

1 S3 Table

Novel ID	Location	Strand	Sequence of 3'	Sequence of 5'	raw data				nomalized data			
					C100	C130	L100	L130	C100	C130	L100	L130
Oe_mir_1	FKYM01055610.1_2875:2902	-		CAGATGCAATCTGCACCACCCAGGGTT	6	0	0	4	3	0	0	1
Oe_mir_2	FKYM01055214.1_1219:1247	+	GAGAGGGATGGGGTTTCTCTCGCTTTTGG		14	0	0	0	6	0	0	0
Oe_mir_3	FKYM01054197.1_28858:28886	+		TTGGGCTCCGGGAGGGGAGATGTAAGAA	36	0	0	0	15	0	0	0
Oe_mir_4	FKYM01050891.1_28768:28796	-		TGTGCTGGGAGTCCCTAATGATTAATA	4	0	0	0	2	0	0	0
Oe_mir_5	FKYM01050338.1_34826:34853	+		GCTTTTGTACCGAGGGTTCGAATCCC	3	3	0	0	2	1	0	0
Oe_mir_6	FKYM01046616.1_5824:5851	-		TGATCTGCCGAAGCCGTGGGATGAAAAA	22	0	0	0	9	0	0	0
Oe_mir_7	FKYM01046010.1_24248:24276	-		CTTGAAGATTATAGAATGTTCTGAGGCC	5	0	0	0	2	0	0	0
Oe_mir_8	FKYM01044337.1_45579:45608	+	TTCTGATGATAAAAAGCGAATCTGAGCCTC		93	86	0	15	38	12	0	4
Oe_mir_9	FKYM01043744.1_6792:6820	-	CAATGATCAGTAGCTGGTTCGAGAGGATG		3	0	0	0	2	0	0	0
Oe_mir_10	FKYM01042526.1_21612:21641	-	CTGTAGCTCAGTTGGTAGAGCACCTCGAAA		23	0	0	0	10	0	0	0
Oe_mir_11	FKYM01041785.1_40379:40407	-		AGTTTAGGTGAGGTATATTTCTGAGCCAG	4	0	0	0	2	0	0	0
Oe_mir_12	FKYM01039660.1_42011:42040	-	AGTAGAGTTGAATAGCGGAAGATCCAATA		3	0	0	0	2	0	0	0
Oe_mir_13	FKYM01038462.1_2970:2999	+	CCCTTAGATGTTCTGGGCCGACGCGCGCT		5	0	0	0	2	0	0	0
Oe_mir_14	FKYM01036130.1_5095:5123	+	TATTTGTACAGATATGGTAGAGGGCACA		27	0	0	0	11	0	0	0
Oe_mir_15	FKYM01035635.1_17875:17902	+	TTGGCATTGAGTTCATGGACCTGTACAG		4	0	0	0	2	0	0	0
Oe_mir_16	FKYM01035436.1_18874:18877	+		GTGCCAAGTGGTAAGGCAACGGGTTTT	35	0	0	0	14	0	0	0
Oe_mir_17	FKYM01033727.1_28864:28894	-	CGTAGCCGCTCATGGTTATTTGGCCGATTA		6	0	0	0	3	0	0	0
Oe_mir_18	FKYM01033545.1_10576:10604	+	ACCCCCATGTGAGAGCCCTGTACAGACC		4	0	0	0	2	0	0	0
Oe_mir_19	FKYM01033405.1_36717:36744	-		GAGAGAAGAGCTGGAGCTGAAAATGAGT	3	3	0	0	2	1	0	0
Oe_mir_20	FKYM01031732.1_27775:27803	-		TAGACCTTGTTGTGTCAGAATTCTTAAT	3	0	0	0	2	0	0	0
Oe_mir_21	FKYM01031203.1_21800:21828	+	TCACTGGTCTAGCTCCATGGCACTAAAAA		7	0	0	6	3	0	0	2
Oe_mir_22	FKYM01031021.1_16233:16261	+		GGACCAGACAGGCGCAGGTAGCTTGACCG	5	0	0	0	2	0	0	0
Oe_mir_23	FKYM01030694.1_9539:9567	-	CGAGGGCTACTATAGTTGATATGTGGAGC		7	26	0	0	3	4	0	0
Oe_mir_24	FKYM01030229.1_16611:16641	-	CCGCGCGATGTCGACGTGAAGAGGAATGCT		51	0	0	0	21	0	0	0
Oe_mir_25	FKYM01029890.1_12726:12756	-	GATGTCGATCCTGAGCGGTTACCTGAGCAAC		23	0	0	15	10	0	0	4
Oe_mir_26	FKYM01025752.1_3968:3999	+	CTTGTGATTGAGAATCGTATCTGTGGCTACT		11	0	0	0	5	0	0	0
Oe_mir_27	FKYM01024763.1_626:656	-	GGGGATGCCGAAGGCAGGCTAGTACTGGA		12	0	0	0	5	0	0	0
Oe_mir_28	FKYM01022360.1_5031:5060	-	CTTGGGCGAGAGTAGTACTAGGATGGGTGA		78	0	0	0	32	0	0	0
Oe_mir_29	FKYM01022360.1_11953:11982	-	CCCCCTGGGAAGTCTCGTGTGCACCCCT		19	150	0	0	8	21	0	0

Oe_mir_30	FKYM01022360.1_16605:16636	-		TAGTACTAGGATGGGTGACCCCTGGGAAGTT	1	0	0	0	1	0	0	0
Oe_mir_31	FKYM01022360.1_16257:16286	-	CCCCCTGGGAAGTCCTCGTGTGTACCCT		8	0	0	0	4	0	0	0
Oe_mir_32	FKYM01022358.1_15344:15372	-		CCCCTGTGAAGTCCTCGTGTGCACCCCT	9	131	34	29	4	18	11	8
Oe_mir_33	FKYM01022358.1_20325:20353	-		CCCCTGGGAAGTTCTCGTGTGCACCCCT	5	48	0	0	2	7	0	0
Oe_mir_34	FKYM01022358.1_31057:31084	-	AGTAGTACTAGGATGGGTGACCCCTCTGG		3	0	0	4	2	0	0	1
Oe_mir_35	FKYM01020931.1_11846:11876	+		AGCCATTCGGCGGTGAATTCGTTCCCGGGCC	5	0	0	0	2	0	0	0
Oe_mir_36	FKYM01014275.1_38979:39010	+		ATTTGGTCTAGTGGTATGATTCTCGCTTTGGG	4	0	0	0	2	0	0	0
Oe_mir_37	FKYM01014067.1_11568:11599	+	AGGTGTAGCAATAATGAACGCTGGTGTATGTTG		12	20	0	0	5	3	0	0
Oe_mir_38	FKYM01007076.1_17288:17316	+	GTGAGGCTGGGCGACCTGATGAGGTGGCA		12	0	0	6	5	0	0	2
Oe_mir_39	FKYM01006094.1_22386:22416	+	AGTGGTAAGAATCCACGTTGTGGCCGTGGA		24	0	0	0	10	0	0	0
Oe_mir_40	FKYM01002722.1_3841:3872	+		GCCTCGTGACTCCGTTGTCTGAATACTGTAT	3	0	0	0	2	0	0	0
Oe_mir_41	FKYM01002369.1_752:781	-	TTATTGTCTATCGTCGGCCCTATGGTAGA		18	0	0	0	8	0	0	0
Oe_mir_42	FKYM01000663.1_3464:3493	-	TACTTCTCCTAGCTGCTTGGCCTGTAGTA		117	0	0	0	47	0	0	0
Oe_mir_43	FKYM01052587.1_72930:72958	+		TGCATCGCGTGACAAATCTATTAAGCTTT	5	24	0	0	2	3	0	0
Oe_mir_44	FKYM01051081.1_21848:21875	-	AGCGACATTATTGTGGAATGTAGACAC		12	0	0	0	5	0	0	0
Oe_mir_45	FKYM01046624.1_613:643	-		CAGACCAAGAACTACGAGATCACCCCTTTCA	16	0	0	0	7	0	0	0
Oe_mir_46	FKYM01044697.1_17668:17695	-	TCTGATTGTGTTCTTTTTTGGTGACAG		7	45	14	6	3	6	5	2
Oe_mir_47	FKYM01044337.1_45577:45608	+	TCTTCTGATGATAAAAGGCGAATCTGAGCCTC		45	0	0	0	18	0	0	0
Oe_mir_48	FKYM01043238.1_8907:8935	+	GTAGCTCATATGGTAGGTCGTTAGTTTTG		4	0	0	0	2	0	0	0
Oe_mir_49	FKYM01042526.1_21613:21641	-	CTGTAGCTCAGTTGGTAGAGCACCTCGAA		38	62	64	44	16	9	21	12
Oe_mir_50	FKYM01040697.1_6755:6784	-		ACTTCGACCAGCGAACTCGAACAGAATTCC	3	0	0	0	2	0	0	0
Oe_mir_51	FKYM01038316.1_13055:13084	-	CTGGAGTTATAACATTTGAATTTCTCTGAC		3	0	0	0	2	0	0	0
Oe_mir_52	FKYM01036652.1_25672:25699	+		CTTGGTGGTGAAATGGTAGACACGCGAG	6	0	0	18	3	0	0	5
Oe_mir_53	FKYM01027114.1_15577:15604	-	AATACTGCTAGACAGGAACATAAGTATG		3	0	0	0	2	0	0	0
Oe_mir_54	FKYM01025620.1_64968:64996	-		TGATATCGTTGTTGGGTTGCAGGAAG	3	0	4	0	2	0	2	0
Oe_mir_55	FKYM01024763.1_629:656	-	GGGGATGCCGAAGGCAGGGCTAGTACT		5	0	0	0	2	0	0	0
Oe_mir_56	FKYM01024477.1_73148:73178	-	CTGGATGATTTCCGACCAGGCTTCATTCCCC		3	0	0	0	2	0	0	0
Oe_mir_57	FKYM01022360.1_11952:11982	-		CCCCCTGGGAAGTCCTCGTGTGCACCCCTT	66	0	0	0	27	0	0	0
Oe_mir_58	FKYM01022360.1_16257:16285	-	CCCCTGGGAAGTCCTCGTGTGTACCCT		15	0	56	21	6	0	18	6
Oe_mir_59	FKYM01022360.1_21160:21189	-		TGAGAGTAGTACTAGGATGGGTGACCCCT	83	0	0	0	34	0	0	0
Oe_mir_60	FKYM01022358.1_10676:10706	-		CCCCTGGGAAGTCCTCGTGTGCCCCCTAT	6	0	0	0	3	0	0	0
Oe_mir_61	FKYM01018308.1_24988:25016	+	TTCAGCAACGGATTTCTTGGCTCTCGCAT		3	0	0	0	2	0	0	0

