

Supplemental Figure S7. Amino acid sequence alignment of the leucine zipper region of 170 ZmbZIP proteins.

The ZmbZIP proteins are categorized into 26 sub-families (BZ1-BZ26) with similar predicted dimerization properties. The leucine zipper region is divided into heptads (*gabcdef*) from L0 to L9 to visualize the potential *g*↔*e*' pairs. Four colors are used to differentiate between different *g*↔*e*' pairs. Attractive basic-acidic pairs (R↔E and K↔E) are colored orange, attractive acidic-basic pairs (E↔R, E↔K, D↔R, and D↔K) are blue, repulsive basic pairs (K↔K, R↔K, R↔Q, Q↔K, and K↔Q) are pink and repulsive acidic pairs (E↔E, E↔D, E↔Q, and Q↔E) are green. If single amino acid at the positions *e* or *g* is charged, the residue is colored pink for basic amino acid and green for acidic amino acid. If the *a* or *d* position is charged, it is colored brown. Asparagines at *a* position are colored red. The prolines and glycines are bold to indicate a potential break in the α -helix. The predicted C-terminal boundary is denoted by the symbol *, other than the natural terminals which are indicated by the symbol ^.

Sub-family	ZmbZIP NO.	Leucine zipper region									
		L0 gabcdef	L1 gabcdef	L2 gabcdef	L3 gabcdef	L4 gabcdef	L5 gabcdef	L6 gabcdef	L7 gabcdef	L8 gabcdef	L9 gabcdef
BZ1	ZmbZIP39	YNELEN	KVSRLEE	ENKKLTT*	KLEENI	LFSEPLP	EPKYQLR	RTGSATF	KQPQPPP	PPPPPIQ	QLPPKHR
	ZmbZIP94	YNELEN	KVARLKE	ENKRKIE*	LKVLPK	EPLEPPP	AEPONKKQ	PPPOPKR	KQPQPPP	PPPPPIQ	QLPPKHR
	ZmbZIP99	YNELEN	KVFRLEE	ENKRLLK*	QQLDEI	LSSAPP	EPKYQLR	RTGSAAF			
BZ2	ZmbZIP14	YMELEA	EVQKLKE	QNEELQK	KQEEMIE*	MQKNQVV	EVISNPY	AQKKRCL	RRTLTGP	W	
	ZmbZIP23	YMELEA	EVQKLRE	QNQELER	KQAEIME*	MQNNEVS	EMLKDPPF	GRKKRLC	RRRTLTA	PW	
	ZmbZIP46.1	YMELEA	EVQKLKE	QNAELQK	KQVPELV*	SNPYAQK	KRCLRRT	LTGPW			
	ZmbZIP46.2	YMELEA	EVQKLKE	QNAELQK	KQEIQME*	MQQNQVP	ELVSNPY	AQKKRCL	RRTLTGP	W	
	ZmbZIP107	YMELEA	EVAKLKD	QNEELQK	KQVEMILK*	KQKDEV	ERNSQH	GPKAKKL	CLRRTL	GPW	
BZ3	ZmbZIP31	YLTELEA	KAKGKLE	RNAELQF	RVSTLQ	ENNTLQ	ILKNTTA*	HASKRS	GGGGGG	KGGDGK	KHHLAKS
	ZmbZIP67	YLTDLEV	KVRDLK	KNSEMIE	RLSTLQ	ENQMLRQ	ILKNTAV*	NRRGS	TASGEFH	GQ	
	ZmbZIP97.1	YLTELEA	RAKDLKE	RNAELQF	RVSTLQ	ENNTLQ	ILKNTTA*	HASKRS	GGGKGGD	GGKKHHF	AKS
