Plant Cell 20 (3), 524-542 (2008)
Catharanthus_roseus.mRNA.EST.fasta	105	16684234	New Phytol. 170 (4), 723-738 (2006)
Catharanthus_roseus.mRNA.EST.fasta	50	17050644	J. Exp. Bot. 57 (14), 3921-3932 (2006)
Cenchrus_ciliaris.mRNA.EST.fasta	8	9002607	Plant Mol. Biol. 32 (6), 1085-1092 (1996)
Centaurea_maculosa.mRNA.EST.fasta	4928	17524143	BMC Plant Biol. 7, 25 (2007)
Ceratonia_siliqua.mRNA.EST.fasta	1377	na	Tree Genet. Genomes 4 (4), 869-879 (2008)
Ceratopteris_richardii.mRNA.EST.fasta	5085	15965014	Plant Physiol. 138 (3), 1734-1745 (2005)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Chamaecyparis_obtusa.mRNA.EST.fasta	5830	16307365	Plant Mol. Biol. 59 (6), 895-907 (2005)
Chamaecyparis_obtusa.mRNA.EST.fasta	67	18595845	Tree Physiol. 28 (9), 1331-1340 (2008)
Chenopodium_album.mRNA.EST.fasta	697	21246286	Mol. Biol. Rep. DOI: 10.1007, 10.1007/s11033-011-0678-5 (2010) In press
Chenopodium_amaranticolor.mRNA.EST.fasta	13	11439122	Plant J. 26 (3), 339-349 (2001)
Chenopodium_quinoa.mRNA.EST.fasta	424	na	Plant Sci. 168 (2), 439-447 (2005)
Chlamydomonas_incerta.mRNA.EST.fasta	5132	16361241	Genetics 172 (3), 1567-1576 (2006)
Chlamydomonas_reinhardtii.mRNA.EST.fasta	26390	11089912	DNA Res. 7 (5), 305-307 (2000)
Chlamydomonas_reinhardtii.mRNA.EST.fasta	12842	na	Phycologia 43, 722-726 (2004)
Chlamydomonas_reinhardtii.mRNA.EST.fasta	11550	10691129	DNA Res. 6 (6), 369-373 (1999)
Chlamydomonas_reinhardtii.mRNA.EST.fasta	2032	19597503	Nat. Methods 6 (8), 589-592 (2009)
Chlamydomonas_sp__W80.mRNA.EST.fasta	82	na	World J. Microbiol. Biotechnol. 16, 23-29 (2000)
Chlamydomonas_sp__W80.mRNA.EST.fasta	11	17391362	FEMS Microbiol. Lett. 271 (1), 48-52 (2007)
Chlorella_vulgaris.mRNA.EST.fasta	263	na	Plant Sci. 175, 238-246 (2008)
Chloris_virgata.mRNA.EST.fasta	1893	20199868	Plant Physiol. Biochem. 48 (4), 247-255 (2010)
Chrysanthemum_x_morifolium.mRNA.EST.fasta	6924	na	Plant Mol. Biol. Rep. 27, 503-510 (2009)
Cicer_arietinum.mRNA.EST.fasta	20162	19912666	BMC Genomics 10, 523 (2009)
Cicer_arietinum.mRNA.EST.fasta	6273	19732460	BMC Genomics 10 (1), 415 (2009)
Cicer_arietinum.mRNA.EST.fasta	5097	18804093	Biochem. Biophys. Res. Commun. 376 (3), 578-583 (2008)
Cicer_arietinum.mRNA.EST.fasta	465	16107212	BMC Plant Biol. 5, 16 (2005)
Cicer_arietinum.mRNA.EST.fasta	108	15247380	Plant Physiol. 135 (3), 1608-1620 (2004)
Cicer_arietinum.mRNA.EST.fasta	62	19821728	Phytopathology 99 (11), 1245-1257 (2009)
Cicer_arietinum.mRNA.EST.fasta	58	na	Plant Sci. 167 (2), 207-216 (2004)
Cicer_arietinum.mRNA.EST.fasta	26	na	Physiol. Mol. Plant Pathol. 66 (5), 192-200 (2005)
Cicer_arietinum.mRNA.EST.fasta	26	na	Physiol. Mol. Plant Pathol. 68 (4-6), 176-188 (2006)
Cicer_arietinum.mRNA.EST.fasta	6	na	Plant Physiol. Biochem. 39, 363-366 (2001)
Cicer_arietinum.mRNA.EST.fasta	1	na	Plant Physiol. Biochem. 38, 1017-1026 (2001)
Cichorium_intybus.mRNA.EST.fasta	12226	19175922	BMC Plant Biol. 9 (1), 14 (2009)
Cichorium_intybus.mRNA.EST.fasta	3422	17553130	BMC Plant Biol. 7, 27 (2007)
Citrullus_colocynthis.mRNA.EST.fasta	18	19415285	Plant Cell Rep. 28 (6), 997-1009 (2009)
Citrullus_lanatus.mRNA.EST.fasta	7876	16802118	Plant Cell Rep. 25 (11), 1233-1245 (2006)
Citrullus_lanatus.mRNA.EST.fasta	15	na	Postharvest Biol. Technol. 48 (3), 372-377 (2008)
Citrus_aurantium.mRNA.EST.fasta	919	18598343	BMC Genomics 9, 318 (2008)
Citrus_aurantium.mRNA.EST.fasta	885	15830128	Plant Mol. Biol. 57 (3), 375-391 (2005)
Citrus_clementina.mRNA.EST.fasta	56019	18598343	BMC Genomics 9, 318 (2008)
Citrus_clementina.mRNA.EST.fasta	13942	15830128	Plant Mol. Biol. 57 (3), 375-391 (2005)
Citrus_clementina_x_Citrus_tangerina.mRNA.EST.fasta	1766	15830128	Plant Mol. Biol. 57 (3), 375-391 (2005)
Citrus_clementina_x_Citrus_tangerina.mRNA.EST.fasta	77	18598343	BMC Genomics 9, 318 (2008)
Citrus_jambhiri.mRNA.EST.fasta	24	14756316	Plant Mol. Biol. 53 (1-2), 189-199 (2003)
Citrus_macrophylla.mRNA.EST.fasta	1076	15830128	Plant Mol. Biol. 57 (3), 375-391 (2005)
Citrus_macrophylla.mRNA.EST.fasta	853	18598343	BMC Genomics 9, 318 (2008)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Citrus_medica.mRNA.EST.fasta	650	15830128	Plant Mol. Biol. 57 (3), 375-391 (2005)
Citrus_medica.mRNA.EST.fasta	465	18598343	BMC Genomics 9, 318 (2008)
Citrus_reshni.mRNA.EST.fasta	3686	18598343	BMC Genomics 9, 318 (2008)
Citrus_reshni.mRNA.EST.fasta	1443	15830128	Plant Mol. Biol. 57 (3), 375-391 (2005)
Citrus_sinensis.mRNA.EST.fasta	3774	na	Plant Sci. 165, 415-422 (2003)
Citrus_sinensis.mRNA.EST.fasta	2149	18598343	BMC Genomics 9, 318 (2008)
Citrus_sinensis.mRNA.EST.fasta	1820	15830128	Plant Mol. Biol. 57 (3), 375-391 (2005)
Citrus_sinensis.mRNA.EST.fasta	1348	20883438	J Integr Plant Biol 52 (10), 856-867 (2010)
Citrus_sinensis.mRNA.EST.fasta	267	19218315	J. Exp. Bot. 60 (3), 801-813 (2009)
Citrus_sinensis.mRNA.EST.fasta	168	21349317	Gene 476 (1-2), 1-9 (2011)
Citrus_sinensis.mRNA.EST.fasta	87	na	Engei Gakkai Zasshi 65, 487-495 (1996)
Citrus_sinensis.mRNA.EST.fasta	66	17525081	J. Exp. Bot. 58 (8), 2203-2216 (2007)
Citrus_sinensis.mRNA.EST.fasta	61	20807411	BMC Plant Biol. 10, 194 (2010)
Citrus_sinensis.mRNA.EST.fasta	3	na	J. Plant Biochem. Biotechnol. 14 (1), 9-14 (2005)
Citrus_sinensis_x_Citrus_trifoliata.mRNA.EST.fasta	1052	15830128	Plant Mol. Biol. 57 (3), 375-391 (2005)
Citrus_trifoliata.mRNA.EST.fasta	323	18930791	Gene 430 (1-2), 95-104 (2009)
Citrus_trifoliata.mRNA.EST.fasta	190	18708326	Tree Physiol. 28 (10), 1449-1457 (2008)
Citrus_trifoliata.mRNA.EST.fasta	13	na	Euphytica 164 (1), 209-219 (2008)
Citrus_unshiu.mRNA.EST.fasta	874	na	J. Am. Soc. Hortic. Sci. 122, 808-812 (1997)
Citrus_unshiu.mRNA.EST.fasta	620	na	Bull. Natl. Inst. Fruit Tree Sci. 2, 127-144 (2003)
Citrus_unshiu.mRNA.EST.fasta	393	na	Plant Sci. 165, 167-168 (2003)
Citrus_unshiu.mRNA.EST.fasta	8	16125877	Gene 359, 111-118 (2005)
Cleome_spinosa.mRNA.EST.fasta	4677	20333207	Genome Biol Evol 2009, 391-399 (2009)
Closterium_peracerosum_strigosum_littorale_complex.mRNA.EST.fasta	2046	16565296	Plant Physiol. 141 (1), 271-279 (2006)
Closterium_peracerosum_strigosum_littorale_complex.mRNA.EST.fasta	1190	14621293	DNA Res. 10 (4), 147-153 (2003)
Coffea_arabica.mRNA.EST.fasta	65163	20864545	Plant Physiol. 154 (3), 1053-1066 (2010)
Coffea_arabica.mRNA.EST.fasta	618	na	Mol. Plant Pathol. 5 (6), 527-536 (2004)
Coffea_arabica.mRNA.EST.fasta	139	17426774	Genome 49 (12), 1594-1605 (2006)
Coffea_arabica.mRNA.EST.fasta	4	15763663	Plant Physiol. Biochem. 43 (1), 27-35 (2005)
Coffea_canephora.mRNA.EST.fasta	46907	16273343	Theor. Appl. Genet. 112 (1), 114-130 (2005)
Coffea_canephora.mRNA.EST.fasta	12835	20864545	Plant Physiol. 154 (3), 1053-1066 (2010)
Coffea_canephora.mRNA.EST.fasta	8787	16924545	Mol. Genet. Genomics 276 (5), 436-449 (2006)
Coffea_eugenioides.mRNA.EST.fasta	18	20864545	Plant Physiol. 154 (3), 1053-1066 (2010)
Coffea_hybrid_cultivar.mRNA.EST.fasta	24	na	Plant Sci. 169 (2), 351-360 (2005)
Coleochaete_orbicularis.mRNA.EST.fasta	4464	20500869	BMC Plant Biol. 10 (1), 96 (2010)
Colocasia_esculenta.mRNA.EST.fasta	9	18622758	Mol. Biol. Rep. 36 (6), 1291-1297 (2009)
Coptis_japonica_var__dissecta.mRNA.EST.fasta	37	17132631	Plant Cell Physiol. 48 (1), 8-18 (2007)
Corchorus_olitorius.mRNA.EST.fasta	10	na	Plant Tissue Cult Biotechnol 16 (2), 95-104 (2006)
Corchorus_olitorius.mRNA.EST.fasta	1	na	Ann Biol Res 1 (3), 120-127 (2010)
Corchorus_trilocularis.mRNA.EST.fasta	2	na	Ann Biol Res 1 (3), 120-127 (2010)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Crocus_sativus.mRNA.EST.fasta	6202	17925031	BMC Plant Biol. 7 (1), 53 (2007)
Cryptomeria_japonica.mRNA.EST.fasta	36011	18691438	BMC Genomics 9 (1), 383 (2008)
Cryptomeria_japonica.mRNA.EST.fasta	8311	16307365	Plant Mol. Biol. 59 (6), 895-907 (2005)
Cryptomeria_japonica.mRNA.EST.fasta	4713	11052197	Plant Mol. Biol. 43 (4), 451-457 (2000)
Cryptomeria_japonica.mRNA.EST.fasta	3655	17169891	Tree Physiol. 26 (12), 1517-1528 (2006)
Cryptomeria_japonica.mRNA.EST.fasta	186	na	Theor. Appl. Genet. 94, 764-772 (1997)
Cryptomeria_japonica.mRNA.EST.fasta	1	12596006	Theor. Appl. Genet. 106 (4), 751-758 (2003)
Cucumis_melo.mRNA.EST.fasta	1800	17928373	J. Exp. Bot. 58 (13), 3645-3656 (2007)
Cucumis_melo.mRNA.EST.fasta	703	21338485	BMC Genomics 12, 122 (2011)
Cucumis_melo.mRNA.EST.fasta	123	na	Mol. Plant Pathol. 8 (1), 9-22 (2007)
Cucumis_melo.mRNA.EST.fasta	34	15060588	Genome 47 (2), 352-360 (2004)
Cucumis_melo.mRNA.EST.fasta	5	10712562	Plant Physiol. 122 (3), 977-983 (2000)
Cucumis_melo_subsp__agrestis.mRNA.EST.fasta	6921	17767721	BMC Genomics 8 (1), 306 (2007)
Cucumis_melo_subsp__melo.mRNA.EST.fasta	22683	17767721	BMC Genomics 8 (1), 306 (2007)
Cucumis_sativus.mRNA.EST.fasta	929	na	Plant Cell Environ. 26 (5), 757-770 (2003)
Cucumis_sativus.mRNA.EST.fasta	187	na	Physiol. Mol. Plant Pathol. 74 (2), 142-150 (2009)
