

**Supplemental data. Köllner, T.G. et al. (2008).
A Maize (*E*)- β -Caryophyllene Synthase Implicated in Indirect
Defense Responses Against Herbivores is Not Expressed in
Most American Maize Varieties.**

Supplemental Figure 1.

20 40 60

TPS23 - Del MAADEARSVSRLHSEEDMHGKHHSTLWGDFFLHHVPCRPQYLIMKDNVEIMKEEVKKMLLDVGSSDLSH
 TPS23 - Mex MAADEARSVSRLHSEEDMHGKHHSTLWGDFFLHHVPCRPQYLIMKDNVEIMKEEVKKMLLDVGSSDLSH
 TPS23 - Hue MAADEARSVSRLHSEEDMHGKHHSTLWGDFFLHHVPCRPQYSIMKDNVKIMKEEVKKMLLDVGSSDLSH
 TPS23 - Par MAADEARSVSRLHSEEDMHGKHHSTLWGDFFLHHVPCRPQYLIMKDNVEIMKEEVKKMLLDVGSSDLSH
 TPS23 - Per MAADEARSVSKLHSEEDMHGKHHSTLWGDFFLHHVPCRPQYLIMKDNVEIMKEEVKKMLLDVGSSDLSH
 TPS23 - Dip MAADEARSVSRLHSEEDMHGKHHSTLWGDFFLHHVPCRPQYLIMKDKVKIMKEEVKKMLLDVGSSDLSH
 TPS23 - Lux MAADEARSVSRLHGEEDMHGKHHSTLWGDFFLHHVPCRPQYLIMKDNVEIMKEEVKKMLLDVGSSDLSH

80 100 120 140

TPS23 - Del KLDCIDTLERLGLDYHYTKEIDELMCNVFEARDQDLDLTTTSQLFYLLRKHGYHISSDVFLKFRRDKGDI
 TPS23 - Mex KLDCIDTLERLGLDYHYTKEIDELMCNVFEARDQDLDLTTTSQLFYLLRKHGYHISSDVFLKFRRDKGDI
 TPS23 - Hue KLECIDTLERLGLDYHYTKEIDELMCNVFEARDQDLDLTTTSQLFYLLRKHGYHISSDVFLKFRRDKGDI
 TPS23 - Par KLDCIDTLERLGLDYHYTKEIDELMCNVFEARDQDLDLTTTSQLFYLLRKHGYHISSDVFLKFRRDKGDI
 TPS23 - Per KLDCIDTLERLGLGYHYTKEIDELMCNVFEARDQDLDLTTTSQLFYLLRKHGYHISSDVFLKFRRDKGDI
 TPS23 - Dip KLDCINTLERLGLDYHYTKEIDELMCNVFEARDQDLDLTTTSQLFYLLRKHGYHISSDVFLKFRRDKGDI
 TPS23 - Lux KLDCIDTGLRLGLDYHYTKEIDELMCNVFEAMDQDLDLTTTSQLFYLLRKHGYHISSDVFLKFRRDKGDI

160 180 200

TPS23 - Del VTNDARCLLRMYEAHVVRVNNGEEILDNIIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRRLNRVQARQ
 TPS23 - Mex VTNDARCLLRMYEAHVVRVNNGEEILDNIIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRRLNRVQARQ
 TPS23 - Hue VTDARCLLRMYEAHVVRVNNGEEILDNIIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRRLNRVQARQ
 TPS23 - Par VTDARCLLRMYEAHVVRVNNGEEILDNIIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRRLNRVQARQ
 TPS23 - Per VTDARCLLRMYEAHVVRVNNGEEILDNIIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRRLNRVQARQ
 TPS23 - Dip VTDARCLLRMYEAHVVRVNNGEEILDNIIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRRLNRVQARQ
 TPS23 - Lux VTDARCLLRMYEAHVVRVNNGEEILDNIIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRRLNRVQARQ

220 240 260 280

TPS23 - Del FISTYEKSTTRINMLLEFSKLDFNILLTLYCEELKDLTLWWKEFQQAQANTITIYARDRMVEMHFWMGVFF
 TPS23 - Mex FISTYEKSTTRINMLLEFSKLDFNILLTLYCEELKDLTLWWKEFQQAQANTITIYARDRMVEMHFWMGVFF
 TPS23 - Hue FISTYEKSTMNNMLLEFSKLDFNILLTLYCEELKDLTMWWKEFQQAQANTITIYARDRMVEMHFWMGVFF
 TPS23 - Par FISTYEKSTTRNNMLLEFSKLDFNILLTLYCEELKDLTLWWKEFQQAQANTITIYARDRMVEMHFWMGVFF
 TPS23 - Per FISTYEKSTTRNNMLLEFSKLDFNILLTLYCEELKDLTLWWKEFQQAQANTITIYARDRMVEMHFWMGVFF
 TPS23 - Dip FISTYEKSTTRNNMLLEFSKLDFNILLTLYCEELKDLTLWWKEFQQAQANTITIYARDRMVEMHFWMGVFF
 TPS23 - Lux FISTYEKSTTRNNMLLEFSKLDFNILLTLYCEELKDLTLWWKEFQQAQANTITIYARDRMVEMHFWMGVFF