	ZmbZIP97.2	YLTELEA	RAKDLKE	RNAELQF	RVSTLQ	ENNTLQ	ILKNTTA*	HASKRS	GGGKGGD	GGKKHHF	AKS
	ZmbZIP108	YLSELEV	RVKELEK	RNSELKE	KLSTLQ	ENQMLRQ	ILKNTTV*	NNRPG	SSSAGGD	SQ	
	ZmbZIP123.1	YLTELEA	KAKGKLE	RNAELQF	RVSTLQ	ENNTLQ	VIDVSGK*	EFKFTWA	SGRGE	DYEEQR	SGEDGNG
	ZmbZIP123.2	YLTELEA	KAKGKLE	RNAELQF	RVSTLQ	ENNTLQ	VIDVSGK*	EFKFTWA	SGRGE	DYEEQR	SGEDGNG
BZ4	ZmbZIP110.1	YIKDLET	KSKHLEA	ECRRLSY	ALQCYAA	ENMALRQ*	SLKDRP	LGAPTAT	QESAVLT	ETLPLVS	LLWLVSI
	ZmbZIP110.2	YIKDLET	KSKHLEA	ECRRLSY	ALQCYAA	ENMALRQ*	SLKDRP	LGAPTAT	QESAVLT	ETLPLVS	LLWLVSI
BZ5	ZmbZIP73	YVQELEK	EVRRRLV	ENLKLK	QCKQKL	DMAALVQ*	QSSKSS	SHIRTS	SSTQL		
	ZmbZIP75	YVENLEK	EVRRRLV	DNKLK	QCKELK	EVAALVL*	PTKSSL	RTSSTQF			
	ZmbZIP95	YVRELET	KVQLLQ	ENESLR	KYDEL*	SVEAVP	MVRKTLQ	RMPSAP			
BZ6	ZmbZIP28	YVEELEG	KVKAMOA	TIADLSA	RISCVTA	ENALKQ*	QLGGA	AAPPMP	MYPTMYS	LPMPWMH	PSYPMRG
	ZmbZIP79.1	YVEELEE	KVKSMHS	VINDLSC	KISFIAA	ENATLRQ*	OLGGVG	SGPPPG	YPPPPG	IHPWV	GYAMRPH
	ZmbZIP79.2	YVEELEE	KVKSMHS	VINDLSC	KISFIAA	ENATLRQ*	OLGGVG	SGPPPG	YPPPPG	IHPWV	GYAMRPH
BZ7	ZmbZIP32	ETEELAT	QVESLAA	ENTSLR	EIGRLTE	SSEKLR	ENSALMV	KLKDTAE*	PSPSKAA	ASPSSPR	ASAENFL
	ZmbZIP43	ETEELAR	RVELLTA	ENTSLR	EISRLR	SSQKLR	ENSALME	KLADGT	DQAQEAS	AGHQTP	TAPSSAR
	ZmbZIP83.1	ETEELAT	QVESLAA	ENTSLR	EIQQLTE	SSEKLR	ENSALMV	KLKDTAE*	PSPIKAS	ASPSSPR	ASAENFL
	ZmbZIP83.2	ETEELAT	QVESLAA	ENTSLR	EIQQLTE	SSEKLR	ENSALMV	KLKDTAE*	PSPIKAS	ASPSSPR	ASAENFL
	ZmbZIP42	ETEELAT	QVESLAA	ENTSLR	EIGRLTE	SSEKLR	ENSALMA*	ISCOSAF	PLTRGAL	SLRSQP	KSKLMHY
BZ8	ZmbZIP124	ETEELAT	QVESLAA	ENTSLR	EIGRLTE	SSEKLR	ENSALMA*	ISCOSAF	PLTRGAL	SLRSQP	KSKLMHY
	ZmbZIP33	ECEELAQ	RADVLKQ	ENASLRD	EVNRIRK	EYEELLS	KNNSLK*	KLEGKQH	KTDEAGL	NNKLQHS	GNDIJKK
	ZmbZIP58	ECEELGQ	RAEELR	ENSSLRA	ELERIRK	EYEQLLS	QNASLKE*	KLGASS	DSLDPDMN	EQNDG	DGGYRKQ
	ZmbZIP63	ECEELGQ	RAETLRS	ENSSLRA	ELERIRK	EYEQLS	QNASLKE*	KLGGSS	PIPDMNE	QNDGN	DKKQSDS
BZ9	ZmbZIP120	ECEELAQ	RADVLKQ	ENASLR	EVNRIRK	EYEELLS	RNNLSS*	KLEGKQH	KTDEAGL	NNKLQHS	GDDSQKK
	ZmbZIP2	EWEEVAN	RADLLKQ	ENSSLKE	ELKQLQE	KCDGLTS	ENTSLHE	KLKALE*	EKSNGNW	YKD	
BZ10	ZmbZIP21.1	ECEELAR	KVADLTT	ENSALRA	ELDNLRK	ACQDMAE	ENSLRLL*	STVPSVT	TTLGMSI	EPPKAQ	HHDEGQ
	ZmbZIP21.2	ECEELAR	KVADLTT	ENSALRA	ELDNLRK	ACQDMAE	ENSLRLL*	STVPSVT	TTLGMSI	EPPKAQ	HHDEGQ
	ZmbZIP86.1	ECEELAR	KVADLTT	ENSALRA	ELDNLRK	ACQDMAE	ENSLRLL*	GVAADQV	PSVITTL	GMSIEPP	KLOLQLQ