Cucumis_sativus.mRNA.EST.fasta	105	12813576	Cell. Mol. Biol. Lett. 8 (2), 421-438 (2003)
Cucumis_sativus.mRNA.EST.fasta	64	na	Sci Hortic (Amsterdam) 117 (2), 89-94 (2008)
Cucumis_sativus.mRNA.EST.fasta	13	na	Plant Biotechnol. 18, 163-167 (2001)
Cunninghamia_lanceolata.mRNA.EST.fasta	407	18059541	Genome 50 (12), 1141-1155 (2007)
Cyamopsis_tetragonoloba.mRNA.EST.fasta	16476	18034910	BMC Plant Biol. 7 (1), 62 (2007)
Cyclamen_persicum.mRNA.EST.fasta	2083	na	J. Plant Growth Regul. 24, 102-115 (2005)
Cynodon_dactylon.mRNA.EST.fasta	15588	na	Funct. Plant Biol. 35 (7), 585-594 (2008)
Cynodon_dactylon.mRNA.EST.fasta	25	na	Plant Sci. 170 (4), 724-731 (2006)
Cynodon_transvaalensis.mRNA.EST.fasta	88	16395568	Theor. Appl. Genet. 112 (4), 727-737 (2006)
Daucus_carota.mRNA.EST.fasta	23	8938424	Plant Physiol. 112 (3), 1365-1374 (1996)
Daucus_carota.mRNA.EST.fasta	12	na	J. Plant Biol. 49 (2), 133-140 (2006)
Daucus_carota.mRNA.EST.fasta	4	na	Breed. Sci. 59, 37-46 (2009)
Daucus_carota.mRNA.EST.fasta	1	18726381	Sci. China, C, Life Sci. 43 (3), 265-271 (2000)
Deschampsia_antarctica.mRNA.EST.fasta	1961	18414009	Mol. Cells 25 (2), 258-264 (2008)
Deschampsia_antarctica.mRNA.EST.fasta	4	12955601	Extremophiles 7 (6), 459-469 (2003)
Dianthus_caryophyllus.mRNA.EST.fasta	235	20308205	J. Exp. Bot. 2010, doi:10.1093/jxb/erq064 (2010)
Dimocarpus_longan.mRNA.EST.fasta	65	na	Plant Sci. 170 (3), 500-510 (2006)
Dioscorea_alata.mRNA.EST.fasta	44134	21303556	BMC Genomics 12, 100 (2011)
Diospyros_kaki.mRNA.EST.fasta	9457	18301901	Plant Cell Rep. 27 (5), 931-938 (2008)
Dunaliella_salina.mRNA.EST.fasta	2831	na	Plant Sci. 179 (5), 437-449 (2010)
Dunaliella_salina.mRNA.EST.fasta	1038	16525865	Mar. Biotechnol. 8 (2), 120-128 (2006)
Dunaliella_salina.mRNA.EST.fasta	2	na	Plant Mol. Biol. Rep. 20 (1), 49-57 (2002)
Elaeis_guineensis.mRNA.EST.fasta	17599	18507865	BMC Plant Biol. 8 (1), 62 (2008)
Elaeis_guineensis.mRNA.EST.fasta	14537	17953740	BMC Genomics 8 (1), 381 (2007)
Elaeis_guineensis.mRNA.EST.fasta	2411	15862313	FEBS Lett. 579 (12), 2709-2714 (2005)
Elaeis_guineensis.mRNA.EST.fasta	1867	19199047	Plant Mol. Biol. 70 (1-2), 173-192 (2009)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Elaeis_guineensis.mRNA.EST.fasta	1605	na	Tree Genet. Genomes (2010) In press
Elaeis_guineensis.mRNA.EST.fasta	37	19706196	BMC Plant Biol. 9, 114 (2009)
Elaeis_guineensis.mRNA.EST.fasta	10	na	Biologia 63 (2), 227-235 (2008)
Eleusine_coracana_subsp_africana.mRNA.EST.fasta	304	17103137	Theor. Appl. Genet. 114 (2), 321-332 (2007)
Eleusine_coracana_subsp_coracana.mRNA.EST.fasta	299	17103137	Theor. Appl. Genet. 114 (2), 321-332 (2007)
Epipremnum_aureum.mRNA.EST.fasta	18	20167611	J. Exp. Bot. 61 (5), 1483-1493 (2010)
Eragrostis_curvula.mRNA.EST.fasta	12295	18196464	Plant Mol. Biol. 67 (1-2), 1-10 (2008)
Eragrostis_pilosa.mRNA.EST.fasta	787	16699556	Genome 49 (4), 365-372 (2006)
Eragrostis_tef.mRNA.EST.fasta	2816	16699556	Genome 49 (4), 365-372 (2006)
Eucalyptus_globulus.mRNA.EST.fasta	1173	19327132	BMC Plant Biol. 9, 36 (2009)
Eucalyptus_globulus.mRNA.EST.fasta	93	na	(in) Savidge,R.A., Barnett,J.R. and Napier,R. (Eds.); CELL AND MOLECULAR BIOLOGY OF WOOD FORMATION; BIOS Scientific, Oxford, UK (2000) In press
Eucalyptus_globulus_subsp_bicostata.mRNA.EST.fasta	54	na	New Phytol. 159 (1), 117-129 (2003)
Eucalyptus_globulus_subsp_bicostata.mRNA.EST.fasta	10	na	Tree Genet. Genomes 5, 317-327 (2009)
Eucalyptus_grandis.mRNA.EST.fasta	1574	15299141	Plant Physiol. 135 (4), 2368-2378 (2004)
Eucalyptus_gunnii.mRNA.EST.fasta	11303	na	Tree Genet. Genomes 5, 317-327 (2009)
Eucalyptus_gunnii.mRNA.EST.fasta	263	16684235	New Phytol. 170 (4), 739-752 (2006)
Eucalyptus_gunnii.mRNA.EST.fasta	224	15604680	Plant Mol. Biol. 55 (2), 263-280 (2004)
Eucalyptus_tereticornis.mRNA.EST.fasta	1131	15299141	Plant Physiol. 135 (4), 2368-2378 (2004)
Euphorbia_esula.mRNA.EST.fasta	524	16786310	Plant Mol. Biol. 61 (1-2), 329-344 (2006)
Euphorbia_tirucalli.mRNA.EST.fasta	9301	na	Plant Biotechnol. 21, 349-353 (2004)
Eutrema_halophilum.mRNA.EST.fasta	35171	19014467	BMC Plant Biol. 8 (1), 115 (2008)
Eutrema_halophilum.mRNA.EST.fasta	1739	15247369	Plant Physiol. 135 (3), 1718-1737 (2004)
Eutrema_halophilum.mRNA.EST.fasta	1038	19017120	J Integr Plant Biol 50 (10), 1327-1335 (2008)
Eutrema_salsugineum.mRNA.EST.fasta	6537	16021339	Plant Mol. Biol. 58 (4), 561-574 (2005)
Fagus_crenata.mRNA.EST.fasta	87	na	Conserv. Genet. 10, 1477-1485 (2009)
Fagus_sylvatica.mRNA.EST.fasta	1149	19290528	Plant Cell Rep. 28 (5), 873-882 (2009)
Fagus_sylvatica.mRNA.EST.fasta	493	11545437	Protist 152 (2), 139-147 (2001)
Fagus_sylvatica.mRNA.EST.fasta	300	16388470	Plant Biol (Stuttg) 7 (6), 670-676 (2005)
Festuca_arundinacea.mRNA.EST.fasta	2450	15710639	J. Exp. Bot. 56 (413), 897-907 (2005)
Festuca_arundinacea.mRNA.EST.fasta	12	na	Physiol. Mol. Plant Pathol. 63 (6), 305-317 (2003)
Festuca_mairei.mRNA.EST.fasta	124	17323082	Mol. Genet. Genomics 277 (5), 571-587 (2007)
Festuca_rubra_subsp_littoralis.mRNA.EST.fasta	3	19106017	J. Plant Physiol. 166 (7), 697-711 (2009)
Festuca_rubra_subsp_rubra.mRNA.EST.fasta	3	17566170	Gene 393 (1-2), 145-152 (2007)
Fragaria_chiloensis.mRNA.EST.fasta	137	20413181	J. Plant Physiol. 167 (14), 1179-1187 (2010)
Fragaria_vesca.mRNA.EST.fasta	2154	19785732	BMC Plant Biol. 9, 122 (2009)
Fragaria_x_ananassa.mRNA.EST.fasta	43	na	Fungal Genet. Biol. 128 (4), 633-650 (2006)
Fragaria_x_ananassa.mRNA.EST.fasta	30	10810141	Plant Cell 12 (5), 647-662 (2000)
Fragaria_x_ananassa.mRNA.EST.fasta	23	9684364	Planta 205 (4), 622-631 (1998)
Fragaria_x_ananassa.mRNA.EST.fasta	10	na	Mol. Ecol. Notes 6 (4), 1195-1197 (2006)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Fragaria_x_ananassa.mRNA.EST.fasta	5	na	Physiol Plant 118, 571-578 (2003)
Fragaria_x_ananassa.mRNA.EST.fasta	1	na	J. Exp. Bot. 62 (3), 1179-1188 (2011)
Galega_orientalis.mRNA.EST.fasta	4	11975733	Physiol Plant 114 (4), 588-593 (2002)
Gerbera_hybrid_cultivar.mRNA.EST.fasta	16994	15781570	Genome Res. 15 (4), 475-486 (2005)
Gerbera_hybrid_cultivar.mRNA.EST.fasta	3	14605235	Plant Physiol. 133 (4), 1831-1842 (2003)
Ginkgo_biloba.mRNA.EST.fasta	2039	na	Tree Genet. Genomes 6 (3), 357-365 (2010)
Glycine_max.mRNA.EST.fasta	68661	18927222	DNA Res. 15 (6), 333-346 (2008)
Glycine_max.mRNA.EST.fasta	29540	14624337	Theor. Appl. Genet. 108 (5), 903-913 (2004)
Glycine_max.mRNA.EST.fasta	3529	17601166	Mol. Plant Microbe Interact. 20 (7), 781-793 (2007)
Glycine_max.mRNA.EST.fasta	3433	15060591	Genome 47 (2), 380-388 (2004)
Glycine_max.mRNA.EST.fasta	1614	18414911	Funct. Integr. Genomics 8 (4), 341-359 (2008)
Glycine_max.mRNA.EST.fasta	801	16307369	Plant Mol. Biol. 59 (6), 965-979 (2005)
Glycine_max.mRNA.EST.fasta	224	10806241	Plant Physiol. 123 (1), 243-254 (2000)
Glycine_max.mRNA.EST.fasta	183	na	Plant Sci. 175 (4), 574-584 (2008)
Glycine_max.mRNA.EST.fasta	105	na	Mol. Plant Pathol. 7 (4), 235-247 (2006)
Glycine_max.mRNA.EST.fasta	88	7753826	Proc. Natl. Acad. Sci. U.S.A. 92, 4457-4461 (1995)
Glycine_max.mRNA.EST.fasta	56	18817859	Gene 426 (1-2), 15-22 (2008)
Glycine_max.mRNA.EST.fasta	18	na	Crop Sci. 43, 1548-1557 (2003)
Glycine_max.mRNA.EST.fasta	15	na	Mol. Plant Microbe Interact. 11, 1258-1263 (1998)
Glycine_max.mRNA.EST.fasta	14	na	Crop Sci. 41 (2), 516-521 (2001)
Glycine_max.mRNA.EST.fasta	14	14586656	Planta 218 (4), 606-614 (2004)
Glycine_max.mRNA.EST.fasta	13	na	Am. J. Bot. 89 (4), 559-565 (2002)
Glycine_soja.mRNA.EST.fasta	2003	16504061	BMC Plant Biol. 6, 4 (2006)
Glycyrrhiza_uralensis.mRNA.EST.fasta	50594	na	Plant Biotechnol. 26, 105-107 (2009)
Gossypium_arboreum.mRNA.EST.fasta	113	15020432	Genetics 166 (1), 389-417 (2004)
Gossypium_arboreum.mRNA.EST.fasta	6	na	Mol. Biol. (N.Y.) 42 (4), 492-498 (2008)
Gossypium_barbadense.mRNA.EST.fasta	10090	na	BMC Genomics (2010) In press
Gossypium_barbadense.mRNA.EST.fasta	889	17377794	Plant Cell Rep. 26 (8), 1309-1320 (2007)
Gossypium_barbadense.mRNA.EST.fasta	82	15020432	Genetics 166 (1), 389-417 (2004)
Gossypium_herbaceum_var__africanum.mRNA.EST.fasta	247	15020432	Genetics 166 (1), 389-417 (2004)
Gossypium_hirsutum.mRNA.EST.fasta	11692	16461577	Plant Cell 18 (3), 651-664 (2006)
Gossypium_hirsutum.mRNA.EST.fasta	9240	na	Physiol Plant 124 (3), 285-294 (2005)
Gossypium_hirsutum.mRNA.EST.fasta	1518	15020432	Genetics 166 (1), 389-417 (2004)
Gossypium_hirsutum.mRNA.EST.fasta	1380	15195948	Mol. Plant Microbe Interact. 17 (6), 654-667 (2004)
Gossypium_hirsutum.mRNA.EST.fasta	1002	na	Physiol. Mol. Plant Pathol. 67 (6), 319-335 (2005)
Gossypium_hirsutum.mRNA.EST.fasta	325	19328593	J. Plant Physiol. 166 (12), 1275-1283 (2009)
Gossypium_hirsutum.mRNA.EST.fasta	242	16429258	Plant Mol. Biol. 60 (2), 167-183 (2006)
Gossypium_hirsutum.mRNA.EST.fasta	210	18775953	J. Exp. Bot. 59 (13), 3661-3674 (2008)
Gossypium_hirsutum.mRNA.EST.fasta	166	12736302	Nucleic Acids Res. 31 (10), 2534-2543 (2003)