300 320 340

TPS23 - Del EPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNAFTAALQRWDEEGVEQCPTYLRTLYTNIRATIKAI
 TPS23 - Mex EPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNAFTAALQRWDEEGVEQCPTYLRTLYTNIRATVKAI
 TPS23 - Hue EPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNAFTAALQRWDEEGVEQCPTYLRTLYTNIRATVKAI
 TPS23 - Par EPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNAFTAALQRWDEEGVEQCPTYLRTLYTNIRATIKAI
 TPS23 - Per EPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNTFTAALQRWDEEGVEQCPTYLRTLYTNIRATVKAI
 TPS23 - Dip EPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNTFTAALQRWDEEGVEQCPTYLRTLYTNIRATVKAI
 TPS23 - Lux EPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNAFTAALQRWDEEGVEQCPTYLRTLYTNIRATVKAI

360 380 400 420

TPS23 - Del EEDLNQNKKHAKLVKGIIIDMAMCYNAETEWRDKKYVPATVDEHLKISARSSGCMHLVSQGFISMGDVA
 TPS23 - Mex EEDLNQNKKHAKLVKGIIIDMAMCYNAETEWRDKKYVPATVDEHLKISARSSGCMHLVSQGFISMGDVA
 TPS23 - Hue EEDLNQNKKHAKLVKGIIIDMAMCYNAETEWRDKKYVPATVDEHLKISARSSGCMHLVSQGFISMGDVA
 TPS23 - Par EEDLNQNKKHAKLVKGIIIDMAMCYNAETEWRDKKYVPATVDEHLKISARSSGCMHLVSQGFISMGDVA
 TPS23 - Per EEDLNQNKKHAKLVKGIIIDMAMCYNAETEWRDKKYVPATVDEHLKISARSSGCMHLVSQGFISMGDVA
 TPS23 - Dip EEDLNQNKKHAKLVKGIIIDMAMCYNAETEWRDKKYVPATVDEHLKISARSSGCMHLVSQGFISMGDVA
 TPS23 - Lux EEDLNQNKKHAKLVKGIIIDMAMCYNAETEWRDKKYVPATVDEHLKISARSSGCMHLVSQGFISMGDVA

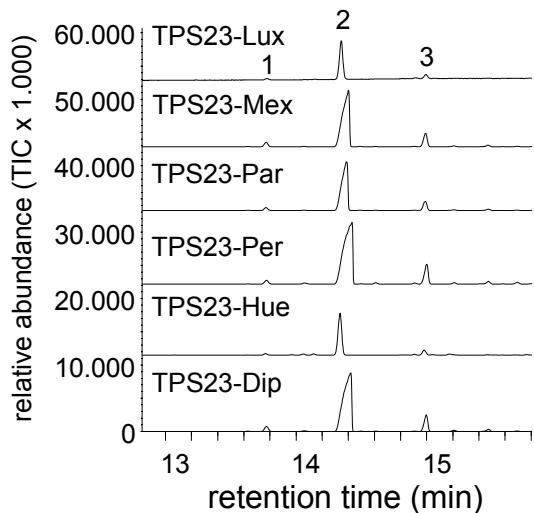
440 460 480

TPS23 - Del TSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTTTVEQAIIEKIRELIEEA
 TSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTTTVEQAIIEKIRELIEEA
 TSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTTTVEQAIIEKIRELIEEA
 TSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTTTVEQAIIEKIRELIEEA
 TSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTTTVEQAIIEKIRELIEEA
 TSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTTTVEQAIIEKIRELIEEA
 TSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTTTVEQAIIEKIRELIEEA

500 520 540

TPS23 - Del WMDTHECLRQPQPKALLERAVNLARTMDFLYKDADGYTDSRSIKGILDSLYVHLID
 TPS23 - Mex WMDTHECLRQPQPKALLERAVNLARTMDFLYKDVGYTDSRSIKGILDSLYVHLID
 TPS23 - Hue WMDTHECLRQPQPKALLERAVNLARTMDFLYKDVGYTDSWSIKGILDSLYVHLID
 TPS23 - Par WMDTHECLRQPQPKALLERAVNLARTMDFLYKDVGYTDSRSIKGILDSLYVHLID
 TPS23 - Per WMDTHECLRQPQPKALLERAVNLARTMDFLYKDVGYTDSRSIKGILDSLYVHLID
 TPS23 - Dip WMDTHECLRQPQPKALLERAVNLARTMDFLYKDVGYTDSRSIKGILDSLYVHLID
 TPS23 - Lux WMDTHECLRQPQPKALLERAVNLARTMDFLYKDVGYTDSRSIKGILDSLYVHLID