Oe_mir_62	FKYM01018276.1_6486:6513	+		CCGAGTTGGCGGCTGAGTTGGATTCTAA	3	0	0	0	2	0	0	0
Oe_mir_63	FKYM01010170.1_31901:31928	-	CAGTTATAATCCAATTTCTCTGATGGTT		18	0	40	0	8	0	13	0
Oe_mir_64	FKYM01002444.1_19730:19757	+		GTTGTCTGGCTCGAGAACTCTTTTGT	1	0	0	0	1	0	0	0
Oe_mir_65	FKYM01002444.1_19792:19819	+	GCGTTGTCGGACCAGGCTTCATCCCCC		171	421	185	95	69	59	60	25
Oe_mir_66	FKYM01001755.1_27381:27411	+		GAATTGATGAGCCGATGCGGTCTGAGTCCA	3	0	0	0	2	0	0	0
Oe_mir_67	FKYM01001671.1_2100:2131	+		GTGCAAAGGTTTCTCGGGCCGGACGGAGATT	674	0	0	0	271	0	0	0
Oe_mir_68	FKYM01000663.1_3463:3493	-	TACTTCTCCTAGCTGCTTGGCCTGTAGTAG		15	0	0	0	6	0	0	0
Oe_mir_69	FKYM01057934.1_3444:3472	+		AGTGGTGTTCAGTGGCGGACGGGTGAG	11	0	0	0	5	0	0	0
Oe_mir_70	FKYM01045039.1_60625:60654	-		CTACTTAACTCAGTGGTTAGAGTATTGCTT	4	0	6	0	2	0	2	0
Oe_mir_71	FKYM01043238.1_8907:8934	+	GTAGCTCATATGGTAGGTCGTTAGTTTT		10	0	0	0	4	0	0	0
Oe_mir_72	FKYM01030230.1_110:140	-	CGACGTCGCGAGAAGTCCACTGAACCTTATC		158	0	0	0	64	0	0	0
Oe_mir_73	FKYM01022360.1_15591:15619	-		CCCCTGGGAAGTCCCTGTGTTGACCCCT	3	0	0	0	2	0	0	0
Oe_mir_74	FKYM01020931.1_11849:11879	+		CATTCGGCGGTGAATTCGTTCCGGGCCCTTG	4	0	0	0	2	0	0	0
Oe_mir_75	FKYM01056150.1_3689:3717	+	AAAGCTGGGTCATGGACGTCACAATGTTT		3	0	0	0	2	0	0	0
Oe_mir_76	FKYM01041963.1_1877:1905	+		AATCAACCTAGTACGAGAGGAACCGTTG	1	0	0	0	1	0	0	0
Oe_mir_77	FKYM01041963.1_1918:1948	+	GGTCATCGCGCTTGGTTGAAAAGCCAGTGGC		3	0	0	0	2	0	0	0
Oe_mir_78	FKYM01041785.1_40380:40407	-		AGTTTAGGTGAGGTATATTCTGAGCCA	3	18	0	5	2	3	0	2
Oe_mir_79	FKYM01030229.1_6373:6403	-		CGGGACATGCTCGGGCTCGGCTCGTCGCTTG	1	0	0	0	1	0	0	0
Oe_mir_80	FKYM01030229.1_5912:5941	-	TGTGAACGACGTTCTTCTCACCCCGGACC		3	0	0	0	2	0	0	0
Oe_mir_81	FKYM01022358.1_16017:16045	-	CCCCCTGGGAAGTCCCTCGTGTGACCCAC		3	0	0	0	2	0	0	0
Oe_mir_82	FKYM01022358.1_37524:37552	-	CCCTGGGAAGTCCCTCGTGTGACCCCTT		37	0	0	0	15	0	0	0
Oe_mir_83	FKYM01011318.1_22461:22490	-	CCTACATCGTATTGGACGTAGTGGGCGATT		3	0	0	0	2	0	0	0
Oe_mir_84	FKYM01010646.1_44282:44313	-	TGTGAAGGCATCGGACCAGGCTTCATCTCT		4	0	0	0	2	0	0	0
Oe_mir_85	FKYM01000663.1_3464:3491	-	CTTCTCTAGCTGCTTGGCCTGTAGTA		32	0	0	0	13	0	0	0
Oe_mir_86	FKYM01053456.1_604:634	+		CTAGTACGAGAGGACCGGGAAGGACGCACCT	4	0	0	0	2	0	0	0
Oe_mir_87	FKYM01045973.1_27091:27119	-	ATGACTTGAGAGGTGTAGGATAAGTGGGA		4	0	0	0	2	0	0	0
Oe_mir_88	FKYM01039660.1_42012:42040	-	AGTAGAGTTGAATAGCGGGAAGATCCAAT		4	0	0	0	2	0	0	0
Oe_mir_89	FKYM01030048.1_65397:65424	-		CAAGACCGTCAATTTTGATACCGTTGG	17	0	0	0	7	0	0	0
Oe_mir_90	FKYM01022360.1_4010:4038	-		CCCCTGGGATGTCTCGTGTGACCCCT	3	0	0	0	2	0	0	0
Oe_mir_91	FKYM01020701.1_5499:5526	+		AAATGATGAGTTTGGCACCTCTGAGTA	3	10	4	12	2	2	2	3
Oe_mir_92	FKYM01010646.1_44284:44313	-	TGTGAAGGCATCGGACCAGGCTTCATCTCT		6	0	0	0	3	0	0	0
Oe_mir_93	FKYM01002117.1_2497:2526	+		GGTGAAGTGTTCGGATCGCGGCGACGTGGG	6	0	0	0	3	0	0	0

Oe_mir_94	FKYM01057936.1_3228:3256	-		CTCCATCTTCTCTTTAATCCATTCTTCC	0	3	0	0	0	1	0	0
Oe_mir_95	FKYM01057852.1_1548:1577	-		CCAGTGTTAACTCCAGGAATCGAACTCTCC	0	3	0	0	0	1	0	0
Oe_mir_96	FKYM01057359.1_22840:22867	+		GGTGGAAATCGAGGTGGGGAGGTAATGG	0	3	0	0	0	1	0	0
Oe_mir_97	FKYM01057189.1_7878:7906	-		CTCCTCGGGCAGCCGATCGTAAAGTATTA	0	3	0	0	0	1	0	0
Oe_mir_98	FKYM01057147.1_46232:46259	+		TTTCAGAGCTTGGGGTCTTAGTTGGAG	0	3	0	0	0	1	0	0
Oe_mir_99	FKYM01057017.1_21057:21086	+	TGATGTCAAAGATTGGGATGGAATTCCTG		0	3	0	0	0	1	0	0
Oe_mir_100	FKYM01056678.1_69919:69947	+	GAGACCTCAGCCTACTAAGTATGCA		0	4	0	0	0	1	0	0
Oe_mir_101	FKYM01056600.1_2135:2164	+	TGATTTGAAGATATGGCAGAGGAAAACGTT		0	7	0	0	0	1	0	0
Oe_mir_102	FKYM01056569.1_38446:38475	-	AATTGTCGCTCCGATTCTTTTTTTTTTTT		0	3	0	0	0	1	0	0
Oe_mir_103	FKYM01056501.1_3277:3305	-		CTTGACTGTGGTCCCTGTTGTGCTGTAAG	0	3	0	0	0	1	0	0
Oe_mir_104	FKYM01055520.1_16794:16823	+	TAGTGTAGACAAGGTGCTTCGGCAGGACGG		0	4	0	0	0	1	0	0
Oe_mir_105	FKYM01055512.1_46275:46302	-		TCTGGAGTTATAGTAAAGGTTACACCTG	0	3	0	0	0	1	0	0
Oe_mir_106	FKYM01055253.1_22607:22636	+		CGGGTTCGAATCTTTGGTTGGCACTCCTT	0	4	0	0	0	1	0	0
Oe_mir_107	FKYM01054560.1_1592:1619	-	CTCGTGACCAAAGGCCTCCGCGTTCACC		0	4	0	0	0	1	0	0
Oe_mir_108	FKYM01053416.1_1717:1746	-		CATTGGTCTAGTGGTATGATTCGCTTT	0	17	0	0	0	3	0	0
Oe_mir_109	FKYM01053369.1_3095:3126	-	CCTGCCACACCTCCACGCGCACTCTGGCCCT		0	3	0	0	0	1	0	0
Oe_mir_110	FKYM01053337.1_30324:30354	-	CCGGACATGGCCTGCACACGGTGCCAGAAT		0	3	0	0	0	1	0	0
Oe_mir_111	FKYM01053185.1_34631:34658	-		CTTGGACCGTTGGATTGACCCGCTGATC	0	19	0	0	0	3	0	0
Oe_mir_112	FKYM01053020.1_54191:54219	+		TTCAGGCCAAGCTCGTCGTCTTACGGCG	0	3	0	0	0	1	0	0
Oe_mir_113	FKYM01052486.1_51784:51811	-		AGACCGGAATTTGAAGCAGAGATGGAGA	0	3	0	0	0	1	0	0
Oe_mir_114	FKYM01052372.1_12262:12292	-		CTTGAGGATGGTCGCACGTTGGCGGATTATA	0	3	0	0	0	1	0	0
Oe_mir_115	FKYM01051749.1_34702:34732	-	CTACTCGACCCCTTCTCTGAATTTAACAGT		0	4	0	0	0	1	0	0
Oe_mir_116	FKYM01051150.1_96483:96510	-		TTATACCTATGAATTCATTGGACCCAG	0	4	0	0	0	1	0	0
Oe_mir_117	FKYM01050899.1_46311:46340	+		GCACACTGTGCGGCTCCATCGGCATGGCAA	0	5	0	0	0	1	0	0
Oe_mir_118	FKYM01050454.1_146736:146763	-		CTTGGCGTGTAGCTGAAGCTCTTGAATA	0	3	0	3	0	1	0	1
Oe_mir_119	FKYM01050453.1_4823:4854	+		TGGGCTGCAGAATGAAATGTTGGAGATTCGA	0	6	0	0	0	1	0	0
Oe_mir_120	FKYM01050353.1_28610:28638	+	ACAGTGGATTGTTGTACTIONACTGGCGAAT		0	3	0	0	0	1	0	0
Oe_mir_121	FKYM01049968.1_32946:32974	+		CAGGTGTGGTCATTGCAAAAAGCCTTGCTC	0	3	0	0	0	1	0	0
Oe_mir_122	FKYM01047776.1_5933:5961	+	ATGTTATCTCACGTCTTGTGGCAGCCGC		0	3	0	0	0	1	0	0
Oe_mir_123	FKYM01047533.1_1444:1472	-	CGACAAGGACTTAATGGAAGGAAAGTTGG		0	5	0	0	0	1	0	0
Oe_mir_124	FKYM01047078.1_645:673	+		TTCTCAATCGACAAAAGTGAAGATAT	0	4	0	0	0	1	0	0
Oe_mir_125	FKYM01047065.1_42827:42857	-	CTCAGATGGATGACACGGACATCGTGA AAA		0	3	0	0	0	1	0	0