Gossypium_hirsutum.mRNA.EST.fasta	73	na	Mol. Biol. (N.Y.) 40 (5), 741-749 (2006)
Gossypium_hirsutum.mRNA.EST.fasta	65	na	Plant Sci. 173 (1), 61-69 (2007)
Gossypium_hirsutum.mRNA.EST.fasta	20	18080126	Plant Cell Rep. 27 (3), 553-561 (2008)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Gossypium_raidmondii.mRNA.EST.fasta	63467	16478941	Genome Res. 16 (3), 441-450 (2006)
Gossypium_raidmondii.mRNA.EST.fasta	110	15020432	Genetics 166 (1), 389-417 (2004)
Gypsophila_perfoliata.mRNA.EST.fasta	31	18504585	Plant Cell Rep. 27 (8), 1411-1422 (2008)
Helianthus_annuus.mRNA.EST.fasta	2448	15007731	Mol. Genet. Genomics 271 (3), 367-375 (2004)
Helianthus_annuus.mRNA.EST.fasta	1690	15821881	Plant Mol. Biol. 57 (2), 255-270 (2005)
Helianthus_annuus.mRNA.EST.fasta	1	na	Am. J. Bot. 91 (2), 184-191 (2004)
Helicosporidium_sp__ex_Simulium_jonesi.mRNA.EST.fasta	1196	16171185	Protist 156 (2), 181-190 (2005)
Hevea_brasiliensis.mRNA.EST.fasta	9861	17545224	J. Exp. Bot. 58 (10), 2429-2440 (2007)
Hevea_brasiliensis.mRNA.EST.fasta	238	20618931	BMC Plant Biol. 10 (1), 140 (2010)
Hevea_brasiliensis.mRNA.EST.fasta	18	na	(in) Jacob,J. (Ed.) Preprints of papers, International Natural Rubber Conference; India 2005, 257-262 (2005)
Hevea_brasiliensis.mRNA.EST.fasta	14	na	(in) Jacob,J., Krishnakumar,R. and Mathew,N.M. (Eds.); TAPPING PANEL DRYNESS OF RUBBER TREES; Kot-tayam, 35-46 (2006) In press
Hevea_brasiliensis.mRNA.EST.fasta	11	na	Tree Physiol. (2010) In press
Hevea_brasiliensis.mRNA.EST.fasta	9	18726169	Mol. Biotechnol. 41 (1), 42-52 (2009)
Hordeum_vulgare.mRNA.EST.fasta	22827	na	Mol. Plant Pathol. 6 (2), 139-151 (2005)
Hordeum_vulgare.mRNA.EST.fasta	7418	15447653	Plant J. 40 (2), 276-290 (2004)
Hordeum_vulgare.mRNA.EST.fasta	84	15361537	J. Exp. Bot. 55 (406), 2213-2218 (2004)
Hordeum_vulgare.mRNA.EST.fasta	25	17175551	J. Exp. Bot. 58 (3), 425-437 (2007)
Hordeum_vulgare.mRNA.EST.fasta	1	15153196	Physiol Plant 121 (2), 282-293 (2004)
Hordeum_vulgare.mRNA.EST.fasta	1	na	Mol. Breed. 14 (4), 349-361 (2004)
Hordeum_vulgare.mRNA.EST.fasta	1	17602190	Plant Cell Physiol. 48 (8), 1132-1147 (2007)
Hordeum_vulgare_subsp__spontaneum.mRNA.EST.fasta	11	17238046	Plant Mol. Biol. 64 (1-2), 17-34 (2007)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	95865	15447653	Plant J. 40 (2), 276-290 (2004)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	13109	12579433	Theor. Appl. Genet. 104 (1), 97-103 (2002)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	10856	na	Plant Mol. Biol. Rep. 18, 123-132 (2000)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	3917	15276436	Phytochemistry 65 (12), 1777-1784 (2004)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	144	12884135	J. Plant Res. 115 (1118), 119-130 (2002)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	48	15604661	Plant Mol. Biol. 55 (1), 1-15 (2004)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	45	19018994	Mol. Plant Pathol. 9 (5), 645-660 (2008)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	42	na	Theor. Appl. Genet. 97 (3), 356-369 (1998)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	23	10628998	Genetics 154, 397-412 (2000)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	6	na	Mol. Plant Pathol. 1, 277-286 (2000)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	1	15361537	J. Exp. Bot. 55 (406), 2213-2218 (2004)
Hordeum_vulgare_subsp__vulgare.mRNA.EST.fasta	1	na	Plant Sci. 133 (2), 191-201 (1998)
Humulus_lupulus.mRNA.EST.fasta	10887	18223037	Plant Cell 20 (1), 186-200 (2008)
Hyoscyamus_niger.mRNA.EST.fasta	2221	16720272	Chem. Biol. 13 (5), 513-520 (2006)
Ipomoea_batatas.mRNA.EST.fasta	2859	12586346	FEBS Lett. 536 (1-3), 101-105 (2003)
Ipomoea_batatas.mRNA.EST.fasta	1034	12692347	Plant Physiol. 131 (4), 1877-1893 (2003)
Ipomoea_batatas.mRNA.EST.fasta	33	12826628	Plant Cell Physiol. 44 (6), 630-636 (2003)
Ipomoea_nil.mRNA.EST.fasta	47	17221229	Plant Cell Rep. 26 (6), 823-835 (2007)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Iris_brevicaulis.mRNA.EST.fasta	2953	19515254	BMC Plant Biol. 9 (1), 72 (2009)
Iris_fulva.mRNA.EST.fasta	3591	19515254	BMC Plant Biol. 9 (1), 72 (2009)
Iris_x_hollandica.mRNA.EST.fasta	201	15082930	Plant Mol. Biol. 53 (6), 845-863 (2003)
Jatropha_curcas.mRNA.EST.fasta	13249	20691070	BMC Genomics 11 (1), 462 (2010)
Jatropha_curcas.mRNA.EST.fasta	6741	20979643	BMC Genomics 11 (1), 606 (2010)
Jatropha_curcas.mRNA.EST.fasta	931	na	Genomics Insights 3, 29-56 (2010)
Juglans_nigra.mRNA.EST.fasta	78	19787353	Plant Cell Rep. 28 (11), 1717-1724 (2009)
Kosteletzkya_virginica.mRNA.EST.fasta	38	19588254	Genetica 137 (2), 189-199 (2009)
Lactuca_sativa.mRNA.EST.fasta	586	17086420	Plant Cell Rep. 26 (4), 507-514 (2007)
Lactuca_sativa.mRNA.EST.fasta	18	18171591	J. Plant Physiol. 165 (17), 1757-1773 (2007)
Lavandula_angustifolia.mRNA.EST.fasta	14	na	Planta (2010) In press
Lavandula_angustifolia.mRNA.EST.fasta	3	20002329	Physiol Plant 2009, doi:10.1111/j.1399-3054.2009.01315.x (2009)
Leymus_chinensis.mRNA.EST.fasta	869	na	Plant Sci. 170 (6), 1081-1086 (2006)
Leymus_cinereus_x_Leymus_triticoides.mRNA.EST.fasta	28786	18543009	Funct. Integr. Genomics 8 (4), 375-386 (2008)
Lilium_longiflorum.mRNA.EST.fasta	886	16571618	Plant Cell Physiol. 47 (6), 698-705 (2006)
Lilium_longiflorum.mRNA.EST.fasta	103	16461387	Plant Physiol. 140 (4), 1374-1383 (2006)
Limonium_bicolor.mRNA.EST.fasta	2358	18640047	Plant Physiol. Biochem. 46 (11), 977-986 (2008)
Limonium_sinense.mRNA.EST.fasta	683	17364815	DNA Seq. 18 (1), 61-67 (2007)
Linum_usitatissimum.mRNA.EST.fasta	4638	15824284	Plant Physiol. 137 (4), 1211-1227 (2005)
Linum_usitatissimum.mRNA.EST.fasta	1462	17503083	Mol. Genet. Genomics 278 (2), 149-165 (2007)
Linum_usitatissimum.mRNA.EST.fasta	927	15666211	Plant Biol (Stuttg) 7 (1), 23-32 (2005)
Linum_usitatissimum.mRNA.EST.fasta	455	18593690	Ann. Bot. 102 (3), 317-330 (2008)
Lolium_multiflorum.mRNA.EST.fasta	116	18821059	Plant Mol. Biol. 69 (1-2), 1-21 (2009)
Lolium_multiflorum.mRNA.EST.fasta	34	na	Plant Breed. 125 (1), 43-51 (2006)
Lolium_perenne.mRNA.EST.fasta	1442	17626623	BMC Plant Biol. 7 (1), 36 (2007)
Lolium_perenne.mRNA.EST.fasta	145	na	Plant Growth Regul. 59 (3), 215-225 (2009)
Lolium_perenne.mRNA.EST.fasta	110	na	Dev. Comp. Immunol. 21 (4), 533-548 (2008)
Lolium_perenne.mRNA.EST.fasta	46	19326093	Theor. Appl. Genet. 118 (8), 1549-1560 (2009)
Lolium_perenne.mRNA.EST.fasta	25	20089196	BMC Mol. Biol. 11 (1), 8 (2010)
Lolium_perenne.mRNA.EST.fasta	22	17207264	Plant Biotechnol. J. 5 (1), 146-161 (2007)
Lolium_perenne.mRNA.EST.fasta	13	na	Euphytica 163 (1), 67-80 (2008)
Lolium_perenne.mRNA.EST.fasta	5	na	Mol. Ecol. Notes 5 (4), 951-957 (2005)
Lolium_perenne_subsp_perenne.mRNA.EST.fasta	10	19484189	Plant Mol. Biol. 70 (6), 709-723 (2009)
Lolium_temulentum.mRNA.EST.fasta	598	na	Insect Biochem. Mol. Biol. 173 (2), 73-83 (2007)
Lolium_temulentum.mRNA.EST.fasta	530	na	BMC Bioinformatics 171 (4), 459-469 (2006)
Lotus_japonicus.mRNA.EST.fasta	74289	15284495	Plant Mol. Biol. 54 (3), 405-414 (2004)
Lotus_japonicus.mRNA.EST.fasta	22751	10819328	DNA Res. 7 (2), 127-130 (2000)
Lotus_japonicus.mRNA.EST.fasta	7320	na	Plant Biotechnol. 25, 173-175 (2008)
Lotus_japonicus.mRNA.EST.fasta	898	10907852	DNA Res. 7, 213-216 (2000)
Lotus_japonicus.mRNA.EST.fasta	152	15980262	Plant Cell 17 (8), 2217-2229 (2005)
Lotus_japonicus.mRNA.EST.fasta	138	11943157	FEBS Lett. 514 (2-3), 229-237 (2002)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Lotus_japonicus.mRNA.EST.fasta	97	19088337	J. Exp. Bot. 60 (2), 641-650 (2009)
Lotus_japonicus.mRNA.EST.fasta	24	9276951	Plant Physiol. 114 (4), 1335-1346 (1997)
Lupinus_albus.mRNA.EST.fasta	5150	19123941	BMC Plant Biol. 9, 1 (2009)
Lupinus_albus.mRNA.EST.fasta	699	21342527	BMC Plant Biol. 11, 36 (2011)
Lupinus_albus.mRNA.EST.fasta	23	na	Plant Soil, DOI:10.1007/s11104-010-0711-9 (2011) In press
Lupinus_albus.mRNA.EST.fasta	7	12207659	Physiol Plant 116 (1), 28-36 (2002)
Lycoris_longituba.mRNA.EST.fasta	16	na	Mol Ecol Resour 9 (1), 278-280 (2009)
Macrotyloma_uniflorum.mRNA.EST.fasta	959	na	Plant Sci. 175 (3), 372-384 (2008)
Malus_x_domestica.mRNA.EST.fasta	558	na	Tree Genet. Genomes 5 (1), 81-91 (2009)
Malus_x_domestica.mRNA.EST.fasta	469	na	Tree Genet. Genomes 5 (1), 27-40 (2009)
Malus_x_domestica.mRNA.EST.fasta	102	na	Postharvest Biol. Technol. 56 (1), 1-11 (2010)
Malus_x_domestica.mRNA.EST.fasta	37	na	Mol. Breed. 22, 95-118 (2008)
Malus_x_domestica.mRNA.EST.fasta	5	na	Mol. Plant Pathol. 4, 447-457 (2003)
Malva_pusilla.mRNA.EST.fasta	813	na	Plant Sci. 167 (3), 481-489 (2004)
Mangifera_indica.mRNA.EST.fasta	37	16307821	J. Plant Physiol. 163 (6), 671-679 (2006)
Manihot_esculenta.mRNA.EST.fasta	35400	18096061	BMC Plant Biol. 7, 66 (2007)
Manihot_esculenta.mRNA.EST.fasta	18166	17541599	Plant Cell Rep. 26 (9), 1605-1618 (2007)