	ZmbZIP86.2	ECEELAR	KVADLTT	ENSALRA	ELDNLRK	ACQDMAE	ENSLRLL*	PSVITTL	TALVAMG	SLQRLQ	QHDEEG
	ZmbZIP122	ECEELAQ	KVTDLTV	VNGMLRS	ELDELKK	ACEDMEA	VTSVMV*			SS	
BZ11	ZmbZIP51	IRDELAR	KVADLSS	QNNENK	EKDVMVM	EYSLKE	ANKQLE	QVARTTA*	KKAPAGS	LAAAAS	
	ZmbZIP118	VRDELAR	KVADLSS	QNATMK	EKDVMVM	EYSLKE	TNEQLKA	AEQAIHH	HHLRLSS*	LF	
	ZmbZIP12	ZmbZIP11	QLSELWA	QVSHLRG	ANRRLLD	DLNRLR	SCADARR	ESARLRE	EKAELTK	KLEQLLQ*	EAKGSL
	ZmbZIP54	QLSELWA	QVSHLRG	ANRRLLD	DLNRLR	SCADARR	ENARLRR	EKAQLAN	RHQLLQ*	IA	
	ZmbZIP57	RLHELSL	RAEELLG	ANQRLL	DLNRVVA	RHGAVAR	ENARLRE	AAEGLR	RIGEVEV*	GEAAAAA	AGAGQPL
BZ12	ZmbZIP59	HLDELT	QAAHLLR	ENAHVAT	ALGLTAQ	GLLAVDA	DNAVLRT	QAAEELA	RGLSLND*	ILACMNT	TNAAAA
	ZmbZIP64	HLDELT	QAAHLLR	ENAHVAT	ALGLTAQ	GLLAVDA	ENAVLRT	QTAELAA	RGLSLND*	ILACMNT	NAVAVA
	ZmbZIP66	RLYELSL	QVAELLG	TNQRLRV	ELNHVTA	KYALLAR	ENAKLRE	AAAGLQR	RLE		
	ZmbZIP72	QLSELWA	QVHHLRS	TNQRLLD	QLNHAIR	DCDRVLR	ENSQRLD	EOTKLOQ*	QLEMPLV	DITTESGA	MSPGS
	ZmbZIP78	QLQQLWD	QVHHLRS	DSRDLD	RNLAIR	DCDRVMR	DNARLRR	ERAGLQR	RLLDIT*	DGDGDDP	PF
BZ13	ZmbZIP109	QLDELSS	QVHHLRS	DSRDLD	RNLAIR	DCDRVMR	DNARLRR	EAGALR	RLAAAE*	AAAAGH	GQVRVVP
	ZmbZIP10	HLEELRG	RAARLRA	GNRDLAA	RIGGQA	RAALAR	ANARLRA	EAGALAR	RLYAARR*	ALALRQV	YAASGG
	ZmbZIP22	HHDDLT	QVDOLKG	ONQQLNL	ALSTTSQ	NLVAVQA	QNSVLT	QRMELAS	RLGALTE*	ILWCSS	STGTAA
	ZmbZIP24	QLTELCA	QVHHLRS	ANRRLLD	DLNRLR	GCSDMCC	ENARLQ	EKTDLST	KLERLTQ*	SSSEPO	
	ZmbZIP44	QLDDLT	QVQALRA	RNGALMD	AARDAR	RCVYQA	ENALMHA	RTVELSA	RQLSLVD*	LICQCMQ	GDAMYQY
BZ14	ZmbZIP87	HLDLDT	QNKQLSD	ONQQLSM	ALSITSQ	NLVAVQA	QNSVLT	QRMELDS	RLGALTE*	ILWYMN	STISTSTA
	ZmbZIP88	QLTELWA	QVQVHLR	ANRRLLD	ELNRAVR	GCSDMIR	ERAKLQK	EKTDLGT	RQLRQLTQ*	PQNAALP	SSSEPO
	ZmbZIP49	QLSELWA	QVQVHLR	TNQRLDQ	OLKCSV	SSETATV	SSVKTPG				
	ZmbZIP124	ETEELAT	QVESLAA	ENTSLR	EIGRLTE	SSEKLR	ENSALMA*	ISCOSAF	PLTRGAL	SLRSQP	KSKLMHY
	ZmbZIP125	ETEELAT	QVESLAA	ENTSLR	EIGRLTE	SSEKLR	ENSALMA*	ISCOSAF	PLTRGAL	SLRSQP	KSKLMHY
BZ15	ZmbZIP112.1	QLSDLES	QVERLKG	ENATLFQ	QLSDAQ	QFSTAVT	DNRILKS	DVEALRI	KVKMAED*	MVARSAV	SCGLGDL
	ZmbZIP112.2	QLSDLES	QVERLKG	ENATLFQ	QLSDAQ	QFSTAVT	DNRILKS	DVEALRI	KVKMAED*	MVARSAV	SCGLGDL
	ZmbZIP65.1	HLADLET	QVDQLRG	ENASLFK	QLTDANQ	QFTTAVT	DNRILKS	DVEALRV	KVKLAED*	MVARGAL	SCGLGSL
	ZmbZIP65.2	HLADLET	QVDQLRG	ENASLFK	QLTDANQ	QFTTAVT	DNRILKS				