Oe_mir_126	FKYM01046860.1_475:502	+	ATGGGCTAATTCATAGATCTTTGGGCTT	0	6	0	0	0	1	0	0
Oe_mir_127	FKYM01046529.1_40911:40940	-	TTTAACCTTGGCTGGTGTCTCGACCATGA	0	3	0	0	0	1	0	0
Oe_mir_128	FKYM01046256.1_65825:65854	-	CCTAAAGAGTAACGGAGGTGTGCGATGGCA	0	5	0	0	0	1	0	0
Oe_mir_129	FKYM01046010.1_24247:24276	-	CTTGAAGAGTTATAGAATGTTCTGAGGCCG	0	132	123	51	0	19	40	13
Oe_mir_130	FKYM01045508.1_19849:19877	+	GTTCACACACTCGACACTTGGCCGCCCT	0	4	0	0	0	1	0	0
Oe_mir_131	FKYM01045431.1_52605:52632	+	GCATCGTGTTTGTAGTGTCTTCCCCTC	0	3	0	0	0	1	0	0
Oe_mir_132	FKYM01045414.1_40636:40665	-	AACATCCTCGGGCCATTGACTAGCTTAAAA	0	3	0	0	0	1	0	0
Oe_mir_133	FKYM01045316.1_9274:9303	+	ACCGACTGCTTGAAATTCCTCCAGTGGCCT	0	3	0	0	0	1	0	0
Oe_mir_134	FKYM01045207.1_84621:84650	+	GTTGACAATTAGACACATGAAGGGGGGCC	0	3	0	0	0	1	0	0
Oe_mir_135	FKYM01045043.1_35421:35449	-	CAATTTGGCGTCTGGATATCTGAATTCTT	0	3	0	0	0	1	0	0
Oe_mir_136	FKYM01044424.1_68296:68327	+	ACCAAAACTCTACCCTTCGGCCACCTTGGCTA	0	4	0	0	0	1	0	0
Oe_mir_137	FKYM01043974.1_52593:52621	-	CATGTGCTCTCCACCCTGGCTCCTGGGAG	0	3	0	0	0	1	0	0
Oe_mir_138	FKYM01043922.1_628:658	+	TTGTGAATGACATCTGGGGTTCATCCCGCA	0	13	0	0	0	2	0	0
Oe_mir_139	FKYM01043922.1_1250:1279	+	GGTGGAAATCGAGGTGGGGAGGTAATGGTG	0	3	0	0	0	1	0	0
Oe_mir_140	FKYM01043625.1_36768:36798	-	CATCTGATGTTTCTGTACGCTGTTTAAGCCG	0	3	0	0	0	1	0	0
Oe_mir_141	FKYM01043549.1_30443:30470	+	TGGAGAGTGCTGTTCGGGTAGCCGGACA	0	4	0	0	0	1	0	0
Oe_mir_142	FKYM01043441.1_63848:63875	+	GGTTATGAAGTAGTAACACTTGACGAAG	0	3	0	0	0	1	0	0
Oe_mir_143	FKYM01043315.1_12502:12530	+	AAAGTTGGTTTTGGTTTTCTTTGTTTAGT	0	3	0	0	0	1	0	0
Oe_mir_144	FKYM01043295.1_24797:24828	-	CATTGACTAATTAGTGATCCCTGGGCTACC	0	3	4	0	0	1	2	0
Oe_mir_145	FKYM01043227.1_12498:12526	+	TCGTTTCATTAACCTGTGCTTGAAAAAT	0	4	0	0	0	1	0	0
Oe_mir_146	FKYM01042830.1_16146:16175	+	GCTTTATAGATCTCCATGATATAGTCGTTT	0	3	0	0	0	1	0	0
Oe_mir_147	FKYM01042502.1_67650:67677	-	TTTGTGGTAGGGCATTCTTCAAGCGTT	0	4	0	0	0	1	0	0
Oe_mir_148	FKYM01042085.1_1948:1978	+	AAGGAGAGAAGTCGATGGGTACCATGCTTTT	0	3	0	0	0	1	0	0
Oe_mir_149	FKYM01042061.1_7863:7891	+	TGTCATCACCGAGCTGGGCCGTCGAGCGA	0	3	0	0	0	1	0	0
Oe_mir_150	FKYM01042027.1_9590:9619	-	TCGAACCGAACCGACCGAACCGAACCCGAC	0	3	0	0	0	1	0	0
Oe_mir_151	FKYM01041892.1_581:608	-	CGCCGATAAAGTATATATTTAAGGTAC	0	5	0	0	0	1	0	0
Oe_mir_152	FKYM01041563.1_61699:61729	-	CCCTTGAAAATCCGAAGGATCGAGTACCCTC	0	5	0	0	0	1	0	0
Oe_mir_153	FKYM01041310.1_64778:64809	+	AGAGTTTCTTAGAGCCACTGCCCTTGGCGT	0	3	0	0	0	1	0	0
Oe_mir_154	FKYM01041219.1_14158:14186	-	CCAGGGCGGCCACCTCGAAATCTTCCGGT	0	3	0	0	0	1	0	0
Oe_mir_155	FKYM01041150.1_361:392	-	CTTTGGTTTATTGACTCCACCACCTCTCTCA	0	13	0	0	0	2	0	0
Oe_mir_156	FKYM01041092.1_51841:51868	+	ACGCCTACTCTTTCGTTCCGGGATTATGA	0	3	0	0	0	1	0	0
Oe_mir_157	FKYM01040974.1_60508:60538	+	TTGGCTCGACTCGGAGATGAACTACACTTTT	0	3	0	0	0	1	0	0

Oe_mir_158	FKYM01040944.1_21036:21067	+		GGTCGTAGGTTCGAATCCTACTGGGGAGATT	0	10	0	0	0	2	0	0
Oe_mir_159	FKYM01040758.1_3058:3085	-	CCCGTGGTGAAGCAGATTTCTGGGCC		0	3	0	0	0	1	0	0
Oe_mir_160	FKYM01040472.1_43462:43492	-		CTGAACTTTGCCGAAATTCCTAGAGAGGTC	0	12	0	0	0	2	0	0
Oe_mir_161	FKYM01039660.1_42762:42789	-		CTCGTGACTCAATTGGAGAAGTATGATGCA	0	3	0	0	0	1	0	0
Oe_mir_162	FKYM01039644.1_34763:34794	+		ATGTGCTCTCCACCCTGGCTCTGGGAGGCAC	0	7	0	0	0	1	0	0
Oe_mir_163	FKYM01039644.1_33671:33700	-		GAGTTGACACCTCGGATGACACTCTACACA	0	3	0	0	0	1	0	0
Oe_mir_164	FKYM01038462.1_1224:1254	+	TGGTTGATCCTGCCAGTAGTCATATGCTTGT		0	####	0	0	0	249	0	0
Oe_mir_165	FKYM01037829.1_36200:36228	+		AGAAGAAAGATGCAGGATTTGATGGGCAA	0	3	0	0	0	1	0	0
Oe_mir_166	FKYM01037652.1_32745:32772	+	GCCCTGCTGCTCGAGACGAAGAGCTCAA		0	3	0	0	0	1	0	0
Oe_mir_167	FKYM01037411.1_2451:2481	+	AGACTATCTAGTTGTTGACGCGCTCCTTAT		0	4	0	0	0	1	0	0
Oe_mir_168	FKYM01037342.1_7411:7442	+	AAAGGAGAGGAGGATGAGGACTTCAAAATTTT		0	3	3	0	0	1	1	0
Oe_mir_169	FKYM01036609.1_19878:19908	+		TTGCATAAACTCTCACATCTCCAAGGACTT	0	3	0	0	0	1	0	0
Oe_mir_170	FKYM01036575.1_61827:61855	-	CCTCTTTGAAATGGGCTAGGGCTAAAACA		0	3	0	0	0	1	0	0
Oe_mir_171	FKYM01036452.1_35378:35406	-	CTAATGAGAGCCACTGATGTGATGATTGC		0	4	0	0	0	1	0	0
Oe_mir_172	FKYM01036277.1_2975:3005	-	CACCCCATGTGAGAGGCCCTGTACAGACCC		0	3	0	0	0	1	0	0
Oe_mir_173	FKYM01036130.1_1873:1901	+	GACTTGGGATCATGTCTTCTTAGGACTAT		0	5	0	0	0	1	0	0
Oe_mir_174	FKYM01035677.1_10375:10403	+		CTGTGATGAATTTTGAATCAGAAGATCA	0	7	0	0	0	1	0	0
Oe_mir_175	FKYM01035436.1_18874:188773	+		GTGGCCAAGTGGAAGGCAACGGGTTTTGG	0	11	0	0	0	2	0	0
Oe_mir_176	FKYM01035104.1_61715:61743	-	CACTGTACAATCGGACATTGAAATTTGCG		0	3	0	0	0	1	0	0
Oe_mir_177	FKYM01035061.1_32346:32376	-		TTGTTGGTCTTCAACGAGGAATTCCTAGTAA	0	14	0	0	0	2	0	0
Oe_mir_178	FKYM01035061.1_32122:32150	-	TGGAAGGATCATTGTCGAAACCTACAAAA		0	4	0	0	0	1	0	0
Oe_mir_179	FKYM01035043.1_111590:111617	+	AATGGGGTGCAGGAGGAATTGGTGAAT		0	3	0	0	0	1	0	0
Oe_mir_180	FKYM01034952.1_10057:10085	+		GTTGTTTGTGGATCTGAAATTTTGAAT	0	4	0	0	0	1	0	0
Oe_mir_181	FKYM01034511.1_156768:156798	-		CCCTTCGAGCAGGATATGCTATCTTTCGCTC	0	8	0	5	0	1	0	2
Oe_mir_182	FKYM01034152.1_1289:1320	+		CACCTTGGCTGCCGACCGACATACCACTTTG	0	3	0	0	0	1	0	0
Oe_mir_183	FKYM01034081.1_29017:29045	+	GAAGATTCTGTGACAACATCGTAGGATAC		0	3	0	0	0	1	0	0
Oe_mir_184	FKYM01033814.1_6356:6387	+	TTTAGATCTGAACGGACTGGTAAATTTGGGCC		0	3	0	0	0	1	0	0
Oe_mir_185	FKYM01033808.1_27257:27284	-	TATGTTGTGGTGGCGTTAAGAATTCATT		0	4	0	0	0	1	0	0
Oe_mir_186	FKYM01033761.1_269336:269364	+	ACCCAACTACTCCAATCAACCCGACTCG		0	3	0	0	0	1	0	0
Oe_mir_187	FKYM01033663.1_137754:137784	-	ACCTTCGCCTTTTACCCTGGAGATGTGCC		0	3	0	0	0	1	0	0
Oe_mir_188	FKYM01033662.1_5592:5620	+		TCGTGAGGACCGTATTGGTGTTCGAAGG	0	3	0	0	0	1	0	0
Oe_mir_189	FKYM01032959.1_25750:25777	-	TATGATGATTTGTGATGGGAATCTCTC		0	9	0	0	0	1	0	0

