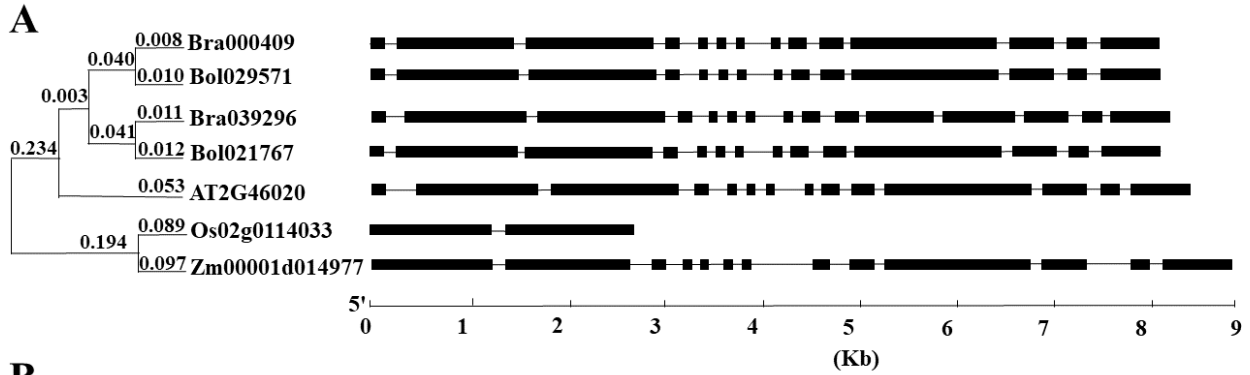


Figure S1. Comparison of the homologs of the AtBRM protein in *Brassica rapa*, *B. oleracea*, *Zea mays*, and *Oryza sativa*.

(A) Phylogenetic relationship and gene structures of the homologs of the AtBRM protein in those species. The unrooted phylogenetic tree was generated by the neighbor-joining method using the MEGA5 program with 1,000 bootstraps. Lines represent the introns while solid boxes represent exons. (B) Comparison of the deduced amino acid sequences of the homologs of the AtBRM protein in those species.



B

Bra000409	MQSGGSGGGPARNPGMGPGGRTNSTSSAASPSSSSSVQQQQQ-----LASRQQ	50
Bol029571Q-----	51
Bra039296HLA-----SRQ..	50
Bol021767	.L.....--...A.....T.....Q-----LASR.	48
AT2G46020A...A...A.....QQQQQQQQQLASRQ..	60
Os02g0114033	-----MQP.EAPPSSGG.P.T..RPPPAVA.G...LG-----FRNQA	37
Zm00001d014977	-----	0
Bra000409	HQLRNSEGNDGMFAYHAGGAQGMGAGSFG---SPSASMQ---PRKFFDSPQLQQ----	99
Bol029571G.....LPQQ.....	104
Bra039296	Q.G...A.....P..V.....G.N.---S.G...QPQQ..RL.....Q.----	103
Bol021767	Q.G...A.....P..V.....G.N.---...G...HPQQ..RL.....Q..QQPQ	105
AT2G46020	Q.H...DT.EN...QP..V.....G.NFA---.SPG...MPQQS.N..E...Q..QQQQ	117
Os02g0114033	MMHHHDQQQQQQQ.PS.APH...G.GSSSFPPS.GP.PPFQGG.NMPLP-GGP.G---	93
Zm00001d014977	MMHHHEQQQ---.F.S.APH...P.GVN-FPQS.GPVSSFQGG.NLPL.SGGP.G---	52
	: : : : : : * . : * * * * * . : : : * . : *	
Bra000409	---QHGEQQSINPMQAYIQFALQAQQQK-----AQQQARMGIMMGSSKQDAR	146
Bol029571	-----H.....	151
Bra039296	--GSSQ....F.....L.....M.-.....	150
Bol021767	--GSSQD....F.....L.....V..H...M.-----	143
AT2G46020	QGSSTQ...NF.....M..H.....MVG--.SVGKDQ	156
Os02g0114033	---LA.GQ.HNPTA...L.YMM.Q...--AHGMLLQQ...K.NMAGP.TR...VA	147
Zm00001d014977	--MV.GQMHNQVA...QFLKL.M.Q...QQKAAQGMLLQQ--.K.NMAGS..R..ML	107
Bra000409	MGVLNMQEMMPMQASNMAQASSSRPAG-----EQFSHGERQMESGPPQQRNETKPHP-QQ	199
Bol029571T.....S.....LA.....G.....	204
Bra039296	..M...GA.....Q..G...N.SA-----AR.....H.....	199
Bol021767	-----GA.-Q...Q..G...N.SA-----AR.....H.....	185
AT2G46020	DARMG.LN.QDLNP.SQP....KPSG-----D..AR...T..SS...S..Q..	210
Os02g0114033	ANTAK...L.SL..QAQ..MFKRQQSEHLQQAQEK.AEQ.QPSNSEQRSGDMRPPM.P.G	207
Zm00001d014977	NNPAK...L.ALH---Q..MYKRQCE-----KEQ.QSSGSEQRSGDMRPP-M.P.G	155
	: * . . . * . . . * : * . . . * *	
Bra000409	VGTGQLMPGNI IREMQAPQAMQGVNMGTNQLALSQQWQAMQAWARERNIDLSHPANANQ	259
Bol029571	A.....K.....S...H.....T.....	264
Bra039296GP.....F.....S.	259
Bol021767T.....GP.....S...F.....S.	245
AT2G46020M.....Q.L...N...FA.....S.	270
Os02g0114033	.PGQ..SSAGMV...PM.GQA.MS.A.A.PM.M-A.L.I...K.H.V...N...VTL	266
Zm00001d014977	..GQ..PSMGM...PI.GQV.MGSA.G.PITP-T.F.I...K.H.F...N...ISA	214
	. ** . : : * * * . * . : * * : * * * * * * * * * * * * * * * * * *	
Bra000409	MAHIL----QGRMAAQKAIIEGNVASQSPTMPVSSQPVSSSALPGENSPRANSASDISGQ	315
Bol029571	.S.....A.....A.....V.....F...	320
Bra039296	.S.....V.....N.....SI.....S...GV.....P...G....	315
Bol021767	.S.....SN.....P..SI.....S...GV.....P...G....	301
AT2G46020A.....G.....SI.I...A...VV.....H.....	326
Os02g0114033	ISQ..PMLQSN...M..QN.VGM..QQ-SVP.QMN--NDA..HSNFPSQG.PSKPR.	322
Zm00001d014977	ISQL.PIWQSN...M..QN.ANM.A.QQQALP.QVN---DT..HVNAPSQG.LLKPR.	271
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Fig. S1 (Cont.)