Manihot_esculenta.mRNA.EST.fasta	12204	15630618	Plant Mol. Biol. 56 (4), 541-554 (2004)
Manihot_esculenta.mRNA.EST.fasta	100	20536786	Physiol Plant 140 (2), 189-198 (2010)
Manihot_esculenta.mRNA.EST.fasta	59	16429258	Plant Mol. Biol. 60 (2), 167-183 (2006)
Manihot_esculenta.mRNA.EST.fasta	40	na	Mol. Plant Pathol. 6 (2), 113-123 (2005)
Manihot_esculenta.mRNA.EST.fasta	36	17318318	Plant Mol. Biol. 64 (1-2), 187-203 (2007)
Manihot_esculenta.mRNA.EST.fasta	7	10701114	Genome 43 (1), 62-67 (2000)
Marchantia_polymorpha.mRNA.EST.fasta	32277	17395720	Proc. Natl. Acad. Sci. U.S.A. 104 (15), 6472-6477 (2007)
Marchantia_polymorpha.mRNA.EST.fasta	970	10231024	DNA Res. 6 (1), 1-11 (1999)
Marchantia_polymorpha.mRNA.EST.fasta	445	10907846	DNA Res. 7, 165-174 (2000)
Marsilea_quadrifolia.mRNA.EST.fasta	14	11785932	Plant Mol. Biol. 47 (6), 703-715 (2001)
Medicago_sativa.mRNA.EST.fasta	5647	15578217	Planta 221 (1), 28-38 (2005)
Medicago_sativa_subsp__x_varia.mRNA.EST.fasta	38	17182124	Biochim. Biophys. Acta 1759 (11-12), 543-551 (2006)
Medicago_sativa_subsp__x_varia.mRNA.EST.fasta	18	10656590	Mol. Plant Microbe Interact. 13 (1), 96-106 (2000)
Medicago_sativa_subsp__x_varia.mRNA.EST.fasta	11	9574504	Mol. Plant Microbe Interact. 11 (5), 358-366 (1998)
Medicago_sativa_subsp__x_varia.mRNA.EST.fasta	2	9738887	Mol. Gen. Genet. 259 (1), 115-122 (1998)
Medicago_truncatula.mRNA.EST.fasta	3339	17378435	Mol. Plant Microbe Interact. 20 (3), 321-332 (2007)
Medicago_truncatula.mRNA.EST.fasta	3074	20209049	PLoS ONE 5 (3), e9519 (2010)
Medicago_truncatula.mRNA.EST.fasta	1775	12744459	Mol. Plant Microbe Interact. 16 (4), 306-314 (2003)
Medicago_truncatula.mRNA.EST.fasta	899	9701588	Plant Physiol. 117 (4), 1325-1332 (1998)
Medicago_truncatula.mRNA.EST.fasta	846	19021865	New Phytol. 181 (3), 563-575 (2009)
Medicago_truncatula.mRNA.EST.fasta	505	na	Physiol. Mol. Plant Pathol. 63 (1), 17-26 (2003)
Medicago_truncatula.mRNA.EST.fasta	450	10656586	Mol. Plant Microbe Interact. 13 (1), 62-71 (2000)
Medicago_truncatula.mRNA.EST.fasta	269	19268390	J. Plant Physiol. 166 (12), 1284-1295 (2009)
Medicago_truncatula.mRNA.EST.fasta	264	na	Physiol. Mol. Plant Pathol. 70 (1-3), 49-59 (2007)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Medicago_truncatula.mRNA.EST.fasta	175	18762986	Mol. Genet. Genomics 280 (5), 427-436 (2008)
Medicago_truncatula.mRNA.EST.fasta	107	17468219	Plant Physiol. 144 (3), 1455-1466 (2007)
Medicago_truncatula.mRNA.EST.fasta	58	16183836	Plant Physiol. 139 (2), 1065-1077 (2005)
Medicago_truncatula.mRNA.EST.fasta	29	15597744	Mol. Plant Microbe Interact. 17 (12), 1385-1393 (2004)
Medicago_truncatula.mRNA.EST.fasta	9	17488237	Plant J. 51 (1), 1-17 (2007)
Megathyrsus_maximus.mRNA.EST.fasta	196	19046615	J. Plant Physiol. 166 (7), 750-761 (2009)
Melaleuca_alternifolia.mRNA.EST.fasta	491	na	Plant Sci. 162 (1), 9-15 (2002)
Mentha_x_piperita.mRNA.EST.fasta	1316	10717007	Proc. Natl. Acad. Sci. U.S.A. 97 (6), 2934-2939 (2000)
Mesembryanthemum_crystallinum.mRNA.EST.fasta	10	10102366	Mol. Gen. Genet. 261 (2), 307-316 (1999)
Mesostigma_viride.mRNA.EST.fasta	10353	16476162	BMC Plant Biol. 6 (1), 2 (2006)
Mesostigma_viride.mRNA.EST.fasta	5619	16476689	Mol. Biol. Evol. 23 (5), 1011-1015 (2006)
Mirabilis_jalapa.mRNA.EST.fasta	8	16778017	Plant Physiol. 141 (4), 1604-1616 (2006)
Morus_alba.mRNA.EST.fasta	14	na	Nong Ye Sheng Wu Ji Shu Xue Bao 11 (4), 375-378 (2003)
Morus_indica.mRNA.EST.fasta	949	na	Tree Genet. Genomes 5 (2), 359-374 (2008)
Musa_acuminata_AAA_Group.mRNA.EST.fasta	48	na	Plant Sci. 172 (5), 1025-1036 (2007)
Musa_acuminata_AAA_Group.mRNA.EST.fasta	15	na	Sci Hortic (Amsterdam) 114 (2), 83-89 (2007)
Musa_acuminata_subsp__burmannicoides.mRNA.EST.fasta	2286	15841358	Theor. Appl. Genet. 110 (8), 1517-1522 (2005)
Musa_x_paradisiaca.mRNA.EST.fasta	1	18006520	J. Biochem. 142 (5), 561-570 (2007)
Nicotiana_attenuata.mRNA.EST.fasta	107	12692347	Plant Physiol. 131 (4), 1877-1893 (2003)
Nicotiana_attenuata.mRNA.EST.fasta	77	na	Physiol Plant 118 (2), 240-252 (2003)
Nicotiana_attenuata.mRNA.EST.fasta	73	12692348	Plant Physiol. 131 (4), 1894-1902 (2003)
Nicotiana_attenuata.mRNA.EST.fasta	54	11161026	Plant Physiol. 125 (2), 683-700 (2001)
Nicotiana_attenuata.mRNA.EST.fasta	26	20398280	BMC Plant Biol. 10 (1), 66 (2010)
Nicotiana_attenuata.mRNA.EST.fasta	18	15078438	Mol. Ecol. 13 (5), 981-995 (2004)
Nicotiana_benthamicana.mRNA.EST.fasta	6995	18931142	Plant Physiol. 148 (4), 1830-1846 (2008)
Nicotiana_benthamicana.mRNA.EST.fasta	20	17249421	Mol. Plant Microbe Interact. 20 (1), 41-52 (2007)
Nicotiana_benthamicana.mRNA.EST.fasta	7	20459311	Mol. Plant Microbe Interact. 23 (6), 715-726 (2010)
Nicotiana_megalosiphon.mRNA.EST.fasta	182	16610743	Mol. Plant Microbe Interact. 19 (4), 399-406 (2006)
Nicotiana_megalosiphon.mRNA.EST.fasta	84	na	Physiol. Mol. Plant Pathol. 67 (3-5), 231-236 (2006)
Nicotiana_megalosiphon.mRNA.EST.fasta	11	na	Plant Sci. 177 (3), 175-180 (2009)
Nicotiana_sp_.mRNA.EST.fasta	8	15078438	Mol. Ecol. 13 (5), 981-995 (2004)
Nicotiana_sp_.mRNA.EST.fasta	1	na	Physiol. Mol. Plant Pathol. 67 (6), 319-335 (2005)
Nicotiana_sylvestris.mRNA.EST.fasta	8580	na	Proc. Jpn. Acad. Ser. B 79, 151-154 (2003)
Nicotiana_tabacum.mRNA.EST.fasta	69021	18086234	Plant Biotechnol. J. 6 (3), 246-263 (2008)
Nicotiana_tabacum.mRNA.EST.fasta	13700	15509851	Plant Cell Physiol. 45 (9), 1280-1289 (2004)
Nicotiana_tabacum.mRNA.EST.fasta	11216	19052150	Plant Physiol. 149 (3), 1211-1230 (2009)
Nicotiana_tabacum.mRNA.EST.fasta	6469	na	Plant J. (2010) In press
Nicotiana_tabacum.mRNA.EST.fasta	797	20699003	BMC Plant Biol. 10, 167 (2010)
Nicotiana_tabacum.mRNA.EST.fasta	350	na	Plant Sci. 167 (4), 679-686 (2004)
Nicotiana_tabacum.mRNA.EST.fasta	206	na	Plant Sci. 170 (3), 417-425 (2006)
Nicotiana_tabacum.mRNA.EST.fasta	96	20004236	Gene 452 (2), 54-62 (2010)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Nicotiana_tabacum.mRNA.EST.fasta	89	17016740	Plant Mol. Biol. 63 (1), 137-149 (2007)
Nicotiana_tabacum.mRNA.EST.fasta	76	na	Plant Sci. 173, 291-301 (2007)
Nicotiana_tabacum.mRNA.EST.fasta	64	16649107	Plant Mol. Biol. 60 (5), 699-716 (2006)
Nicotiana_tabacum.mRNA.EST.fasta	53	12122207	Proc. Natl. Acad. Sci. U.S.A. 99 (16), 10870-10875 (2002)
Nicotiana_tabacum.mRNA.EST.fasta	49	15854833	Plant Physiol. Biochem. 43 (3), 249-259 (2005)
Nicotiana_tabacum.mRNA.EST.fasta	38	15837708	J. Exp. Bot. 56 (416), 1575-1590 (2005)
Nicotiana_tabacum.mRNA.EST.fasta	38	17547674	New Phytol. 175 (1), 140-154 (2007)
Nicotiana_tabacum.mRNA.EST.fasta	21	9210336	Plant Cell Physiol. 38 (5), 638-642 (1997)
Nicotiana_tabacum.mRNA.EST.fasta	15	9845441	Cytometry 33 (4), 460-468 (1998)
Nicotiana_tabacum.mRNA.EST.fasta	13	16456648	Plant Cell Rep. 25 (7), 728-740 (2006)
Nicotiana_tabacum.mRNA.EST.fasta	13	7787185	Plant Mol. Biol. 28, 27-38 (1995)
Nicotiana_tabacum.mRNA.EST.fasta	9	na	Plant Growth Regul. 32 (2-3), 307-313 (2000)
Nicotiana_tabacum.mRNA.EST.fasta	8	na	Plant Biotechnol. 24, 409-416 (2007)
Nicotiana_tabacum.mRNA.EST.fasta	8	9725022	Mol. Plant Microbe Interact. 11 (9), 895-905 (1998)
Nicotiana_tabacum.mRNA.EST.fasta	7	16643858	Biochem. Biophys. Res. Commun. 344 (3), 1001-1007 (2006)
Nicotiana_tabacum.mRNA.EST.fasta	7	11844105	Plant J. 29 (3), 269-279 (2002)
Nicotiana_tabacum.mRNA.EST.fasta	4	9617809	Plant Mol. Biol. 37 (2), 385-389 (1998)
Noccaea_caerulescens.mRNA.EST.fasta	4289	16684236	New Phytol. 170 (4), 753-766 (2006)
Ocimum_basilicum.mRNA.EST.fasta	5690	11161012	Plant Physiol. 125 (2), 539-555 (2001)
Ocimum_basilicum.mRNA.EST.fasta	5422	14657409	Plant Physiol. 134 (1), 370-379 (2004)
Ocimum_basilicum.mRNA.EST.fasta	4806	15516500	Plant Physiol. 136 (3), 3724-3736 (2004)
Ocimum_basilicum.mRNA.EST.fasta	28	17263464	J. Agric. Food Chem. 55 (3), 706-713 (2007)
Oenothera_elata_subsp__hookeri.mRNA.EST.fasta	3530	16829020	Genomics 88 (3), 372-380 (2006)
Olea_europaea.mRNA.EST.fasta	1102	19852839	BMC Plant Biol. 9, 128 (2009)
Oryza_longistaminata.mRNA.EST.fasta	71367	21159186	BMC Genomics 11, 705 (2010)
Oryza_minuta.mRNA.EST.fasta	5199	14968263	Plant Cell Rep. 22 (11), 839-847 (2004)
Oryza_minuta.mRNA.EST.fasta	109	14648106	Plant Cell Rep. 22 (8), 599-607 (2004)
Oryza_sativa.mRNA.EST.fasta	27959	17187943	Gene 389 (2), 107-113 (2007)
Oryza_sativa.mRNA.EST.fasta	8190	15685292	PLoS Biol. 3 (2), e38 (2005)
Oryza_sativa.mRNA.EST.fasta	463	18922162	BMC Genomics 9 (1), 485 (2008)
Oryza_sativa.mRNA.EST.fasta	107	14527726	Gene 314, 141-148 (2003)
Oryza_sativa.mRNA.EST.fasta	49	15558307	Planta 221 (1), 105-112 (2004)
Oryza_sativa.mRNA.EST.fasta	39	12384290	Gene 297 (1-2), 93-102 (2002)
Oryza_sativa.mRNA.EST.fasta	10	12595990	Theor. Appl. Genet. 106 (4), 620-628 (2003)
Oryza_sativa.mRNA.EST.fasta	8	17585380	Theor. Appl. Genet. 115 (4), 537-547 (2007)
Oryza_sativa.mRNA.EST.fasta	8	17219102	Plant Cell Rep. 26 (7), 1097-1110 (2007)