Oe_mir_190	FKYM01032593.1_39528:39558	+		AGAATGTTTTGGCTGCTGTAACCTTGATTT	0	3	0	0	0	1	0	0
Oe_mir_191	FKYM01032455.1_173952:173983	-		CTGACATTGATTCCACTGATTGTTGCTTCCAG	0	9	0	0	0	1	0	0
Oe_mir_192	FKYM01032398.1_38958:38986	-	CTTTGAGTCTTGGGAAACCTTAGTGGGG		0	3	0	0	0	1	0	0
Oe_mir_193	FKYM01032261.1_34412:34439	+	GCTTGGGCTAAGATATATGGAACCTCGA		0	3	0	0	0	1	0	0
Oe_mir_194	FKYM01032181.1_16889:16920	-	CTGAACTGCTGGCTGAGAAGAGATGTGTAGCT		0	3	0	0	0	1	0	0
Oe_mir_195	FKYM01032104.1_25851:25882	+	ACTGGATCAAGGTCTGGATTATGTCGCCACAT		0	6	0	3	0	1	0	1
Oe_mir_196	FKYM01031956.1_33577:33607	+		AAGATGGGTACACTCAAGATGGGACTGTAGA	0	3	0	0	0	1	0	0
Oe_mir_197	FKYM01031715.1_553:583	-	CCCCTGTTGGGAGTACGTTTGCAAGAATGAA		0	3	0	0	0	1	0	0
Oe_mir_198	FKYM01031697.1_32005:32032	-		CTGGCATACTTTGATCATGGAGGCATCC	0	4	0	0	0	1	0	0
Oe_mir_199	FKYM01031615.1_24175:24202	+		CTTAACCTTTGAAAATTAGAGTTTGACTC	0	3	0	0	0	1	0	0
Oe_mir_200	FKYM01031548.1_41101:41128	-	CTCTTCTCCCAAATCTCTCTGCGTGAA		0	3	0	0	0	1	0	0
Oe_mir_201	FKYM01031303.1_105321:105351	+		AATCAGTACATGAACCCCGTCCTTAGAAA	0	3	0	0	0	1	0	0
Oe_mir_202	FKYM01030905.1_721:748	+		GGCTCATACAGAACCGTTGAAGATATAT	0	4	0	0	0	1	0	0
Oe_mir_203	FKYM01030870.1_51900:51928	+		CGTTGTTGGAGTTGGATCTCAGTGGCAAT	0	7	4	0	0	1	2	0
Oe_mir_204	FKYM01030824.1_348:375	-	CGCCTCCGCTTATTCTGTATCGGTACAC		0	4	0	0	0	1	0	0
Oe_mir_205	FKYM01030424.1_7122:7152	+	AGAAAAAGAAGAAGAAGAGAAAAAGAAAT		0	3	0	0	0	1	0	0
Oe_mir_206	FKYM01030327.1_64018:64047	+	AGGGCAGAAGAGAATGATTGAACTTGGTAT		0	3	0	0	0	1	0	0
Oe_mir_207	FKYM01030115.1_2788:2817	-		CTGCCATTGAGATAAGAGATCAAGATTGT	0	3	0	0	0	1	0	0
Oe_mir_208	FKYM01029967.1_18286:18315	+		ATACCTTTTGTCCACTTCCAGTTCCTATG	0	5	0	0	0	1	0	0
Oe_mir_209	FKYM01028810.1_612:642	-		CTAGGAAGAGCAGCATACTTGGAGGATTGGC	0	3	0	0	0	1	0	0
Oe_mir_210	FKYM01028100.1_2654:2684	+	AAATCAAACAGGGCTTCTGTAGATTGGGCT		0	3	0	0	0	1	0	0
Oe_mir_211	FKYM01027885.1_29180:29209	-	ACAATGGAATCAGTTTCTTAGGAAGGCCAC		0	7	0	0	0	1	0	0
Oe_mir_212	FKYM01027527.1_7775:7802	+	AACGTTACTGTTATTATCATGAACAAAT		0	4	0	0	0	1	0	0
Oe_mir_213	FKYM01027278.1_291:318	-	TGTTAAATCTATTTCCTGCACATTGGTT		0	3	0	0	0	1	0	0
Oe_mir_214	FKYM01026684.1_21524:21551	-	GTAAGCCATGTCTTTCATTGAGGTGCGC		0	4	0	0	0	1	0	0
Oe_mir_215	FKYM01026603.1_7850:7877	-		TGGATGTGAGAGATAAAAAATAAAGGGTT	0	3	0	0	0	1	0	0
Oe_mir_216	FKYM01026590.1_3110:3137	+	CATTTACTGCTCTCATTGTACACTAATG		0	4	0	0	0	1	0	0
Oe_mir_217	FKYM01026399.1_2058:2089	+	TAAGATTAGAGCGCATGTCTAAGTTGTTCCGGG		0	5	0	0	0	1	0	0
Oe_mir_218	FKYM01026253.1_22880:22909	+	GGTGTCCTGTTTCAGTCAAAGCCGGTCTGCA		0	3	0	0	0	1	0	0
Oe_mir_219	FKYM01026244.1_11719:11748	+	AGTGGCAAGAAAAGGACAGTCGAGAAGAGA		0	3	0	0	0	1	0	0
Oe_mir_220	FKYM01026217.1_6151:6179	+		TTGCAGTTCGTGGTGTGTTACTCATT	0	4	0	0	0	1	0	0
Oe_mir_221	FKYM01026167.1_1447:1475	-	TATCGACACCTAAGATTGATGACCAGACC		0	3	0	0	0	1	0	0

Oe_mir_222	FKYM01025872.1_11227:11257	+		TTGGTCATCTGGGCGTTTGTAGATCAAGAAT	0	4	0	0	0	1	0	0
Oe_mir_223	FKYM01025820.1_24211:24239	-	CAGTATTAGACGCCCCAGAAACTGAAGGA		0	4	0	3	0	1	0	1
Oe_mir_224	FKYM01025779.1_287:315	-		CGTTGGAATGCAAATGGTCGTCGAACCTT	0	3	0	0	0	1	0	0
Oe_mir_225	FKYM01025563.1_59213:59240	+	CTTTGTCTTTGGGGCTTACAATTCACCT		0	14	0	0	0	2	0	0
Oe_mir_226	FKYM01025129.1_10921:10952	-		GGGTATCTTGAAAGTCTCGGGTTCGAACCTTT	0	3	0	0	0	1	0	0
Oe_mir_227	FKYM01025050.1_77572:77602	+	CCTCTGGCTGTCGCGTGCTTGTGAACCCTTG		0	4	0	0	0	1	0	0
Oe_mir_228	FKYM01024992.1_111314:111342	+		GGTAGGCTTTAGTCATTGGCGCTCGAGTG	0	4	0	0	0	1	0	0
Oe_mir_229	FKYM01024767.1_64355:64384	-		CACTCACCACCACCTTCCAATTCATTGCA	0	3	0	0	0	1	0	0
Oe_mir_230	FKYM01023731.1_32271:32298	+	CATGGGCCAGGCTTGATGCGAACCGGCC		0	4	0	0	0	1	0	0
Oe_mir_231	FKYM01023446.1_7389:7418	+		TTTGCAGGATGGGGAACAAATGAGAAATTG	0	3	0	0	0	1	0	0
Oe_mir_232	FKYM01023343.1_42295:42322	-		GGGGCTGAAGGATCATGAAGACTTGATT	0	3	0	0	0	1	0	0
Oe_mir_233	FKYM01023310.1_82754:82781	+		CTTTCATGGTGTAGTTCACATTGCTCCA	0	4	0	0	0	1	0	0
Oe_mir_234	FKYM01022965.1_14715:14742	-	ATTATTACTGCTGCTACTACTACTTGAC		0	3	0	0	0	1	0	0
Oe_mir_235	FKYM01022919.1_11172:11200	+		ATTGGCGGAGGCTGATGATAATGGGAGCA	0	3	0	0	0	1	0	0
Oe_mir_236	FKYM01022889.1_105454:105483	-	GGAGAGACTGAGTGCTCAGTTATCATGTTT		0	4	5	0	0	1	2	0
Oe_mir_237	FKYM01022630.1_148806:148834	-		CGAGTGGGGATTGGTACTGATGGTTTCAT	0	3	0	0	0	1	0	0
Oe_mir_238	FKYM01022360.1_18270:18298	-	GCCCTGGGAAGTCCCTCGTGTGCACCCCT		0	149	0	0	0	21	0	0
Oe_mir_239	FKYM01022360.1_21839:21868	-	GTAAAGCGTGCTTGGGCGAGAGTAGTACTA		0	6	0	0	0	1	0	0
Oe_mir_240	FKYM01022359.1_6030:6057	-		CTTGGGCGAGAGTAGTACTAGGATGGAT	0	6	0	0	0	1	0	0
Oe_mir_241	FKYM01022358.1_21056:21086	+		TAACTTCGGAGTTCTGATGGGATCCGAGGC	0	3	0	0	0	1	0	0
Oe_mir_242	FKYM01022358.1_10020:10047	-	CCCCTGGGAAGTCCCTCGTGTGCACCCC		0	###	0	0	0	406	0	0
Oe_mir_243	FKYM01022358.1_10678:10706	-		CCCCTGGGAAGTCCCTCGTGTGCACCCCT	0	33	0	0	0	5	0	0
Oe_mir_244	FKYM01022358.1_22439:22467	-		CCCGTGGGAAGTCCCTCGTGTGCACCCCT	0	17	0	0	0	3	0	0
Oe_mir_245	FKYM01022012.1_27180:27211	+	GACTAGTCTTGTGGTGTATTGTCTTTG		0	3	0	0	0	1	0	0
Oe_mir_246	FKYM01021721.1_37399:37428	+		TACTGAGACGCAGCACTGGAGATTTATGAT	0	4	0	0	0	1	0	0
Oe_mir_247	FKYM01021364.1_134468:134498	+		CTCCATTCACAGACTTCCCTGTTTTGTACCAA	0	3	0	0	0	1	0	0
Oe_mir_248	FKYM01021342.1_14557:14585	-		CTTTGTCGTTGATTTTGTCTGTTGCAG	0	3	0	0	0	1	0	0
Oe_mir_249	FKYM01020992.1_55162:55190	-		TAGGCTGGTTCGGATTGGTTATACATAGT	0	4	0	0	0	1	0	0
Oe_mir_250	FKYM01020931.1_11967:11998	-		ATTGTACACCCCGGTCACACTATGGGAGC	0	15	0	0	0	2	0	0
Oe_mir_251	FKYM01020887.1_50338:50365	-		TTCCCTTGCTTGTGATTGCAGTATCTGT	0	3	0	0	0	1	0	0
Oe_mir_252	FKYM01020836.1_54001:54029	-		CTGGATTTAGAACAAAACAAGACAGCC	0	4	0	0	0	1	0	0
Oe_mir_253	FKYM01020806.1_47918:47946	+		GCTACATTTTACATTTTCTTGTGGATA	0	5	0	0	0	1	0	0