Supplemental Figure 1. Amino acid alignment of the putative TPS23 orthologs from maize and the teosinte species *Z. m. parviflora*, *Z. luxurians*, *Z. m. mexicana*, *Z. diploperennis*, *Z. perennis*, *Z. m. huehuetenangensis*. The complete ORFs were aligned using CLUSTAL W (see methods section). Amino acids identical in all sequences are marked by black boxes and amino acids identical in at least four sequences are marked by grey boxes.



Supplemental Figure 2. Sesquiterpene products of the putative TPS23 orthologs. Putative orthologs of TPS23 were cloned from the teosinte species *Z. m. parviflora* (Par), *Z. luxurians* (Lux), *Z. m. mexicana* (Mex), *Z. diploperennis* (Dip), *Z. perennis* (Per), *Z. m. huehuetenangensis* (Hue). The genes were expressed in *E. coli*, extracted, partially purified and incubated with the substrate (*E,E*)-FPP. The resulting terpene products were collected with a SPME fiber and analyzed by GC-MS. The products were identified as δ -elemene (1), (*E*)- β -caryophyllene (2) and α -humulene (3) by comparison of their retention times and mass spectra to those of authentic standards.

Supplemental_Dataset_1.txt

>TPS10-B73

533 weight: 0.57

-----MDATAFHPSLWGDFVKYK-PPTAPKRGHMT
ERAELLKEEVRKTLKAAA-NQITNALDL--IITLQLRLGDHHYENEISEL
LRFVYSSSDYDDK-----DLYVVSLRFYLLRKHGCVSSDVFTSFK
DEEG--NFV---VDDTKCLLSLYNAAYVRTHGEKLDEAITFTRRQLEAS
LLD-PL----EPALADEVHLTQTPLFRRRLRILEAINYI-----
--PIYGKEAGRNEAILEAKLFNLAQLIYCEELKEVTLWWKQLNVETNL
S-FIRDRIVECHFWMTGACCEPQYSLSRVIATKMTALITVLDDMMDTYST
TEEAMLLAEAIYRWEENAAELLP--RYMKDFYLKLKTIDSCGDELGPN-
RSFRTFYLKEMLKVLVRGSSQEIKWRNENYVPKTISEHLEHSGPTVGAFQ
VACSSFVGMGD-SITKESFEWLTYPELAKSMLNISRLLNDTASTKREQN
AGQHVSTVQCYMLKHGTT-MDEACEKIKELTEDSWKDMMELYLTPTE--H
PKLIAQTIVDFARTADYMYKETDGFTFSH-TIKDMIAKLFVDPISLF---

>TPS4-B73 554 weight: 0.57

-MASPPAHRSSKADEELPKASSTFHPSSLWGSFFLTYQ-PPTAPQRANMK
ERAEVLRERVRKVLKGSTTDQLPETVNL--ILTQLRQLGYYYENEIDKL
LHQIYSNSDYNVK-----DLNLVSQRFYLLRKNGYDVPSDVFLSK
TEEG--GFA-CAAADTRSLLSLYNAAYLRKHGEVLDEAISSTRRLQDL
LGR-LLP---ESPFAKEVSSSLRTPLFRRVGILEARNYI-----
--PIYETEATRNEAVLEAKLNPNLQQLDFCEELKHCSAWNEMIAKSKL
T-FVRDRIVEEYFWMNGACYDPPYLSRIILTKITGLITIIDDMFDTHGT
TEDCMKFAEAFCRWDESAIHLLP--EYMKDFYILMLETFQSfedALGPE
KSYRVLYLKQAMERLVELYSKEIKWRDDDYVP-TMSEHLQVSAETIATIA
LTCSAYAGMGDMSIRKETFEWLSFPQFIRTFGSFVRLSNDVVSTKREQT
KDHSPTVHCYMKEHGT-MDDACEKIKELIEDSWKDMLEQLSALKG--L
PKVVPQLVFDFSRRTDNMYRDRDALTSSE-ALKEMIQLLFVEPIPE---