Oryza_sativa.mRNA.EST.fasta	2	na	Acta Bot. Sin. 42 (1), 55-58 (2000)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	77946	15685292	PLoS Biol. 3 (2), e38 (2005)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	27566	19000321	BMC Plant Biol. 8, 114 (2008)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	19684	15918889	Plant J. 42 (5), 772-780 (2005)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	18424	15888683	Plant Physiol. 138 (1), 105-115 (2005)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Oryza_sativa_Indica_Group.mRNA.EST.fasta	5815	17132712	J. Exp. Bot. 58 (2), 253-265 (2007)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	988	14564505	Mol. Genet. Genomics 270 (3), 253-262 (2003)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	283	7896104	Genetics 138 (4), 1251-1274 (1994)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	132	na	Theor. Appl. Genet. 97 (3), 356-369 (1998)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	114	18315879	BMC Plant Biol. 8, 26 (2008)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	111	na	Sci. China, C, Life Sci. 45 (5), 449-467 (2002)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	65	15791452	Theor. Appl. Genet. 110 (7), 1177-1186 (2005)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	47	14605807	Theor. Appl. Genet. 108 (3), 525-534 (2004)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	7	12595990	Theor. Appl. Genet. 106 (4), 620-628 (2003)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	1	12897975	Sheng Wu Hua Xue Yu Sheng Wu Wu Li Xue Bao 35 (8), 768-773 (2003)
Oryza_sativa_Indica_Group.mRNA.EST.fasta	1	17522955	Plant Mol. Biol. 65 (4), 403-415 (2007)
Oryza_sativa_Japonica_Group.mRNA.EST.fasta	50513	15888683	Plant Physiol. 138 (1), 105-115 (2005)
Oryza_sativa_Japonica_Group.mRNA.EST.fasta	2080	7987417	Plant J. 6 (4), 615-624 (1994)
Oryza_sativa_Japonica_Group.mRNA.EST.fasta	785	14564505	Mol. Genet. Genomics 270 (3), 253-262 (2003)
Oryza_sativa_Japonica_Group.mRNA.EST.fasta	754	15659629	Plant Cell 17 (2), 389-403 (2005)
Oryza_sativa_Japonica_Group.mRNA.EST.fasta	396	15514441	Genes Genet. Syst. 79 (4), 213-226 (2004)
Oryza_sativa_Japonica_Group.mRNA.EST.fasta	222	18315879	BMC Plant Biol. 8, 26 (2008)
Oryza_sativa_Japonica_Group.mRNA.EST.fasta	105	16987952	Mol. Biol. Evol. 23 (12), 2505-2520 (2006)
Oryza_sativa_Japonica_Group.mRNA.EST.fasta	56	11332734	Mol. Plant Microbe Interact. 14 (5), 685-692 (2001)
Oryza_sativa_Japonica_Group.mRNA.EST.fasta	37	8541495	Plant Mol. Biol. 29 (4), 685-689 (1995)
Oryza_sativa_Japonica_Group.mRNA.EST.fasta	2	na	Plant Biotechnol. 15, 77-81 (1998)
Oryza_sativa_Japonica_Group.mRNA.EST.fasta	2	na	J. Gen. Plant Pathol. 68, 128-133 (2002)
Paeonia_suffruticosa.mRNA.EST.fasta	2204	19292824	Physiol Plant 135 (4), 439-449 (2009)
Panax_quinquefolius.mRNA.EST.fasta	4719	19947964	Physiol Plant 138 (2), 134-149 (2010)
Panax_quinquefolius.mRNA.EST.fasta	269	na	Physiol. Mol. Plant Pathol. 72 (1-3), 10-20 (2008)
Pandanus_odoratissimus.mRNA.EST.fasta	977	20149480	J. Plant Physiol. 167 (11), 914-919 (2010)
Panicum_virgatum.mRNA.EST.fasta	5734	na	Bioenergy Res 2010, doi10.1007/s12155-010-9080-8 (2010)
Papaver_somniferum.mRNA.EST.fasta	4612	17077972	Planta 225 (5), 1085-1106 (2007)
Papaver_somniferum.mRNA.EST.fasta	10	15386001	Nature 431 (7007), 413-414 (2004)
Paullinia_cupana_var_sorbilis.mRNA.EST.fasta	14888	17917729	Plant Cell Rep. 27 (1), 117-124 (2008)
Pennisetum_glaucum.mRNA.EST.fasta	2493	17558562	Plant Mol. Biol. 64 (6), 713-732 (2007)
Pennisetum_glaucum.mRNA.EST.fasta	66	19236576	Mol. Plant Pathol. 10 (2), 291-304 (2009)
Persicaria_minor.mRNA.EST.fasta	72	na	Acta Physiol. Plant., DOI10.1007/s11738-010-0546-2 (2010) In press
Petunia_axillaris_subsp_axillaris.mRNA.EST.fasta	9379	21143680	Plant J. 64 (6), 1002-1017 (2010)
Petunia_axillaris_subsp_axillaris.mRNA.EST.fasta	1696	na	Plant Sci. 173, 495-500 (2007)
Petunia_x_hybrida.mRNA.EST.fasta	36404	21143680	Plant J. 64 (6), 1002-1017 (2010)
Petunia_x_hybrida.mRNA.EST.fasta	3347	na	Plant Sci. 173, 495-500 (2007)
Petunia_x_hybrida.mRNA.EST.fasta	284	na	Sci. Hortic. 127 (3), 398-404 (2011)
Phalaenopsis_equestris.mRNA.EST.fasta	5604	na	Plant Sci. 170 (3), 426-432 (2006)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Phalaenopsis_violacea.mRNA.EST.fasta	2359	16836766	BMC Plant Biol. 6 (1), 14 (2006)
Phaseolus_vulgaris.mRNA.EST.fasta	37919	19397807	BMC Plant Biol. 9 (1), 46 (2009)
Phaseolus_vulgaris.mRNA.EST.fasta	15333	15824284	Plant Physiol. 137 (4), 1211-1227 (2005)
Phaseolus_vulgaris.mRNA.EST.fasta	5243	16121253	Genome 48 (3), 562-570 (2005)
Phaseolus_vulgaris.mRNA.EST.fasta	247	19646251	BMC Plant Biol. 9 (1), 100 (2009)
Phaseolus_vulgaris.mRNA.EST.fasta	147	17449651	Plant Physiol. 144 (2), 752-767 (2007)
Phaseolus_vulgaris.mRNA.EST.fasta	136	21416283	Plant Cell Rep. 10.1007, DOI:10.1007/s00299-011-1055-5 (2011) In press
Phaseolus_vulgaris.mRNA.EST.fasta	99	17701202	Planta 227 (1), 151-165 (2007)
Phaseolus_vulgaris.mRNA.EST.fasta	78	20300860	Mol. Biol. Rep. 38 (1), 75-81 (2010)
Phaseolus_vulgaris.mRNA.EST.fasta	41	18321191	Mol. Plant Microbe Interact. 21 (4), 459-468 (2008)
Phaseolus_vulgaris.mRNA.EST.fasta	32	na	Mol. Breed. 21 (2), 159-172 (2008)
Phaseolus_vulgaris.mRNA.EST.fasta	20	16568285	Theor. Appl. Genet. 112 (8), 1452-1457 (2006)
Phoenix_dactylifera.mRNA.EST.fasta	74	na	Plant Sci. 179 (4), 325-332 (2010)
Physcomitrella_patens.mRNA.EST.fasta	510	18047796	J. Biochem. Mol. Biol. 40 (6), 986-1001 (2007)
Physcomitrella_patens.mRNA.EST.fasta	172	10394633	Plant Cell Physiol. 40 (4), 378-387 (1999)
Physcomitrella_patens.mRNA.EST.fasta	81	na	Bot. Acta 111, 145-151 (1998)
Physcomitrella_patens.mRNA.EST.fasta	45	15322883	Planta 220 (3), 384-394 (2005)
Physcomitrella_patens.mRNA.EST.fasta	29	17601824	Plant Cell 19 (6), 1750-1769 (2007)
Physcomitrella_patens.mRNA.EST.fasta	7	na	Plant Biol. 4, 595-602 (2002)
Physcomitrella_patens.mRNA.EST.fasta	7	14586556	Curr. Genet. 44 (6), 339-347 (2004)
Physcomitrella_patens.mRNA.EST.fasta	1	15310072	J. Plant Physiol. 161 (7), 823-835 (2004)
Physcomitrella_patens_subsp_patens.mRNA.EST.fasta	82313	12808149	Proc. Natl. Acad. Sci. U.S.A. 100 (13), 8007-8012 (2003)
Picea_abies.mRNA.EST.fasta	7892	17764001	Plant Mol. Biol. 65 (3), 311-328 (2007)
Picea_abies.mRNA.EST.fasta	4	na	Tree Genet. Genomes 2 (1), 39-52 (2006)
Picea_abies.mRNA.EST.fasta	1	na	Theor. Appl. Genet. 103 (8), 1223-1231 (2001)
Picea_engelmannii_x_Picea_glauca.mRNA.EST.fasta	16043	16545489	J. Plant Physiol. 164 (1), 78-89 (2007)
Picea_engelmannii_x_Picea_glauca.mRNA.EST.fasta	12127	18854048	BMC Genomics 9, 484 (2008)
Picea_glauca.mRNA.EST.fasta	49486	16545489	J. Plant Physiol. 164 (1), 78-89 (2007)
Picea_glauca.mRNA.EST.fasta	25934	18854048	BMC Genomics 9, 484 (2008)
Picea_sitchensis.mRNA.EST.fasta	68724	16545489	J. Plant Physiol. 164 (1), 78-89 (2007)
Picea_sitchensis.mRNA.EST.fasta	12061	18854048	BMC Genomics 9, 484 (2008)
Pinus_densiflora.mRNA.EST.fasta	2779	19203959	Tree Physiol. 29 (3), 411-421 (2009)
Pinus_elliottii.mRNA.EST.fasta	142	15221139	Theor. Appl. Genet. 109 (5), 922-932 (2004)
Pinus_elliottii.mRNA.EST.fasta	8	14711673	Appl. Environ. Microbiol. 70 (1), 441-451 (2004)
Pinus_pinaster.mRNA.EST.fasta	196	20140468	Amino Acids 2010, doi:10.1007/s00726-010-0483-9 (2010) In press
Pinus_pinaster.mRNA.EST.fasta	9	12602883	Plant Mol. Biol. 51 (2), 249-262 (2003)
Pinus_radiata.mRNA.EST.fasta	6389	19159482	BMC Genomics 10 (1), 41 (2009)
Pinus_radiata.mRNA.EST.fasta	217	20561208	New Phytol. 187 (3), 764-776 (2010)
Pinus_radiata.mRNA.EST.fasta	82	18406157	Plant Physiol. Biochem. 46 (5-6), 559-568 (2008)
Pinus_taeda.mRNA.EST.fasta	55076	17001497	Plant Mol. Biol. 62 (4-5), 485-501 (2006)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Pinus_taeda.mRNA.EST.fasta	28579	na	Plant Sci. 170 (3), 417-425 (2006)
Pinus_taeda.mRNA.EST.fasta	1097	9689143	Proc. Natl. Acad. Sci. U.S.A. 95 (16), 9693-9698 (1998)
Pinus_taeda.mRNA.EST.fasta	44	15221139	Theor. Appl. Genet. 109 (5), 922-932 (2004)
Pinus_taeda.mRNA.EST.fasta	35	15821878	Plant Mol. Biol. 57 (2), 203-224 (2005)
Pinus_taeda.mRNA.EST.fasta	3	na	(in) Dean,J.F.D. (ed.) Proceedings of the 26th Biennial Southern Forest Tree Improvement Conference. University of Georgia, Athens, GA 26, 198-201 (2001)
Piper_longum.mRNA.EST.fasta	11	na	J Theor Appl Inf Technol 4 (5), 459-465 (2008)
Pisum_sativum.mRNA.EST.fasta	10346	19706781	J. Exp. Bot. 60 (14), 4201-4213 (2009)
Pisum_sativum.mRNA.EST.fasta	2735	18590528	BMC Plant Biol. 8 (1), 73 (2008)
Pisum_sativum.mRNA.EST.fasta	53	na	Theor. Appl. Genet. 95 (8), 1289-1299 (1997)
Pisum_sativum.mRNA.EST.fasta	4	9341677	Mol. Gen. Genet. 256 (1), 37-44 (1997)
Pisum_sativum.mRNA.EST.fasta	1	10517839	Plant Physiol. 121 (2), 479-488 (1999)