Oe_mir_254	FKYM01020739.1_18088:18118	-	CTGGGTCATGAAATTGAGAATATTGTAGAGC	0	3	0	0	0	1	0	0
Oe_mir_255	FKYM01020737.1_19863:19892	+	AACATCAGCTGCTGCTTCTCCCATGGGGGG	0	3	0	0	0	1	0	0
Oe_mir_256	FKYM01020655.1_53523:53550	-	CAAGTGGAATTGTGTTGACAGTTGCTGT	0	7	3	0	0	1	1	0
Oe_mir_257	FKYM01020576.1_26883:26914	-	CTGCCTGCCAATGTTGCTGATCTCCTGTAAGG	0	4	0	0	0	1	0	0
Oe_mir_258	FKYM01020304.1_32011:32038	-	CCTGGACCTGTTTGATAAECTTGATGTAG	0	3	0	0	0	1	0	0
Oe_mir_259	FKYM01020216.1_29780:29808	-	CCAGTGGTCTAGTGGTAGAATAGTACCCT	0	74	0	0	0	10	0	0
Oe_mir_260	FKYM01020215.1_15652:15679	-	CGTGGCCAAGTGGTAAGGCAACGGGTTT	0	72	0	15	0	10	0	4
Oe_mir_261	FKYM01019908.1_177537:177565	-	CTGTCTGGATGGATATATGAATATAGCGA	0	3	0	0	0	1	0	0
Oe_mir_262	FKYM01019888.1_34696:34724	+	TAGGCATCGATGGATGGGACTGATTGAA	0	5	0	0	0	1	0	0
Oe_mir_263	FKYM01018286.1_1424:1453	-	CATTTGTGGAGAACAGAAGACTTGCTTATT	0	3	0	0	0	1	0	0
Oe_mir_264	FKYM01018039.1_6767:6795	-	CTGTGAGTCGGGACTTGCTAAGGTCAATC	0	3	0	0	0	1	0	0
Oe_mir_265	FKYM01017651.1_96711:96738	+	GAGAAGGAATTAGCTGCTGTTAAGATTA	0	3	0	0	0	1	0	0
Oe_mir_266	FKYM01017270.1_1650:1677	+	CAACAGATCGTCGCATTTTAGTCCTCT	0	3	0	0	0	1	0	0
Oe_mir_267	FKYM01016520.1_1151:1179	-	CAATTACCAGACTCTGAGAGCCCGGTATT	0	4	0	0	0	1	0	0
Oe_mir_268	FKYM01016251.1_16190:16220	+	AGGGAATAAGAGCTGTCTGTACACTGCGCCT	0	3	0	0	0	1	0	0
Oe_mir_269	FKYM01016140.1_893:921	-	ATGAGAGAGAAAAATGAAATGTAATAAAA	0	3	0	0	0	1	0	0
Oe_mir_270	FKYM01016107.1_50797:50825	-	AGGGTATGCAAAATGAACACTCGGCGCCC	0	3	0	0	0	1	0	0
Oe_mir_271	FKYM01015889.1_3257:3286	-	CAGGTTGTTGGAGGCAGCAGATGAGAGGAT	0	3	0	0	0	1	0	0
Oe_mir_272	FKYM01015691.1_7881:7909	+	TAGTGTGGACGTTGGAGATTAGTAAGTC	0	6	0	0	0	1	0	0
Oe_mir_273	FKYM01015362.1_13420:13447	-	TTTTGTCTCTGATGTCTTTTGATGCAG	0	3	0	0	0	1	0	0
Oe_mir_274	FKYM01015154.1_43247:43275	-	TCTGTACCGATTGAACTTTGCTACTTGTA	0	4	0	0	0	1	0	0
Oe_mir_275	FKYM01014890.1_29876:29905	-	GGTGAAGCGGAGGAAGTTGGACTAAGGAG	0	4	0	0	0	1	0	0
Oe_mir_276	FKYM01014814.1_23059:23086	+	TCTGAACTGAGAACACAGGCAGATTAT	0	3	0	0	0	1	0	0
Oe_mir_277	FKYM01014745.1_573:601	+	TCTCTAGACTGTGTCGGAATTGGCTGAAA	0	3	0	0	0	1	0	0
Oe_mir_278	FKYM01014521.1_2636:2664	+	GTATTGATGTGTAGCTTGGAGGAGTTGTA	0	4	0	0	0	1	0	0
Oe_mir_279	FKYM01014450.1_36446:36473	+	CTCCATACTGACGAACCATCTGTTGATG	0	3	0	0	0	1	0	0
Oe_mir_280	FKYM01014275.1_38979:39006	+	ATTTGGTCTAGTGGTATGATTCTCGCTT	0	61	0	0	0	9	0	0
Oe_mir_281	FKYM01014275.1_39321:39349	-	CATTTGGTCTAGTGGTATGATTCTCGCTT	0	13	0	0	0	2	0	0
Oe_mir_282	FKYM01014226.1_36524:36552	+	TGATGGAAGTGGCGAAGGACGGTGACGGT	0	3	0	0	0	1	0	0
Oe_mir_283	FKYM01014067.1_11352:11380	+	ACCAGCGTTCGATTGCTTCACATGTCTC	0	6	0	0	0	1	0	0
Oe_mir_284	FKYM01014066.1_42739:42770	+	ATTTTTCTATCCAACCCACAACCTATAAGTA	0	3	0	0	0	1	0	0
Oe_mir_285	FKYM01013831.1_112003:112031	+	GGATTGAGCGAGGGGAGAGGACGATAAT	0	3	0	0	0	1	0	0

Oe_mir_286	FKYM01013681.1_43255:43283	+	AGAAGCTCTGACCCTAATAATCCGCAAAG		0	3	0	0	0	1	0	0
Oe_mir_287	FKYM01013460.1_48785:48813	+	GTGGTTGAAAACACTGAGAGGGGATGGCG		0	3	0	0	0	1	0	0
Oe_mir_288	FKYM01012939.1_24989:25019	+	TGGACCAAAGAGGTGGACTATAACCGCAGAGA		0	3	0	0	0	1	0	0
Oe_mir_289	FKYM01012927.1_2686:2717	-	CTGAGATTGTCGTTGGCCTTATTGATCACCTA		0	3	0	0	0	1	0	0
Oe_mir_290	FKYM01012386.1_11060:11089	-		CAGAGCAGATTCACGTACAGGTTCAAATC	0	3	0	0	0	1	0	0
Oe_mir_291	FKYM01012275.1_39013:39041	+	TCTAGTTCGCAAAGGCCTGGAAAGAAGCTG		0	3	0	0	0	1	0	0
Oe_mir_292	FKYM01012177.1_74001:74029	+	TGGAACGCCGTCGCGCTCTGGGCTTGGGG		0	5	0	0	0	1	0	0
Oe_mir_293	FKYM01011956.1_31804:31832	+	GTGGAGGATTAGATCCAGTAACTGGGGGT		0	3	0	0	0	1	0	0
Oe_mir_294	FKYM01011839.1_22220:22247	+	GAGACCGAGAAGTTGCTTTAGCTGACTC		0	3	0	0	0	1	0	0
Oe_mir_295	FKYM01011479.1_96921:96952	-	CATGTGATGATTTGCTGGATATCTAGATGGCT		0	10	4	0	0	2	2	0
Oe_mir_296	FKYM01011318.1_22461:22489	-	CTACATCGTATTGGACGTAGTGGGCGATT		0	20	0	0	0	3	0	0
Oe_mir_297	FKYM01011217.1_3441:3472	+		TTTTATGCTTGATGAGATTGTTGATGGAGGG	0	6	0	0	0	1	0	0
Oe_mir_298	FKYM01010646.1_44324:44355	-		GGCTGGCTCGAAGCAATCACAGTTTGTCTGTTA	0	9	0	2	0	1	0	1
Oe_mir_299	FKYM01010646.1_44283:44313	-	TGTGAAGGCATCGGACCAGGCTTCATTCTC		0	6	0	3	0	1	0	1
Oe_mir_300	FKYM01010420.1_5154:5184	-	CAGGAGATGATGACACTGGCGCGGAAGAGA		0	4	0	0	0	1	0	0
Oe_mir_301	FKYM01010102.1_550:579	-	CTTGTTTCTGATGATGACGAAACACATATG		0	11	0	0	0	2	0	0
Oe_mir_302	FKYM01010006.1_13689:13718	-		CATTGGGCTAGAATACTTGTGAGGGATTG	0	4	0	0	0	1	0	0
Oe_mir_303	FKYM01009893.1_35512:35542	-		CAACGCGGGCAAATCGGAATCTACTTGCCA	0	3	0	0	0	1	0	0
Oe_mir_304	FKYM01009841.1_22720:22750	-		CTACTGCTTCTAATCCATTGAAGTCAGATG	0	3	0	0	0	1	0	0
Oe_mir_305	FKYM01009823.1_3172:3201	+		ATGCCTATGAAATGTGGACCGATAAACCA	0	5	0	0	0	1	0	0
Oe_mir_306	FKYM01008166.1_3621:3651	-		TTTTGGGTTACCCGACCCACCCTCCTCTC	0	5	0	0	0	1	0	0
Oe_mir_307	FKYM01007534.1_28559:28588	-	TTTTCATTTCACTTTTCTTATCGTTCACA		0	3	0	0	0	1	0	0
Oe_mir_308	FKYM01007076.1_17287:17316	+	GGTGAGGCTGGGCGACCTGATGAGGTGGCA		0	35	15	0	0	5	5	0
Oe_mir_309	FKYM01006718.1_20647:20676	-	CTCTTGCGGAATGTTTGTAGATTCTGGAT		0	3	0	0	0	1	0	0
Oe_mir_310	FKYM01006580.1_94466:94497	-		CTTAAGAAAGCAATTCGGCTCCGGCGATAAC	0	3	0	0	0	1	0	0
Oe_mir_311	FKYM01006545.1_48432:48460	+	CTTCAGGACTGTGAGAAGTGTGATGAGT		0	6	0	0	0	1	0	0
Oe_mir_312	FKYM01006022.1_8662:8693	+		GTCGCGAAGCCCGATTTCAGATTCTGGCCG	0	3	0	0	0	1	0	0
Oe_mir_313	FKYM01005955.1_4763:4791	+		GGGGTGAACGGCCGTAAGTTCCTGGTCCC	0	3	0	0	0	1	0	0
Oe_mir_314	FKYM01005945.1_5994:6021	+	GTCAATCGAAGCCTCAGTGTCCAGAT		0	3	0	0	0	1	0	0
Oe_mir_315	FKYM01005943.1_45544:45572	-		CCAGTTCCTCCGAGGCCTCCTCATGTACT	0	3	0	0	0	1	0	0
Oe_mir_316	FKYM01005859.1_33696:33725	+		CGTCTTGATTGGAAGCACATGTCTACACTT	0	5	0	0	0	1	0	0
Oe_mir_317	FKYM01005528.1_1435:1463	+		CGCTGCGGATAAGGACAAGGCACCAAAT	0	3	0	0	0	1	0	0