Bra000409	FHDWFAQPFQREGPAHN-IEDDWLETEKKVIVIHRLHQILEPFMLRRRVEDVEGSLPPK-	1194
Bo1029571-.....-	1200
Bra039296S.....-	1201
Bo1021767S.....-	1187
AT2G46020A.....-	1212
Os02g0114033	.Q...SK...DV.T.SEE.....I.....R.V	1178
Zm00001d014977	.S...SK...D..T..EE.....I.....R.-	1159
	* **::**::*: * . *****:***** * *	
Bra000409	-----VSVVLRCRMSAIQSAVYDWIKATGTLRVDP	1224
Bo1029571	-----	1230
Bra039296	-----S.....	1231
Bo1021767	-----S.....	1217
AT2G46020	-----	1242
Os02g0114033	DCSALVTLSDAICHSAFVCYIVNSSCHLQE.I.....G..G.I.....S...I....	1238
Zm00001d014977	-----D.I.....V.G.I.....S...I....	1189
	*:*****:*. *:*****:***:****	
Bra000409	DDEKLRAQKSSIIYQAKIYRTLNNRCMELRKT CNHPLLNYPYFNDLSKDFLVRSCGKLWIL	1284
Bo1029571T.....	1290
Bra039296K...NP.....K.....	1291
Bo1021767K...NP.....K.....	1277
AT2G46020NP.....A.....F.....	1302
Os02g0114033	E...A.I.RNAM...T.KN...K.....V.....S.FM.YYG...II.....N.	1298
Zm00001d014977	E...R...RNPM..V.T.KN...K.....V.....S.FL.-HG...MI.....N.	1248
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Bra000409	DRILIKLQRTGHRVLLFSTMTKLLDILEEYLQWRRLVYRRIDGTTSLLEDRESAIVDFNDP	1344
Bo1029571	1350
Bra039296	1351
Bo1021767	1337
AT2G46020	1362
Os02g0114033H.S.....Q.....R.	1358
Zm00001d014977HKS.....M.D.....R.	1308
	*****: :*****:*****:*****:*****:***** *	
Bra000409	DTDCFIFLLSIRAAGRGLNLQTDADTVVIYDPDPNPKNEEQAVARAHRIQGTREVKVIYME	1404
Bo1029571	1410
Bra039296	1411
Bo1021767	1397
AT2G46020	1422
Os02g0114033	NS.....S.....Q.....D.....	1418
Zm00001d014977	GS.....S.....Q.....	1368
	:*****:*****:*****:*****:*****:***** *	
Bra000409	AVVEKFSSHQKEDELRSRGGSIDVEDDLAGKDRYIGSIEGLIRNNIQQYKIDMADEVINAG	1464
Bo1029571	1470
Bra039296I.....M.L.....L.....	1447
Bo1021767M.....V.L.....	1457
AT2G46020L.....V.L...M.....	1482
Os02g0114033	...DNI..Y.....N..G.L.....M...S.....	1478
Zm00001d014977	...DNI..Y.....N..G.L.....M.....	1428
	: : **::**.*** *:***:**** :****	
Bra000409	RFDQRTTHEERRMTLETLLHDEERYQETVHDVPSLHEVNRMIARSEEEVELFDQMDEEFD	1524
Bo1029571	1530
Bra039296D.....	1507
Bo1021767	1517
AT2G46020	1542
Os02g0114033QQ.....T.....	1538
Zm00001d014977DS.....Q.....T.S.....D..	1488
	*****:*****:*****:*****:*****:***** *	