Pisum_sativum.mRNA.EST.fasta	1	na	Plant Mol. Biol. 4, 81-86 (1985)
Pisum_sativum.mRNA.EST.fasta	1	9845441	Cytometry 33 (4), 460-468 (1998)
Plantago_major.mRNA.EST.fasta	3	17041024	Plant Physiol. 142 (4), 1427-1441 (2006)
Plumbago_zeylanica.mRNA.EST.fasta	1522	19500307	Plant J. 60 (1), 33-47 (2009)
Plumbago_zeylanica.mRNA.EST.fasta	258	16307821	J. Plant Physiol. 163 (6), 671-679 (2006)
Poa_secunda.mRNA.EST.fasta	30	na	J. Plant Physiol. 159, 661-670 (2002)
Populus_alba.mRNA.EST.fasta	162	19464753	J. Plant Physiol. 166 (14), 1544-1556 (2009)
Populus_alba_x_Populus_tremula_var__glandulosa.mRNA.EST.fasta	7595	na	Plant Sci. 169 (6), 1118-1124 (2005)
Populus_alba_x_Populus_tremula_var__glandulosa.mRNA.EST.fasta	720	19962907	Plant Physiol. Biochem. 48 (2-3), 136-141 (2010)
Populus_euphratica.mRNA.EST.fasta	13845	16356264	Genome Biol. 6 (12), R101 (2005)
Populus_euphratica.mRNA.EST.fasta	58	14704136	Tree Physiol. 24 (3), 265-276 (2004)
Populus_nigra.mRNA.EST.fasta	36104	18053163	BMC Genomics 8, 448 (2007)
Populus_nigra.mRNA.EST.fasta	15257	15653793	Plant Cell Physiol. 45 (12), 1738-1748 (2004)
Populus_tomentosa.mRNA.EST.fasta	209	19426563	BMC Genomics 10 (1), 215 (2009)
Populus_tremula.mRNA.EST.fasta	17156	15353603	Proc. Natl. Acad. Sci. U.S.A. 101 (38), 13951-13956 (2004)
Populus_tremula_x_Populus_alba.mRNA.EST.fasta	10062	15095135	Plant Biol (Stuttg) 6 (1), 55-64 (2004)
Populus_tremula_x_Populus_alba.mRNA.EST.fasta	279	15908438	Plant Cell Physiol. 46 (8), 1213-1225 (2005)
Populus_tremula_x_Populus_alba.mRNA.EST.fasta	77	12729894	FEBS Lett. 542 (1-3), 37-41 (2003)
Populus_tremula_x_Populus_alba.mRNA.EST.fasta	2	19943172	Planta 231 (3), 525-535 (2010)
Populus_tremula_x_Populus_tremuloides.mRNA.EST.fasta	26226	15353603	Proc. Natl. Acad. Sci. U.S.A. 101 (38), 13951-13956 (2004)
Populus_tremula_x_Populus_tremuloides.mRNA.EST.fasta	4809	9789088	Proc. Natl. Acad. Sci. U.S.A. 95 (22), 13330-13335 (1998)
Populus_tremula_x_Populus_tremuloides.mRNA.EST.fasta	28	11724959	Proc. Natl. Acad. Sci. U.S.A. 98 (25), 14732-14737 (2001)
Populus_trichocarpa.mRNA.EST.fasta	31797	19121102	Physiol Plant 135 (1), 84-97 (2009)
Populus_trichocarpa.mRNA.EST.fasta	27835	16626454	Mol. Ecol. 15 (5), 1275-1297 (2006)
Populus_trichocarpa.mRNA.EST.fasta	6594	15353603	Proc. Natl. Acad. Sci. U.S.A. 101 (38), 13951-13956 (2004)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Populus_trichocarpa.mRNA.EST.fasta	883	9789088	Proc. Natl. Acad. Sci. U.S.A. 95 (22), 13330-13335 (1998)
Populus_trichocarpa_x_Populus_deltoides.mRNA.EST.fasta	19982	19121102	Physiol Plant 135 (1), 84-97 (2009)
Populus_trichocarpa_x_Populus_deltoides.mRNA.EST.fasta	16625	16626454	Mol. Ecol. 15 (5), 1275-1297 (2006)
Populus_trichocarpa_x_Populus_deltoides.mRNA.EST.fasta	6335	12729894	FEBS Lett. 542 (1-3), 37-41 (2003)
Populus_trichocarpa_x_Populus_deltoides.mRNA.EST.fasta	945	15605173	Planta 219 (6), 936-947 (2004)
Populus_trichocarpa_x_Populus_deltoides.mRNA.EST.fasta	103	na	Plant Cell Environ. 26 (5), 757-770 (2003)
Populus_trichocarpa_x_Populus_nigra.mRNA.EST.fasta	14281	16626454	Mol. Ecol. 15 (5), 1275-1297 (2006)
Populus_trichocarpa_x_Populus_nigra.mRNA.EST.fasta	5849	19121102	Physiol Plant 135 (1), 84-97 (2009)
Primula_vulgaris.mRNA.EST.fasta	19	17561923	Plant J. 51 (1), 18-31 (2007)
Prototheca_wickerhamii.mRNA.EST.fasta	5646	15862313	FEBS Lett. 579 (12), 2709-2714 (2005)
Prototheca_wickerhamii.mRNA.EST.fasta	259	15701787	Eukaryotic Cell 4 (2), 253-261 (2005)
Prototheca_wickerhamii.mRNA.EST.fasta	1	16612544	J. Mol. Evol. 62 (5), 588-599 (2006)
Prunus_armeniaca.mRNA.EST.fasta	24	16564142	Gene 374, 96-103 (2006)
Prunus_armeniaca.mRNA.EST.fasta	2	10811281	Parasitology 120 (Pt 4), 399-407 (2000)
Prunus_domestica.mRNA.EST.fasta	10	na	Phytopathol Pol 36, 91-96 (2005)
Prunus_mume.mRNA.EST.fasta	4473	20626882	BMC Genet. 11 (1), 66 (2010)
Prunus_persica.mRNA.EST.fasta	7862	18661259	Plant Mol. Biol. 68 (4-5), 379-397 (2008)
Prunus_persica.mRNA.EST.fasta	5862	16351742	BMC Bioinformatics 6 4, S16 (2005)
Prunus_persica.mRNA.EST.fasta	4983	18387211	BMC Bioinformatics 9 (Suppl 2), S9 (2008)
Prunus_persica.mRNA.EST.fasta	222	na	J. Am. Soc. Hortic. Sci. 131 (4), 551-563 (2006)
Pseudotsuga_menziesii.mRNA.EST.fasta	87	na	Theor. Appl. Genet. 97 (5-6), 771-776 (1998)
Pseudotsuga_menziesii_var__menziesii.mRNA.EST.fasta	6721	15454556	Genetics 168 (1), 447-461 (2004)
Puccinellia_tenuiflora.mRNA.EST.fasta	2366	16545489	J. Plant Physiol. 164 (1), 78-89 (2007)
Puccinellia_tenuiflora.mRNA.EST.fasta	1850	na	Plant Sci. 173 (3), 309-320 (2007)
Pueraria_montana_var__lobata.mRNA.EST.fasta	6365	na	Planta (2010) In press
Pyrus_pyrifolia.mRNA.EST.fasta	307	na	J. Gen. Plant Pathol. 75, 119-124 (2009)
Pyrus_pyrifolia.mRNA.EST.fasta	73	na	Breed. Sci. 59, 391-400 (2009)
Pyrus_pyrifolia.mRNA.EST.fasta	36	na	Sci. Hortic. 124, 195-203 (2010)
Pyrus_pyrifolia.mRNA.EST.fasta	8	na	Acta Hortic. 814, 645-649 (2009)
Quercus_mongolica_subsp__crispula.mRNA.EST.fasta	3385	18506101	Genes Genet. Syst. 83 (2), 179-187 (2008)
Quercus_petraea.mRNA.EST.fasta	811	16684234	New Phytol. 170 (4), 723-738 (2006)
Quercus_robur.mRNA.EST.fasta	26	na	New Phytol. 163 (1), 149-157 (2004)
Quercus_suber.mRNA.EST.fasta	236	17351057	Plant Physiol. 144 (1), 419-431 (2007)
Quercus_suber.mRNA.EST.fasta	2	18316306	Tree Physiol. 28 (5), 743-751 (2008)
Raphanus_sativus_var__niger.mRNA.EST.fasta	88	20553608	BMC Plant Biol. 10, 111 (2010)
Rehmannia_glutinosa.mRNA.EST.fasta	199	19774491	Mol. Biol. Rep. 37 (2), 1069-1079 (2010)
Rhododendron_catawbiense.mRNA.EST.fasta	863	15933892	Planta 221 (3), 406-416 (2005)
Rhododendron_catawbiense.mRNA.EST.fasta	378	16167139	Theor. Appl. Genet. 111 (7), 1347-1356 (2005)
Rhododendron_luteum.mRNA.EST.fasta	13	na	Mol. Breed. 23, 171-178 (2009)
Rhododendron_simsii.mRNA.EST.fasta	32	na	Mol. Breed. 23, 171-178 (2009)
Ricinus_communis.mRNA.EST.fasta	4288	17672910	BMC Plant Biol. 7, 42 (2007)
Ricinus_communis.mRNA.EST.fasta	750	12228533	Plant Physiol. 108 (3), 1141-1150 (1995)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Ricinus_communis.mRNA.EST.fasta	158	16936221	J. Exp. Bot. 57 (12), 3183-3193 (2006)
Rosa_chinensis.mRNA.EST.fasta	1794	11943190	FEBS Lett. 515 (1-3), 35-38 (2002)
Rosa_hybrid_cultivar.mRNA.EST.fasta	3511	12368489	Plant Cell 14 (10), 2325-2338 (2002)
Rubus_idaeus_subsp__strigosus.mRNA.EST.fasta	6	na	J. Am. Soc. Hortic. Sci. (2010) In press
Rubus_ulmifolius_var__inermis_x_Rubus_thyrsgiger.mRNA.EST.fasta	2678	18570660	BMC Plant Biol. 8 (1), 69 (2008)
Saccharum_hybrid_cultivar.mRNA.EST.fasta	4564	na	Genet. Mol. Biol. 24 (1-4), 1-7 (2001)
Saccharum_hybrid_cultivar.mRNA.EST.fasta	14	12679075	Mol. Phylogenet. Evol. 27 (1), 103-112 (2003)
Saccharum_hybrid_cultivar_CoS_767.mRNA.EST.fasta	13450	15454556	Genetics 168 (1), 447-461 (2004)
Saccharum_hybrid_cultivar_CoS_767.mRNA.EST.fasta	11740	19809842	Funct. Integr. Genomics Oct 7, DOI:10.1007/s10142-009-0144-9 (2009)
Saccharum_hybrid_cultivar_Co_1148.mRNA.EST.fasta	1069	15454556	Genetics 168 (1), 447-461 (2004)
Saccharum_hybrid_cultivar_Co_62175.mRNA.EST.fasta	5	na	Acta Physiol. Plant., DOI10.1007/s11738-010-0676-6 (2010) In press
Saccharum_hybrid_cultivar_Q117.mRNA.EST.fasta	8115	15316286	Plant Mol. Biol. 54 (4), 503-517 (2004)
Saccharum_hybrid_cultivar_Q117.mRNA.EST.fasta	829	na	Plant Sci. 168 (3), 761-772 (2005)
Saccharum_hybrid_cultivar_Q117.mRNA.EST.fasta	197	12856943	Plant Mol. Biol. 52 (2), 371-386 (2003)
Saccharum_hybrid_cultivar_SP70_1143.mRNA.EST.fasta	24313	na	Genet. Mol. Biol. 24 (1-4), 1-7 (2001)
Saccharum_hybrid_cultivar_SP80_3280.mRNA.EST.fasta	134261	na	Genet. Mol. Biol. 24 (1-4), 1-7 (2001)
Saccharum_hybrid_cultivar_SP80_3280.mRNA.EST.fasta	1219	12913139	Plant Physiol. 132 (4), 1811-1824 (2003)
Saccharum_hybrid_cultivar_SP80_3280.mRNA.EST.fasta	54	12733061	Mol. Genet. Genomics 269 (3), 406-419 (2003)
Saccharum_hybrid_cultivar__mixed_.mRNA.EST.fasta	73778	na	Genet. Mol. Biol. 24 (1-4), 1-7 (2001)
Salicornia_brachiata.mRNA.EST.fasta	930	19556705	Genes Genet. Syst. 84 (2), 111-120 (2009)
Salix_matsudana.mRNA.EST.fasta	17	17765407	Gene 402 (1-2), 68-80 (2007)
Salsola_tragus.mRNA.EST.fasta	400	na	Plant Signal Behav (2010) In press
Salvia_miltiorrhiza.mRNA.EST.fasta	10288	na	Sci China Life Sci 53 (1), 273-285 (2010)
Scherffelia_dubia.mRNA.EST.fasta	710	17442103	BMC Evol. Biol. 7, 65 (2007)
Scherffelia_dubia.mRNA.EST.fasta	356	11545437	Protist 152 (2), 139-147 (2001)
Sciadopitys_verticillata.mRNA.EST.fasta	11	na	Conserv. Genet. 10, 1997-1999 (2009)
Selaginella_lepidophylla.mRNA.EST.fasta	1046	na	Plant Sci. 170 (6), 1173-1184 (2006)
Selaginella_moellendorffii.mRNA.EST.fasta	2181	15938755	BMC Genomics 6 (1), 85 (2005)
Selenastrum_capricornutum.mRNA.EST.fasta	41	19462152	Bull Environ Contam Toxicol 83 (3), 301-306 (2009)