Oe_mir_318	FKYM01005233.1_459:487	+	TTCAGTCGCCTCCAGTAAATCAATGGGAA	0	3	0	0	0	1	0	0
Oe_mir_319	FKYM01005150.1_42797:42825	-	CTGTCAGGGTTGGCTTACTTGCACAGCAA	0	3	0	0	0	1	0	0
Oe_mir_320	FKYM01004445.1_30721:30748	+	TTATGATCTTGAAGGCAAGACCGTTGGA	0	4	0	0	0	1	0	0
Oe_mir_321	FKYM01004346.1_45994:46022	-	CTCCATACTGCAAAGGGCTGCTGAGAAGA	0	3	0	0	0	1	0	0
Oe_mir_322	FKYM01004241.1_35582:35609	-	CAGAAGTTTTACGTGGAATGCAAAAAGA	0	6	0	0	0	1	0	0
Oe_mir_323	FKYM01004020.1_12618:12647	+	TGTGATAATGGAAGTGAATGGTGAAGGTT	0	4	0	0	0	1	0	0
Oe_mir_324	FKYM01003810.1_57390:57417	+	CGCTGAGATTCAGCCCTTTGTCGCTCA	0	4	0	4	0	1	0	1
Oe_mir_325	FKYM01003810.1_125141:125169	+	ATAGTGTACATGTAGCAAGGTCCCAACCG	0	4	0	0	0	1	0	0
Oe_mir_326	FKYM01003456.1_62278:62305	-	AACAGTAGTAATGAACAGTAACAGAAGA	0	3	0	0	0	1	0	0
Oe_mir_327	FKYM01003264.1_942:972	+	TTTGCAATGTTTGACGGATTGAGGGCCCAA	0	6	0	0	0	1	0	0
Oe_mir_328	FKYM01003182.1_22696:22723	-	CAGTGGCAACGGAATCTAATGGATCCTT	0	4	0	0	0	1	0	0
Oe_mir_329	FKYM01002918.1_16521:16552	-	TATACCCAGAGAGAGAGAGAGAGAGAGGGA	0	3	0	0	0	1	0	0
Oe_mir_330	FKYM01002789.1_15586:15616	+	CCATTGCCTGGTCCGAGCCTCGGACTTCTA	0	3	0	0	0	1	0	0
Oe_mir_331	FKYM01002677.1_29530:29557	-	CAAGACCGTGAATTTTGATACCGTTGG	0	4	0	0	0	1	0	0
Oe_mir_332	FKYM01002526.1_49982:50013	+	CTGTCATCGCTGTTGATTTATCTGGGGGATC	0	3	0	0	0	1	0	0
Oe_mir_333	FKYM01002507.1_947:976	+	TTGGAGGAATTGTGGACTAGGCCTGGGAA	0	4	0	0	0	1	0	0
Oe_mir_334	FKYM01002091.1_17328:17356	-	CCCCTGTTACCAAAACTGTCATACTGT	0	4	0	0	0	1	0	0
Oe_mir_335	FKYM01002017.1_7876:7903	+	CGTGCTGGTACGGACTGAGGTTTTAT	0	3	0	7	0	1	0	2
Oe_mir_336	FKYM01001997.1_45585:45612	+	TGGTGTTAAGAGGCAAGAAAGTGAGAT	0	3	0	0	0	1	0	0
Oe_mir_337	FKYM01001859.1_141973:142000	-	GAGGAACCTCCTGAGCTGATGATGTGA	0	3	0	0	0	1	0	0
Oe_mir_338	FKYM01001075.1_1073:1101	-	CCAAGTCAAGATCTCCCGTCGGTTATCGT	0	3	0	0	0	1	0	0
Oe_mir_339	FKYM01000829.1_14769:14800	+	TCTTCTGGATTGAGCTCATCTGAAACTAAGCT	0	4	0	0	0	1	0	0
Oe_mir_340	FKYM01000423.1_115924:115951	-	TGTTACCTGGACTGCCTGCCATCATCC	0	4	0	0	0	1	0	0
Oe_mir_341	FKYM01000307.1_2869:2899	+	CTCACTGTAGTACCATTGGAAGAATACTGG	0	18	0	0	0	3	0	0
Oe_mir_342	FKYM01000306.1_55757:55785	-	CAACAAGAATGTTGAGTACTGGCAGCAGG	0	3	0	0	0	1	0	0
Oe_mir_343	FKYM01000034.1_132491:132518	-	CGAAGTTGATATTATTCATCATAGCTG	0	3	0	0	0	1	0	0
Oe_mir_344	FKYM01056917.1_4069:4096	+	GGGTGAACCTTGATAAAGACTGATAAACT	0	0	3	0	0	0	1	0
Oe_mir_345	FKYM01056571.1_23321:23350	+	CGGTGATATGACCTCTTCTCTGAGGCAGT	0	0	4	0	0	0	2	0
Oe_mir_346	FKYM01055655.1_115505:115533	+	TCTACCCTTCGACCACCTTAGCCGTTGGA	0	0	3	0	0	0	1	0
Oe_mir_347	FKYM01055504.1_37807:37834	-	CATTGGGACATGATGTCGTCTGATCCAC	0	0	3	0	0	0	1	0
Oe_mir_348	FKYM01055053.1_33715:33742	-	TTCTTTGACATGACAATCGGCGGTCAAC	0	0	3	0	0	0	1	0
Oe_mir_349	FKYM01054928.1_4030:4057	-	ACTTGGTAGGAATGTAGTATTCTGTTTC	0	0	3	5	0	0	1	2

Oe_mir_350	FKYM01054349.1_17122:17150	-	CTCTGTAGTGGATTGAGTGTGCATCCTTT		0	0	21	0	0	0	7	0
Oe_mir_351	FKYM01053416.1_1393:1420	+		TAGTGGTATGATTCTCGCTTAGGGTGCG	0	0	6	0	0	0	2	0
Oe_mir_352	FKYM01052820.1_9440:9468	+		CGTGGTAGAAACAACCACACCTCTGACTT	0	0	3	0	0	0	1	0
Oe_mir_353	FKYM01052084.1_102596:102623	-	CGAGAATGGAGCTAAGGGTTGTGAGGTA		0	0	3	0	0	0	1	0
Oe_mir_354	FKYM01052049.1_2914:2945	+		TGGCTGATTAGGAATCCTCTTGAATTTAGGTC	0	0	4	0	0	0	2	0
Oe_mir_355	FKYM01052049.1_3096:3127	+	CGTGAGGATTTCTCTCCGTCAGTATTCTCAT		0	0	1	0	0	0	1	0
Oe_mir_356	FKYM01051315.1_42025:42054	+	GGAGGAGATGAGCCTGAGTAAATCAAAGTG		0	0	3	0	0	0	1	0
Oe_mir_357	FKYM01051081.1_22229:22258	-		CTTCAAGGAATCCCATGAAGTCCCATGATC	0	0	4	0	0	0	2	0
Oe_mir_358	FKYM01048149.1_20186:20214	-		CATTGAGAGACTCCGTGTTGTTGACATC	0	0	3	0	0	0	1	0
Oe_mir_359	FKYM01047269.1_37537:37565	-	TTGATATATTTGATTACATTAAGAGAATG		0	0	3	0	0	0	1	0
Oe_mir_360	FKYM01047076.1_60339:60370	+		ATTTTCGCGACTGTTGGAGTGTGTAGTTAT	0	0	3	0	0	0	1	0
Oe_mir_361	FKYM01046746.1_51420:51447	+		ATGGACATGAACCTGGAGATGAAGGTGA	0	0	3	0	0	0	1	0
Oe_mir_362	FKYM01046385.1_14681:14712	+		CCCAGTCTCAAGTGCAACCTGACCCTATCATC	0	0	9	7	0	0	3	2
Oe_mir_363	FKYM01046256.1_64432:64462	-	CCTTCGGACCCTTGACATTTTCATCGCCTTAC		0	0	9	0	0	0	3	0
Oe_mir_364	FKYM01044424.1_72272:72299	+		CTATGATGTTGATCTCAAATTTGTGCCA	0	0	3	0	0	0	1	0
Oe_mir_365	FKYM01042757.1_1085:1113	-	AGGATGAAAATGGTGGAGAGCAGTGAAGA		0	0	3	0	0	0	1	0
Oe_mir_366	FKYM01041975.1_2987:3014	-	GGAGGTGGTCTTAATGAAAAACAAAGCG		0	0	3	0	0	0	1	0
Oe_mir_367	FKYM01041563.1_53782:53810	-	TGAGAAGTCCACTGAACCTTATCATTTAG		0	0	3	0	0	0	1	0
Oe_mir_368	FKYM01041531.1_9292:9319	+		ACTGAACTCTGTTCTTGAATTTGTGTCT	0	0	4	0	0	0	2	0
Oe_mir_369	FKYM01041155.1_13199:13226	+	GGATACTCTGAGATAGAGCGAGAAAAAT		0	0	8	0	0	0	3	0
Oe_mir_370	FKYM01040723.1_27723:27753	-		CGGGTTCGAATCTTTGGTTGGCACTTCCTTT	0	0	4	0	0	0	2	0
Oe_mir_371	FKYM01040102.1_14957:14985	+	CTCCCCAAATTAGCCTTGCCCAAGTATCCC		0	0	3	0	0	0	1	0
Oe_mir_372	FKYM01038735.1_2416:2443	-	CTTGTTTACAAGTAGATAGTGGCTAGCT		0	0	3	0	0	0	1	0
Oe_mir_373	FKYM01035905.1_70113:70141	-		CAGTTGGTTAGAGCGTGGCTGTTAACC	0	0	5	6	0	0	2	2
Oe_mir_374	FKYM01035436.1_188744:188772	+		GTGGCCAAGTGGTAAGGCAACGGGTTTTG	0	0	5	10	0	0	2	3
Oe_mir_375	FKYM01034996.1_25393:25420	+	ATACTCTGAGGATGTTCTGCATTGGCTT		0	0	3	0	0	0	1	0
Oe_mir_376	FKYM01034384.1_70207:70234	+	TCTCTGAGCACCCGATGTCTGCTACTAT		0	0	3	0	0	0	1	0
Oe_mir_377	FKYM01034242.1_1585:1612	+	ATAGGTTTAATTTCTGTAGACTCACAT		0	0	5	0	0	0	2	0
Oe_mir_378	FKYM01034207.1_114302:114330	-	TTACTCCGAGGCATCTCCAGCTTGCTATT		0	0	3	0	0	0	1	0
Oe_mir_379	FKYM01033615.1_75822:75849	-	CTGCCACTTGTTGATTTTGTAAACCCATT		0	0	3	0	0	0	1	0
Oe_mir_380	FKYM01033607.1_64785:64814	-		AGACAAAGATTGAGCCCTAAGATTTTCAGA	0	0	3	0	0	0	1	0
Oe_mir_381	FKYM01032996.1_78923:78950	+		GAACTTGACAGTCTCCTCACAAATCGC	0	0	3	0	0	0	1	0