Senecio_aethnensis.mRNA.EST.fasta	131	15969730	Mol. Ecol. 14 (8), 2493-2510 (2005)
Senecio_cambrensis.mRNA.EST.fasta	146	15969730	Mol. Ecol. 14 (8), 2493-2510 (2005)
Senecio_chrysanthemifolius.mRNA.EST.fasta	115	15969730	Mol. Ecol. 14 (8), 2493-2510 (2005)
Senecio_squalidus.mRNA.EST.fasta	111	20813907	Plant Physiol. 154 (3), 1347-1360 (2010)
Senecio_squalidus_subsp__squalidus.mRNA.EST.fasta	154	15969730	Mol. Ecol. 14 (8), 2493-2510 (2005)
Senecio_vulgaris_subsp__vulgaris.mRNA.EST.fasta	132	15969730	Mol. Ecol. 14 (8), 2493-2510 (2005)
Sesbania_drummondii.mRNA.EST.fasta	100	19245025	Environ. Sci. Technol. 43 (3), 843-850 (2009)
Sesbania_rostrata.mRNA.EST.fasta	102	na	Plant Sci. 167 (2), 207-216 (2004)
Sesbania_rostrata.mRNA.EST.fasta	87	18557786	FEMS Microbiol. Lett. 285 (1), 16-24 (2008)
Sesbania_rostrata.mRNA.EST.fasta	66	na	Plant Sci. 162 (6), 915-921 (2002)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Sesbania_rostrata.mRNA.EST.fasta	19	11522814	Nucleic Acids Res. 29 (17), 3459-3468 (2001)
Setaria_italica.mRNA.EST.fasta	2097	17498921	Genomics 90 (1), 121-131 (2007)
Setaria_italica.mRNA.EST.fasta	327	20171162	Biochem. Biophys. Res. Commun. 393 (4), 720-727 (2010)
Setaria_italica.mRNA.EST.fasta	290	na	J. Plant Physiol. 168 (3), 280-287 (2011)
Setaria_italica.mRNA.EST.fasta	27	18592419	Mol. Biotechnol. 40 (3), 241-251 (2008)
Silene_latifolia.mRNA.EST.fasta	3662	19467153	BMC Genomics 10 (1), 243 (2009)
Simmondsia_chinensis.mRNA.EST.fasta	385	na	J. Biol. Chem. 63 (1-3), 137-146 (2008)
Solanum_chacoense.mRNA.EST.fasta	7751	16235114	Plant Mol. Biol. 59 (3), 515-532 (2005)
Solanum_chacoense.mRNA.EST.fasta	1	16461387	Plant Physiol. 140 (4), 1374-1383 (2006)
Solanum_habrochaites.mRNA.EST.fasta	17038	21098679	Plant Physiol. 155 (1), 524-539 (2011)
Solanum_lycopersicum.mRNA.EST.fasta	57422	20350329	BMC Genomics 11, 210 (2010)
Solanum_lycopersicum.mRNA.EST.fasta	35824	15975739	Gene 356, 127-134 (2005)
Solanum_lycopersicum.mRNA.EST.fasta	27216	16307365	Plant Mol. Biol. 59 (6), 895-907 (2005)
Solanum_lycopersicum.mRNA.EST.fasta	4336	16273343	Theor. Appl. Genet. 112 (1), 114-130 (2005)
Solanum_lycopersicum.mRNA.EST.fasta	4045	21098679	Plant Physiol. 155 (1), 524-539 (2011)
Solanum_lycopersicum.mRNA.EST.fasta	774	12669794	Genome 46 (1), 28-47 (2003)
Solanum_lycopersicum.mRNA.EST.fasta	663	na	Postharvest Biol. Technol. 50 (2), 117-124 (2008)
Solanum_lycopersicum.mRNA.EST.fasta	596	17210988	J. Exp. Bot. 58 (3), 507-520 (2007)
Solanum_lycopersicum.mRNA.EST.fasta	344	9724330	Genome Res. 8 (8), 842-847 (1998)
Solanum_lycopersicum.mRNA.EST.fasta	252	na	Plant Sci. 173 (6), 609-620 (2007)
Solanum_lycopersicum.mRNA.EST.fasta	193	16059720	Planta 222 (6), 947-956 (2005)
Solanum_lycopersicum.mRNA.EST.fasta	177	na	Physiol. Mol. Plant Pathol. 73, 67-77 (2009)
Solanum_lycopersicum.mRNA.EST.fasta	166	11553762	Plant Physiol. 127 (1), 345-359 (2001)
Solanum_lycopersicum.mRNA.EST.fasta	156	na	Plant Sci. 170 (5), 976-985 (2006)
Solanum_lycopersicum.mRNA.EST.fasta	141	19236572	Mol. Plant Pathol. 10 (2), 237-248 (2009)
Solanum_lycopersicum.mRNA.EST.fasta	117	na	Euphytica 12 (2), 229-236 (2003)
Solanum_lycopersicum.mRNA.EST.fasta	106	20618710	Mol. Plant Pathol. 11 (4), 531-544 (2010)
Solanum_lycopersicum.mRNA.EST.fasta	67	15553246	Mol. Plant Microbe Interact. 17 (11), 1212-1222 (2004)
Solanum_lycopersicum.mRNA.EST.fasta	51	12913147	Plant Physiol. 132 (4), 1901-1912 (2003)
Solanum_lycopersicum.mRNA.EST.fasta	50	na	Postharvest Biol. Technol. 53 (3), 131-137 (2009)
Solanum_lycopersicum.mRNA.EST.fasta	35	12410809	Plant J. 32 (3), 299-315 (2002)
Solanum_lycopersicum.mRNA.EST.fasta	22	17153921	Mol. Plant Microbe Interact. 19 (12), 1368-1377 (2006)
Solanum_lycopersicum.mRNA.EST.fasta	20	11011101	Plant Sci. 159 (1), 135-148 (2000)
Solanum_lycopersicum.mRNA.EST.fasta	6	10417710	Plant J. 18 (6), 589-600 (1999)
Solanum_lycopersicum.mRNA.EST.fasta	6	20459311	Mol. Plant Microbe Interact. 23 (6), 715-726 (2010)
Solanum_lycopersicum.mRNA.EST.fasta	2	20096428	Phytochemistry 71 (5-6), 605-613 (2010)
Solanum_lycopersicum.mRNA.EST.fasta	2	8980540	Plant Mol. Biol. 32 (5), 891-900 (1996)
Solanum_lycopersicum_var__cerasiforme.mRNA.EST.fasta	1160	16183847	Plant Physiol. 139 (2), 750-769 (2005)
Solanum_melongena.mRNA.EST.fasta	98086	19857557	Gene 450 (1-2), 76-84 (2010)
Solanum_pennellii.mRNA.EST.fasta	2600	21098679	Plant Physiol. 155 (1), 524-539 (2011)
Solanum_sogarandinum.mRNA.EST.fasta	16	na	Plant Sci. 124 (1), 68-78 (1997)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Solanum_sogarandinum.mRNA.EST.fasta	9	na	Plant Sci. 133 (1), 57-67 (1998)
Solanum_torvum.mRNA.EST.fasta	28379	19837731	J. Exp. Bot. 61 (2), 423-427 (2010)
Solanum_tuberosum.mRNA.EST.fasta	20758	16094426	Genome 48 (4), 598-605 (2005)
Solanum_tuberosum.mRNA.EST.fasta	9881	na	Am. J. Potato Res. 87 (1), 50-59 (2010)
Solanum_tuberosum.mRNA.EST.fasta	6075	11591384	FEBS Lett. 506 (2), 123-126 (2001)
Solanum_tuberosum.mRNA.EST.fasta	703	na	Physiol. Mol. Plant Pathol. 75 (1-2), 3-12 (2010)
Solanum_tuberosum.mRNA.EST.fasta	533	14966214	J. Exp. Bot. 55 (397), 613-622 (2004)
Solanum_tuberosum.mRNA.EST.fasta	382	na	Physiol. Mol. Plant Pathol. 67 (3-5), 237-247 (2006)
Solanum_tuberosum.mRNA.EST.fasta	316	19236574	Mol. Plant Pathol. 10 (2), 263-275 (2009)
Solanum_tuberosum.mRNA.EST.fasta	182	na	Mol. Plant Pathol. 5 (3), 191-201 (2004)
Solanum_tuberosum.mRNA.EST.fasta	166	12753591	Plant J. 34 (4), 529-541 (2003)
Solanum_tuberosum.mRNA.EST.fasta	153	na	Plant Sci. 166 (2), 423-433 (2004)
Solanum_tuberosum.mRNA.EST.fasta	113	21249504	Planta Jan 20, DOI:10.1007/s00425-011-1350-y (2011) In press
Solanum_tuberosum.mRNA.EST.fasta	77	16738852	Plant Cell Rep. 25 (10), 1094-1103 (2006)
Solanum_tuberosum.mRNA.EST.fasta	36	na	Mol. Plant Microbe Interact. 12 (4), 356-361 (1999)
Solanum_tuberosum.mRNA.EST.fasta	29	17609532	J. Exp. Bot. 58 (11), 2761-2774 (2007)
Solanum_tuberosum.mRNA.EST.fasta	26	na	J Integr Plant Biol 52 (5), 453-467 (2010)
Solanum_tuberosum.mRNA.EST.fasta	1	na	Plant Physiol. 155 (1), 421-432 (2011)
Sorghum_bicolor.mRNA.EST.fasta	5468	17998204	J. Biol. Chem. 283 (6), 3231-3247 (2007)
Sorghum_bicolor.mRNA.EST.fasta	269	16292568	Planta 223 (5), 932-947 (2005)
Sorghum_bicolor.mRNA.EST.fasta	89	na	Theor. Appl. Genet. 97 (5-6), 771-776 (1998)
Sorghum_bicolor.mRNA.EST.fasta	82	14701914	Plant Physiol. 134 (1), 420-431 (2004)
Sorghum_bicolor.mRNA.EST.fasta	47	14760522	Mol. Genet. Genomics 271 (3), 308-316 (2004)
Sorghum_bicolor.mRNA.EST.fasta	38	na	Weed Sci. 56, 356-363 (2008)
Sorghum_bicolor.mRNA.EST.fasta	14	9738887	Mol. Gen. Genet. 259 (1), 115-122 (1998)
Sorghum_bicolor.mRNA.EST.fasta	4	9617804	Plant Mol. Biol. 37 (2), 319-335 (1998)
Sorghum_bicolor.mRNA.EST.fasta	3	16473662	J. Plant Physiol. 163 (5), 570-576 (2006)
Sorghum_halepense.mRNA.EST.fasta	729	16998090	Plant Physiol. 142 (3), 1148-1159 (2006)
Sorghum_propinquum.mRNA.EST.fasta	393	16998090	Plant Physiol. 142 (3), 1148-1159 (2006)
Spartina_alterniflora.mRNA.EST.fasta	1227	18305970	Funct. Integr. Genomics 8 (3), 287-300 (2008)
Spartina_alterniflora.mRNA.EST.fasta	28	na	Plant Sci. 170 (6), 1141-1149 (2006)
Spirogyra_pratensis.mRNA.EST.fasta	7294	20500869	BMC Plant Biol. 10 (1), 96 (2010)
Stenogyne_rugosa.mRNA.EST.fasta	666	16928278	BMC Plant Biol. 6, 16 (2006)
Streptocarpus_rexii.mRNA.EST.fasta	1	15659624	Plant Cell 17 (2), 430-443 (2005)
Suaeda_maritima.mRNA.EST.fasta	169	19497134	BMC Plant Biol. 9 (1), 69 (2009)
Syntrichia_ruralis.mRNA.EST.fasta	9159	15546486	BMC Genomics 5 (1), 89 (2004)
Syntrichia_ruralis.mRNA.EST.fasta	685	19523028	Physiol Plant 136 (4), 437-460 (2009)
Syntrichia_ruralis.mRNA.EST.fasta	141	10394632	Plant Cell Physiol. 40 (4), 361-368 (1999)
Syntrichia_ruralis.mRNA.EST.fasta	6	na	Bryologist 103 (1), 44-51 (2000)
Tagetes_erecta.mRNA.EST.fasta	2	10571881	Plant J. 20 (2), 217-230 (1999)
Taiwania_cryptomerioides.mRNA.EST.fasta	436	na	Plant Sci. 167 (4), 955-957 (2004)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Tamarix_androssowii.mRNA.EST.fasta	2566	na	Plant Sci. 170 (1), 28-36 (2006)
Tamarix_androssowii.mRNA.EST.fasta	18	15599053	Zhi Wu Sheng Li Yu Fen Zi Sheng Wu Xue Xue Bao 30 (2), 229-233 (2004)
Tamarix_hispida.mRNA.EST.fasta	9447	18058243	Plant Mol. Biol. 66 (3), 245-258 (2008)
Tamarix_hispida.mRNA.EST.fasta	7726	19146931	Gene 433 (1-2), 65-71 (2009)
Taxus_cuspidata.mRNA.EST.fasta	22	11876470	Mol. Biotechnol. 20 (2), 137-143 (2002)
Theobroma_cacao.mRNA.EST.fasta	85327	18973681	BMC Genomics 9 (1), 512 (2008)
Theobroma_cacao.mRNA.EST.fasta	4443	12447539	Planta 216 (2), 255-264 (2002)
Theobroma_cacao.mRNA.EST.fasta	2927	17557832	Ann. Bot. 100 (1), 129-140 (2007)
Theobroma_cacao.mRNA.EST.fasta	2114	15340758	Plant Cell Rep. 23 (6), 404-413 (2004)
Theobroma_cacao.mRNA.EST.fasta	187	na	Mol. Plant Pathol. 8 (3), 279-292 (2007)