Oe_mir_382	FKYM01032621.1_5808:5838	-		TTGGTTAAAGGCTTGTGGATTATATAGT	0	0	3	0	0	0	1	0
Oe_mir_383	FKYM01032104.1_25853:25882	+	TGGATCAAGGTCTGGATTATGTCGCCACAT		0	0	8	0	0	0	3	0
Oe_mir_384	FKYM01031755.1_83362:83390	+		AATGGCAAGAGTAGCGGGCACCGATACT	0	0	6	0	0	0	2	0
Oe_mir_385	FKYM01030426.1_12797:12825	-		TGGAAGGGACGCATTTATTAGATAAAATGG	0	0	4	0	0	0	2	0
Oe_mir_386	FKYM01030409.1_2386:2414	+	ATAGTTACGTGCGGATGGTTGCATCTATT		0	0	3	0	0	0	1	0
Oe_mir_387	FKYM01029890.1_12727:12756	-	GATGTCGATCCTGAGCGGTTACCTGAGCAA		0	0	57	0	0	0	19	0
Oe_mir_388	FKYM01029816.1_35171:35198	-		TGATTTTATGCAAGCTGCTAAGTCCGT	0	0	3	0	0	0	1	0
Oe_mir_389	FKYM01028429.1_562:589	+		CATCGACTCCGATCTGTATTGTCTCGTT	0	0	3	0	0	0	1	0
Oe_mir_390	FKYM01027163.1_47489:47518	+		GTCCCGTGATTGGTGTTCGTGCCAAGCTGA	0	0	3	0	0	0	1	0
Oe_mir_391	FKYM01025940.1_24229:24256	+	GCGATTTCTTGGCTACGGAGTGTGTTTC		0	0	3	0	0	0	1	0
Oe_mir_392	FKYM01025563.1_59213:59241	+	CTTTGTCTTTGGGGCTTACAATCACCTA		0	0	5	22	0	0	2	6
Oe_mir_393	FKYM01025050.1_77586:77615	+	GTGCTTGTGAACCTTGTCTCTTGAAGCC		0	0	3	0	0	0	1	0
Oe_mir_394	FKYM01024844.1_50311:50339	-		TCGGTGGTGATGAAGTCGAAAACGGAGTG	0	0	3	0	0	0	1	0
Oe_mir_395	FKYM01023070.1_36911:36938	-		CTTGGACCGTTGGATTGATCCGCTGATC	0	0	3	0	0	0	1	0
Oe_mir_396	FKYM01022924.1_253:282	+		AACGTCGAGGAACATGTACTAGGGTGTATG	0	0	4	0	0	0	2	0
Oe_mir_397	FKYM01022360.1_18268:18298	-	GCCCTGGGAAGTCTCGTGTGCACCCCTTT		0	0	73	0	0	0	24	0
Oe_mir_398	FKYM01022359.1_9984:10015	-	GGGTGACCCCTGGGAAGTCCTCGTGTGCC		0	0	7	0	0	0	3	0
Oe_mir_399	FKYM01022358.1_11341:11368	-		CACCTGGGAAGTCTCGTGTGCCCCCT	0	0	4	0	0	0	2	0
Oe_mir_400	FKYM01022358.1_11011:11039	-	CCTCTGGGAAGTCTCGTGTGCACCCC		0	0	11	0	0	0	4	0
Oe_mir_401	FKYM01022014.1_1538:1568	+		TCTGTGATGGTAAATGGGTGTGGTCTGAGCC	0	0	4	0	0	0	2	0
Oe_mir_402	FKYM01021782.1_22601:22628	+	TTCCATCACGCTTTGTTGTAGGGCAGG		0	0	8	9	0	0	3	3
Oe_mir_403	FKYM01020958.1_18271:18300	+		GCATTTGGTCTAGTGGTATGATTCTCGCTT	0	0	21	0	0	0	7	0
Oe_mir_404	FKYM01019895.1_7600:7630	-		CTTTATTACGTCTGGGAGCTGGAGCTTGTGC	0	0	4	0	0	0	2	0
Oe_mir_405	FKYM01018308.1_24987:25016	+	TTTCAGCAACGGATTTCTGGCTCTCGCAT		0	0	5	0	0	0	2	0
Oe_mir_406	FKYM01017665.1_47362:47391	-	ATAAATGTTGGGCTGTTATTCTTGTGCAC		0	0	4	0	0	0	2	0
Oe_mir_407	FKYM01015766.1_8152:8179	+		TGGCAGCTGGAGAAGTGAACGCAAT	0	0	3	0	0	0	1	0
Oe_mir_408	FKYM01015060.1_2770:2799	+	GGATCTGAGAAAAGCTCCAGTCTGAAGGA		0	0	4	0	0	0	2	0
Oe_mir_409	FKYM01014142.1_11310:11340	-		CACTGCCAGAGAAGTGTGGAGCAACCCATA	0	0	3	0	0	0	1	0
Oe_mir_410	FKYM01013352.1_54315:54343	-	GCTGTCTATCATCTGAATTCGGTTTAGA		0	0	4	6	0	0	2	2
Oe_mir_411	FKYM01013088.1_48462:48490	+	GGGAGGAGCAACTGAAGGAACAGATGGCT		0	0	3	0	0	0	1	0
Oe_mir_412	FKYM01012168.1_18221:18252	-		CAACGAAGACTGCAGTGTAGCTGAGCCTATAG	0	0	3	0	0	0	1	0
Oe_mir_413	FKYM01011530.1_43070:43097	+		GTTGGTCTTCAACGAGGAATTCCTAGTA	0	0	3	0	0	0	1	0

Oe_mir_414	FKYM01010415.1_37449:37477	+		GTTTGACAAACTCGACACTTGGCGGCCCC	0	0	3	0	0	0	1	0
Oe_mir_415	FKYM01007418.1_15487:15515	-		CCGTGTTTATGGATTGGATCGTTTCTGCC	0	0	7	0	0	0	3	0
Oe_mir_416	FKYM01006167.1_127537:127566	-	CAAGAAGAGGAATGTAAAACCTTGTGCCATG		0	0	3	0	0	0	1	0
Oe_mir_417	FKYM01004872.1_28641:28669	-	CAGAGAACTTACAATAATACAACGCCGCA		0	0	3	0	0	0	1	0
Oe_mir_418	FKYM01004493.1_7151:7181	-		CTCCTCCTCTCACATGGGGTTCGGACTCCAC	0	0	3	0	0	0	1	0
Oe_mir_419	FKYM01003996.1_54815:54844	-	CTGGTTGCTCTGGTTTATTTGGTGTCTGTT		0	0	3	0	0	0	1	0
Oe_mir_420	FKYM01003711.1_24032:24059	+		CCTACCTATTGACCGCAACTTCTGTAT	0	0	21	0	0	0	7	0
Oe_mir_421	FKYM01002812.1_8506:8533	+		TTTTGGTCTTTCAAGGGTGTTCATTGTC	0	0	5	0	0	0	2	0
Oe_mir_422	FKYM01002408.1_65648:65677	+	AGTGTGAAAATCCAATGGTGTGGGCTCCTG		0	0	4	0	0	0	2	0
Oe_mir_423	FKYM01001628.1_127866:127894	+		TCACATCGACGGGAAGGTTTGGCACCTCG	0	0	3	0	0	0	1	0
Oe_mir_424	FKYM01001537.1_46654:46681	+		AGAGTAACTATTGTATGTCGTGTAGA	0	0	4	0	0	0	2	0
Oe_mir_425	FKYM01000034.1_133738:133766	-	CTCATCGGGGTTGCCATGCAGATTGGCG		0	0	3	0	0	0	1	0
Oe_mir_426	FKYM01055738.1_692:719	-		CGGTGCATCGAAACTCTATTTAGTAAG	0	0	0	4	0	0	0	1
Oe_mir_427	FKYM01055253.1_22607:22638	+		CGGGTTCGAATCTTTGGTTGGCACTTCCTTAG	0	0	0	7	0	0	0	2
Oe_mir_428	FKYM01054914.1_2650:2677	+	CTTTTCGTTGAGTTGGGTCGGCAAGGA		0	0	0	4	0	0	0	1
Oe_mir_429	FKYM01054349.1_17121:17150	-	CTCTGTAGTGGATTGAGTGTGCATCCTTTT		0	0	0	6	0	0	0	2
Oe_mir_430	FKYM01053846.1_27861:27890	+		CACCACCATTTTCAACCTATCAAACCTGC	0	0	0	3	0	0	0	1
Oe_mir_431	FKYM01053416.1_1715:1746	-		CATTTGGTCTAGTGGTATGATTTCTCGCTTTGG	0	0	0	3	0	0	0	1
Oe_mir_432	FKYM01052374.1_12957:12987	-	GGTGCTTGTTCGCTTGAATAAAAAATACAGT		0	0	0	3	0	0	0	1
Oe_mir_433	FKYM01052158.1_37789:37817	-		TCCATAAECTCAAAGGATTCGGTCGGAA	0	0	0	4	0	0	0	1
Oe_mir_434	FKYM01052107.1_7040:7067	-	CTCGGACCAGGCTTCAITCCCCCAACT		0	0	0	3	0	0	0	1
Oe_mir_435	FKYM01051969.1_19399:19427	+		TCAAGGTGATGCTGATGATAAACTGGTGC	0	0	0	3	0	0	0	1
Oe_mir_436	FKYM01051778.1_52:81	-	CAAGGCTCTTAACCTTGTGGTCTGGGGTTC		0	0	0	3	0	0	0	1
Oe_mir_437	FKYM01050294.1_1154:1182	-	CTCTGAGTTCGACGTTGAATAGCCATAGT		0	0	0	4	0	0	0	1
Oe_mir_438	FKYM01047025.1_8816:8844	+	TACTTGTGAAAACCTGAATGCAAGATGTTT		0	0	0	3	0	0	0	1
Oe_mir_439	FKYM01046256.1_64435:64464	-	CTCCTTCGGACCCTTGACATTTTCATCGCCT		0	0	0	11	0	0	0	3
Oe_mir_440	FKYM01046040.1_6951:6979	-	CAGAGGTCGTGAGTTCGAAACATGGGGCT		0	0	0	3	0	0	0	1
Oe_mir_441	FKYM01044089.1_10441:10469	+		GTGTGATTAATAATGAAAGAGAAAAA	0	0	0	3	0	0	0	1
Oe_mir_442	FKYM01043838.1_1480:1507	+		TCGTTCAATTTCCATTTCCATCGACCATT	0	0	0	4	0	0	0	1
Oe_mir_443	FKYM01042756.1_3717:3745	-		TCTTAACCACTGTCCACCTTGAATCAC	0	0	0	3	0	0	0	1
Oe_mir_444	FKYM01042675.1_28157:28185	+	CGAGGAGGGGCTCATGACGGGGTGTGG		0	0	0	4	0	0	0	1
Oe_mir_445	FKYM01042141.1_13034:13062	+		CCAACATGTCAGAAATAAGAGCTTGCTC	0	0	0	3	0	0	0	1