Theobroma_cacao.mRNA.EST.fasta	31	15933892	Planta 221 (3), 406-416 (2005)
Theobroma_cacao.mRNA.EST.fasta	16	16832690	Planta 224 (6), 1449-1464 (2006)
Theobroma_cacao.mRNA.EST.fasta	6	15979314	Plant Physiol. Biochem. 43 (6), 611-622 (2005)
Thlaspi_arvense.mRNA.EST.fasta	647	16972165	Plant Mol. Biol. 63 (2), 171-184 (2007)
Torenia_fournieri.mRNA.EST.fasta	256	19295610	Nature 458 (7236), 357-361 (2009)
Trifolium_pratense.mRNA.EST.fasta	38109	16769692	DNA Res. 12 (5), 301-364 (2005)
Triphysaria_versicolor.mRNA.EST.fasta	7220	16288663	BMC Plant Biol. 5, 24 (2005)
Triphysaria_versicolor.mRNA.EST.fasta	2085	11553755	Plant Physiol. 127 (1), 272-282 (2001)
Triticum_aestivum.mRNA.EST.fasta	243462	16832693	Mol. Genet. Genomics 276 (3), 304-312 (2006)
Triticum_aestivum.mRNA.EST.fasta	116232	16306141	Plant Physiol. 139 (4), 1870-1880 (2005)
Triticum_aestivum.mRNA.EST.fasta	4433	12590341	Funct. Integr. Genomics 3 (1-2), 33-38 (2003)
Triticum_aestivum.mRNA.EST.fasta	2743	19995415	BMC Genomics 10 (1), 586 (2009)
Triticum_aestivum.mRNA.EST.fasta	2564	na	Plant Mol. Biol. Rep. 18, 123-132 (2000)
Triticum_aestivum.mRNA.EST.fasta	1274	na	Physiol Plant 129 (2), 375-385 (2007)
Triticum_aestivum.mRNA.EST.fasta	1216	16391697	Genome 48 (5), 913-923 (2005)
Triticum_aestivum.mRNA.EST.fasta	1194	15703054	Plant J. 41 (5), 660-672 (2005)
Triticum_aestivum.mRNA.EST.fasta	179	17207255	Plant Biotechnol. J. 5 (1), 38-49 (2007)
Triticum_aestivum.mRNA.EST.fasta	177	16027977	Plant Mol. Biol. 58 (2), 247-267 (2005)
Triticum_aestivum.mRNA.EST.fasta	171	16053709	J. Biochem. Mol. Biol. 38 (4), 420-431 (2005)
Triticum_aestivum.mRNA.EST.fasta	99	na	Theor. Appl. Genet. 78, 495-504 (1989)
Triticum_aestivum.mRNA.EST.fasta	82	19171027	BMC Genomics 10 (1), 48 (2009)
Triticum_aestivum.mRNA.EST.fasta	72	na	J Integr Plant Biol 47 (7), 839-848 (2005)
Triticum_aestivum.mRNA.EST.fasta	31	na	Physiol. Mol. Plant Pathol. 71 (4-6), 251-259 (2007)
Triticum_aestivum.mRNA.EST.fasta	20	19453509	Physiol Plant 136 (2), 209-222 (2009)
Triticum_aestivum.mRNA.EST.fasta	15	12582923	Theor. Appl. Genet. 105 (6-7), 972-979 (2002)
Triticum_aestivum.mRNA.EST.fasta	14	na	Plant Syst. Evol. 278 (1-2), 125-131 (2009)
Triticum_aestivum.mRNA.EST.fasta	14	19082576	Theor. Appl. Genet. 118 (4), 753-764 (2009)
Triticum_aestivum.mRNA.EST.fasta	13	16893592	J. Plant Physiol. 164 (7), 913-922 (2007)
Triticum_aestivum.mRNA.EST.fasta	11	na	Physiol. Mol. Plant Pathol. 65 (1), 21-30 (2004)
Triticum_aestivum.mRNA.EST.fasta	10	18592282	Funct. Integr. Genomics 8 (4), 421-427 (2008)
Triticum_aestivum.mRNA.EST.fasta	9	12587874	Pest Manag. Sci. 59 (2), 202-214 (2003)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Triticum_aestivum.mRNA.EST.fasta	7	na	Mai Lei Zuo Wu Xue Bao 28 (2), 249-253 (2008)
Triticum_aestivum.mRNA.EST.fasta	6	16636822	Funct. Integr. Genomics 7 (1), 69-77 (2007)
Triticum_aestivum.mRNA.EST.fasta	3	12175074	Genome 45 (4), 706-718 (2002)
Triticum_aestivum.mRNA.EST.fasta	2	na	Theor. Appl. Genet. 97 (3), 356-369 (1998)
Triticum_aestivum.mRNA.EST.fasta	2	10102366	Mol. Gen. Genet. 261 (2), 307-316 (1999)
Triticum_aestivum.mRNA.EST.fasta	1	19154231	Plant Biotechnol. J. 7 (2), 200-209 (2009)
Triticum_aestivum.mRNA.EST.fasta	1	na	Mol. Breed. 14 (4), 349-361 (2004)
Triticum_aestivum.mRNA.EST.fasta	1	na	Mol. Plant Breed. 3 (6), 857-863 (2005)
Triticum_aestivum_Thinopyrum_intermedium_alien_addition_line.mRNA.EST.fasta	9	15644969	Genome 47 (6), 1114-1121 (2004)
Triticum_turgidum_subsp__durum.mRNA.EST.fasta	432	na	Minerva Biotechnologica 18 (4), 159-164 (2006)
Triticum_turgidum_subsp__durum.mRNA.EST.fasta	48	19171027	BMC Genomics 10 (1), 48 (2009)
Tropaeolum_majus.mRNA.EST.fasta	10507	17488821	Proc. Natl. Acad. Sci. U.S.A. 104 (20), 8550-8555 (2007)
Ulmus_americana.mRNA.EST.fasta	535	na	Physiol. Mol. Plant Pathol. 74 (3-4), 251-266 (2010)
Ulva_fasciata.mRNA.EST.fasta	78	na	Mar. Biotechnol. (2011) In press
Ulva_linza.mRNA.EST.fasta	1898	na	J. Phycol. 41 (6), 1219-1226 (2005)
Vaccinium_corymbosum.mRNA.EST.fasta	1282	na	Mol Biol (Mosk) 166 (4), 863-872 (2004)
Vanda_hybrid_cultivar.mRNA.EST.fasta	2133	21116862	Mol. Biol. Rep. DOI: 10.1007, 10.1007/s11033-010-0506-3 (2010) In press
Vicia_monantha.mRNA.EST.fasta	53	na	Egypt J Genet Cytol 38, 137-152 (2009)
Vicia_villosa.mRNA.EST.fasta	25	na	Egypt J Genet Cytol 39 (2010) In press
Vigna_radiata.mRNA.EST.fasta	811	18060406	Plant Cell Rep. 27 (3), 535-552 (2008)
Vigna_unguiculata.mRNA.EST.fasta	104	18095375	Proteomics 8 (1), 149-159 (2008)
Vigna_unguiculata.mRNA.EST.fasta	18	11943192	FEBS Lett. 515 (1-3), 44-50 (2002)
Viola_baoshanensis.mRNA.EST.fasta	43	19106016	J. Plant Physiol. 166 (8), 862-870 (2009)
Vitis_arizonica_x_Vitis_rupestris.mRNA.EST.fasta	5421	17316447	BMC Plant Biol. 7, 8 (2007)
Vitis_berlandieri_x_Vitis_riparia.mRNA.EST.fasta	168	19660119	BMC Genomics 10 (1), 363 (2009)
Vitis_berlandieri_x_Vitis_riparia.mRNA.EST.fasta	28	20192835	Mol. Plant Microbe Interact. 23 (4), 485-496 (2010)
Vitis_hybrid_cultivar.mRNA.EST.fasta	6	na	Plant Growth Regul. 61 (2), 127-134 (2010)
Vitis_labrusca.mRNA.EST.fasta	5	na	Plant Sci. 162, 621-628 (2002)
Vitis_vinifera.mRNA.EST.fasta	77583	17761391	Gene 402 (1-2), 40-50 (2007)
Vitis_vinifera.mRNA.EST.fasta	15796	na	Plant Sci. 173 (4), 446-457 (2007)
Vitis_vinifera.mRNA.EST.fasta	7331	15856347	Funct. Integr. Genomics 5 (4), 208-217 (2005)
Vitis_vinifera.mRNA.EST.fasta	760	na	Plant Sci. 179 (5), 510-519 (2010)
Vitis_vinifera.mRNA.EST.fasta	275	na	J. Plant Physiol. 158 (12), 1575-1583 (2001)
Vitis_vinifera.mRNA.EST.fasta	193	na	Eur. J. Plant Pathol. 129, 281-301 (2011)
Vitis_vinifera.mRNA.EST.fasta	170	11011096	Plant Sci. 159 (1), 87-95 (2000)
Vitis_vinifera.mRNA.EST.fasta	24	na	Plant Sci. 171 (1), 132-138 (2006)
Vitis_vinifera.mRNA.EST.fasta	7	na	Plant Mol. Biol. Rep. 22, 225-236 (2004)
Vitis_vinifera.mRNA.EST.fasta	2	16134894	Mol. Plant Microbe Interact. 18 (8), 819-829 (2005)
Vitis_vinifera.mRNA.EST.fasta	1	na	Vitis 40, 191-196 (2001)
Vitis_vinifera.mRNA.EST.fasta	1	17942184	J. Plant Physiol. 165 (5), 471-481 (2008)

File name (species, subspecies, cultivars, etc.)	Sequences	PMID	Source
Xerophyta_humilis.mRNA.EST.fasta	347	na	Physiol Plant 122, 39-53 (2004)
Xerophyta_humilis.mRNA.EST.fasta	53	na	Physiol Plant 122 (1), 39-53 (2004)
Zea_mays.mRNA.EST.fasta	537518	18937034	Plant Mol. Biol. 69 (1-2), 179-194 (2009)
Zea_mays.mRNA.EST.fasta	287917	17662031	Plant J. 51 (5), 910-918 (2007)
Zea_mays.mRNA.EST.fasta	260736	17095711	Genome Res. 17 (1), 69-73 (2007)
Zea_mays.mRNA.EST.fasta	30553	16247562	Plant Mol. Biol. 59 (2), 363-374 (2005)
Zea_mays.mRNA.EST.fasta	23300	15466291	Genome Res. 14 (10), 1932-1937 (2004)
Zea_mays.mRNA.EST.fasta	16762	na	Environ. Exp. Bot. 69 (2), 158-166 (2010)
Zea_mays.mRNA.EST.fasta	15867	18803597	Plant Biotechnol. J. 6 (9), 855-869 (2008)
Zea_mays.mRNA.EST.fasta	14822	18024554	Plant Physiol. 146 (1), 32-44 (2008)
Zea_mays.mRNA.EST.fasta	5174	12787250	Plant J. 34 (5), 697-707 (2003)
Zea_mays.mRNA.EST.fasta	4070	16718485	Planta 224 (5), 1004-1014 (2006)
Zea_mays.mRNA.EST.fasta	1064	8012409	Plant J. 5, 605-610 (1994)
Zea_mays.mRNA.EST.fasta	932	7811968	Plant Mol. Biol. 26, 1085-1101 (1994)
Zea_mays.mRNA.EST.fasta	701	17766395	Plant Physiol. 145 (3), 575-588 (2007)
Zea_mays.mRNA.EST.fasta	190	15604718	Plant Mol. Biol. 55 (6), 807-823 (2004)
Zea_mays.mRNA.EST.fasta	111	na	Plant Mol. Biol. 31, 23-34 (1996)
Zea_mays.mRNA.EST.fasta	75	14657407	Plant Physiol. 134 (1), 380-387 (2004)
Zea_mays.mRNA.EST.fasta	63	15856942	Mol. Biol. (Mosk.) 39 (2), 198-206 (2005)
Zea_mays.mRNA.EST.fasta	62	16167904	Plant J. 44 (1), 167-178 (2005)
Zea_mays.mRNA.EST.fasta	60	17082872	Plant Mol. Biol. 63 (3), 381-391 (2007)
Zea_mays.mRNA.EST.fasta	19	15695436	Plant Cell Physiol. 46 (2), 332-338 (2005)
Zea_mays.mRNA.EST.fasta	18	10634908	Plant Cell 12 (1), 65-80 (2000)
Zea_mays.mRNA.EST.fasta	17	17869108	Curr. Biol. 17 (19), 1686-1691 (2007)
Zea_mays.mRNA.EST.fasta	5	12124387	J. Biol. Chem. 277 (40), 37741-37746 (2002)
Zea_mays.mRNA.EST.fasta	3	na	Sex. Plant Reprod. 13, 21-27 (2000)
Zingiber_officinale.mRNA.EST.fasta	10	17928987	Theor. Appl. Genet. 116 (1), 123-134 (2007)
Zingiber_zerumbet.mRNA.EST.fasta	10	17928987	Theor. Appl. Genet. 116 (1), 123-134 (2007)
Zinnia_violacea.mRNA.EST.fasta	17461	12438691	Proc. Natl. Acad. Sci. U.S.A. 99 (24), 15794-15799 (2002)
Zinnia_violacea.mRNA.EST.fasta	446	16306148	Plant Physiol. 139 (4), 1821-1839 (2005)
Zinnia_violacea.mRNA.EST.fasta	62	12417703	Plant Cell 14 (11), 2813-2824 (2002)
Zostera_marina.mRNA.EST.fasta	7226	18239962	Mar. Biotechnol. 10 (3), 297-309 (2008)
Zostera_marina.mRNA.EST.fasta	1161	18086234	Plant Biotechnol. J. 6 (3), 246-263 (2008)
Zostera_marina.mRNA.EST.fasta	15	na	Mar. Biol. 157 (1), 81-89 (2010)
Zostera_marina.mRNA.EST.fasta	14	na	Mol. Ecol. Notes 7 (5), 777-780 (2007)