Oe_mir_446	FKYM01041563.1_53968:53996	-		TGTGATGGGGATAGATCATTGCAATTGTT	0	0	0	3	0	0	0	1
Oe_mir_447	FKYM01041226.1_24538:24566	+	ACCCATGGGGATTACAAGTCTGAGATGC		0	0	0	14	0	0	0	4
Oe_mir_448	FKYM01040730.1_160344:160373	-	CGCTCGGCTCGCCGTTGACTCTGGTGGTGA		0	0	0	3	0	0	0	1
Oe_mir_449	FKYM01040669.1_64843:64874	-		CGGACGAGAAAGACAGTGTGACGGTGGAGAG	0	0	0	3	0	0	0	1
Oe_mir_450	FKYM01039618.1_3599:3629	-	CTTCCAGCAATTTATAATAATCCAGACGTAC		0	0	0	3	0	0	0	1
Oe_mir_451	FKYM01038484.1_2158:2187	+		GTTGTCCCTTCTCGTTCTATTATTATCT	0	0	0	3	0	0	0	1
Oe_mir_452	FKYM01038462.1_1224:1252	+	TGGTTGATCCTGCCAGTAGTCATATGCTT		0	0	0	10	0	0	0	3
Oe_mir_453	FKYM01036119.1_83981:84012	-		CCGGGTTTGAATCTTTGGTTGGCACTTCCTTT	0	0	0	62	0	0	0	16
Oe_mir_454	FKYM01035439.1_6502:6533	-		TTGTTGTCGATGTTGTGGTAAACTGAGCCCGA	0	0	0	5	0	0	0	2
Oe_mir_455	FKYM01034081.1_28300:28331	-	CTTTGGAAATTTAGGCTCTGTGTTTGTGGG		0	0	0	3	0	0	0	1
Oe_mir_456	FKYM01033960.1_62696:62723	+	CGAACCTTGACGCGGCAGCCTCAATCA		0	0	0	3	0	0	0	1
Oe_mir_457	FKYM01033524.1_70787:70817	-	CTGAATGATCTCGGACCAGGCTTCATCCCC		0	0	0	3	0	0	0	1
Oe_mir_458	FKYM01032968.1_33117:33144	-	GTGGCACTTACTTACTTGTATAGAGATA		0	0	0	5	0	0	0	2
Oe_mir_459	FKYM01032634.1_9323:9351	+	ACTCCCTGTTGTAATAATGTTTAGATCAT		0	0	0	3	0	0	0	1
Oe_mir_460	FKYM01031741.1_33313:33343	+		TTTGGGTGCGAGAGGTCCCAGTTCGATTCT	0	0	0	3	0	0	0	1
Oe_mir_461	FKYM01031732.1_27776:27803	-		TAGACCTTGTGTGTGTCAGAATCTTAA	0	0	0	10	0	0	0	3
Oe_mir_462	FKYM01030017.1_17868:17897	+		TTGGCGAGGAAGGTGTCTCTGTTTAGGGCG	0	0	0	3	0	0	0	1
Oe_mir_463	FKYM01030017.1_18270:18299	+	TGGAACGGTATAAACCCCTGCTAGGAGCTT		0	0	0	2	0	0	0	1
Oe_mir_464	FKYM01027470.1_5567:5595	-	CTCATGGCCTAGTATTTACATCGTTAAC		0	0	0	3	0	0	0	1
Oe_mir_465	FKYM01026946.1_9147:9175	-		TACATGGAAATACTCAGTAGAATATAAAA	0	0	0	3	0	0	0	1
Oe_mir_466	FKYM01025050.1_61504:61532	+		AAGTCTCAGGATCGTATTATTGGAGCCCA	0	0	0	3	0	0	0	1
Oe_mir_467	FKYM01025050.1_77573:77604	+	CTCTGGCTGTCGCGTGCTTGGAACCCCTTGCT		0	0	0	3	0	0	0	1
Oe_mir_468	FKYM01025035.1_62024:62052	+	CCCTAATCGGCTTGTGTAGACGAGGCCG		0	0	0	3	0	0	0	1
Oe_mir_469	FKYM01024763.1_628:656	-	GGGGATGCCGAAGGCAGGGCTAGTACTG		0	0	0	16	0	0	0	4
Oe_mir_470	FKYM01023719.1_39867:39896	-		TGTTGGGGCTGACTACTTTGACAGGCCACA	0	0	0	3	0	0	0	1
Oe_mir_471	FKYM01022884.1_67927:67955	-		CCGAACGCTACGGTCCACTCATGATGCTT	0	0	0	4	0	0	0	1
Oe_mir_472	FKYM01022360.1_19589:19618	-	CCCCCTGGGAAGTCCCTCGTGTGTACCTCT		0	0	0	6	0	0	0	2
Oe_mir_473	FKYM01022359.1_5334:5362	-		CCCATGGGAAGTCCCTCGTGTGCACCCCT	0	0	0	9	0	0	0	3
Oe_mir_474	FKYM01022358.1_22800:22827	-	CTTGGGCGAGAGTAGAACTAGGATGGGT		0	0	0	3	0	0	0	1
Oe_mir_475	FKYM01021684.1_47380:47409	-	CCGACAGGAAGGATAGGAATGGCTGCCCA		0	0	0	3	0	0	0	1
Oe_mir_476	FKYM01020992.1_55159:55190	-		TAGGCTGGTTCGGATTGGTTATACATAGTTCC	0	0	0	13	0	0	0	4
Oe_mir_477	FKYM01020978.1_1376:1407	-	CTAATGAATTCCTGGGAACCTTCTATAGTAAA		0	0	0	4	0	0	0	1

Oe_mir_478	FKYM01014973.1_4534:4561	-	CGGGCCTGGGTTCACTGGAAGAAGCTAG		0	0	0	3	0	0	0	1
Oe_mir_479	FKYM01013831.1_82961:82990	+		TTCAATCCAATAGATCTGATCTGAGCCTCA	0	0	0	14	0	0	0	4
Oe_mir_480	FKYM01012340.1_10912:10941	+	GGCAGGGGCTATATTAAGAGCAGAGACCTT		0	0	0	3	0	0	0	1
Oe_mir_481	FKYM01011433.1_17511:17539	+		CCCTATTTGTTGGTCCGTTCTAGTGATA	0	0	0	16	0	0	0	4
Oe_mir_482	FKYM01008235.1_32541:32568	-		TTTTGTTGATTGAACTTTTGTATCAG	0	0	0	3	0	0	0	1
Oe_mir_483	FKYM01006869.1_3254:3284	-		CCCCGTGAAACTCTCCATGAACTTTTGAAAC	0	0	0	3	0	0	0	1
Oe_mir_484	FKYM01004084.1_58055:58083	+		AGTCGACTTTAGCGAAACACCCTTTGACA	0	0	0	4	0	0	0	1
Oe_mir_485	FKYM01003121.1_36001:36030	+	AATGGCCTCCACCATCCATCACCCGGCT		0	0	0	3	0	0	0	1
Oe_mir_486	FKYM01002413.1_7066:7097	+	TTTCATAGCTCAGTTGGTTAGAGCACCCGTTT		0	0	0	4	0	0	0	1
Oe_mir_487	FKYM01002296.1_87009:87040	+		ATCGTGGATCGGTGGACAGCATTCTTTAACC	0	0	0	3	0	0	0	1
Oe_mir_488	FKYM01002105.1_14741:14770	-	CTTGAGGATGGCCGCACTCTTGCCGATTAC		0	0	0	3	0	0	0	1
Oe_mir_489	FKYM01001537.1_46651:46681	+		CTCAGAGTTAACTATTTGTATGTCGTGTAGA	0	0	0	3	0	0	0	1
Oe_mir_490	FKYM01000977.1_18441:18469	+		TGAGACTGGTGTGGATATGGGGAGTGGT	0	0	0	3	0	0	0	1
Oe_mir_491	FKYM01000760.1_40439:40467	+		GGTTTACCGGTGTGTACCCCTTGACCCT	0	0	0	7	0	0	0	2
Oe_mir_492	FKYM01000760.1_40646:40673	+	GGATTGAATGGACTGAGACCGAGGAACT		0	0	0	1	0	0	0	1