>TPS23-Del 547 weight: 1.11

-MAADEARSVSLRHSEE--DMHGKHHSTLWGDFLHHV-PCRPGQYILMK
DNVEIMKEEVKKMLLDVGSSDLSHKLDC--IDTLERLGLDYHYTKEIDEL
MCNVFEARDQDL-----DLTTSQLFYLLRKHGHISSDVFLKFR
DDKG--DIV---TNNDARCLLRMYEAAHVRVNGEIILDNLIHTKRQLQCI
VDD-L----EPTLQEERYALETPLFRRRLNRVQARQFI-----
--STYEKSTTRINMLLEFSKLDNFNILLTLYCEELKDLTLWWKEFQAQANT
TIYARDRMVEMHFWMGMGVFFEPQYSYSRKMLTQLFMIVSVLDDLYDSHCT
TEEGNAFTAALQRWDEEGVEQCP--TYLRTLTYTNIRATIKAIIEEDLFQZ
NNKHAKLVKGLIIDMAMCYNAETEWRDKKYVPATVDEHLKISARSSGCMH
LVSQGFISMGDVA-TSEALEWASTYPKIVRAVCIIARLANDIMSYKREAS
NNTMVSTVQTCAKEYGTTTVEQAIKEIRELIEEAWMDITHECLRQPQ---
PKALLERAVNLARTMDFLYKDADGYTDSR-SIKGILDSDLVHLD-----

>At_TPS27 547 weight: 1.04

-----MGSEVRPLADFPANIWEDPLTSFSKSDLGTET--FK
EKHSTLKEAVKEAFMSSKANPIENIKF---IDALCRLGVSCFHFEKDIVEQ
LDKSFDCDFPQMVRQEGC---DLYTVGIIFQVFRQFGFKLSADVFEKF
DENG--KFKGHLVTDAYGMLSLEYAAQWGTHGEDIIDEALAFSRSHLEEI
S---SRS---SPHLAIRIKNALKHYPYHKGISRIETRQYI-----

Supplemental_Dataset_1.txt

--SYYEEEESCDPTLLEFAKIDFNLLQILHREELACVTRWHHEMEFKSKV
T-YTRHRITEAYLWLSLGTYFEPQYSQARVITTMALILFTALDDMYDAYGT
MEELELFDTADMEWLVPVPDEIPIPDSMKFIYNTVEFYDKLDEELEKEG
RSGCGFHLKKSLQKTANGYMQEAKWLKDYIA-TFDEYKENAILSSGGYYG
LIAMTFVRMTDVA-KLDAFEWLNSHPKIRVASEIIISRFTDDISSYEFHK
REHVATGIDCYMQQFGVS-KERAVEVMGNIVSDAWKDLNQELMRPHV--F
PFPLLMRVLNLNSRVIDVFYRYQDAYTNPK-LLKEHIVSLLIETIPI---
--

>CSCS 567 weight: 0.82

MSSHFPASIMKTDIYDTKRSLANFHPTIWKEHFLSFTFDALVKVDGGMK
ERIEKLKEEIRMMVIASVQNPLVKLNL---VDSIQRLGVSYHFEDEIDQF
LEHMYVSYNNSSLFSSNDSDQDDDLHISALLFRLLRQHGYRISCDIFLKFM
DNNG--KFKESELVEDERGILSLYEASHMRGHGEALLEEALEFTTHLKAY
IHLYSNI---NPNFASEVSNALKLPIRKCVPRVKAREYF-----
--EIYQQQPSPHNETLLTFSKLDNFNILQKLHQKEIAEICRWKDLNVSTNF
P-FARDRIVECYFWILSIYFEPYFKFGRKILTAKVIAINTSIMDDIYDAYGT
FEELQLFTLAIRWDMSMVNLLP--QYMKVHYTLLDLFEEMDKGIVNDG
ISYRSCFGKEAMKRQAESYFKEAEWLKNYKP-KYGEYMEVALASSGYEL
LSTISFVCMGRIA-TKEVFELFDCPQILKASTTISRLMDDVSYKFEKE
REHIVSAVECYMSNHGRS-EDETCAELLKQVEDAWKTINECCLHPMN--V
PMPFLICLLNLTRVMALLYSHEDGYTNKGRTKLLIQSLLIDPLHL---
--

>QHS1 548 weight: 0.82

MSVKE-EKVIRPIVHFPPSVADQFL--IFDDKQAEQANVE
QVVNELREDVRKDLVSSLVQTEHTNLLKLIDAIQRLGIAYHFEEEIEQA
LQHIYDTYGDDWKGRS-----PSLWFRILRQQGFYVSCDIFKNYK
KEDG--SFKESELTDVEGLLELYEATYLRLVQGEGVLDALVFTRTCLEKI
AKDLVHT---NPTLSTYIQEALKQPLHKRLTRLEALRYI-----
--PMYEQQASHNESLLKLAQKLGFLQSLHRKELSEVSRWKGLDVPNNL
P-YARDRMVECYFWALGVYFEPKYSQARIFLAKVISLATVLDITYDAYGT
YEELKIFTEAIQRWSITCIDMLP--EYLKLLYQGVLDIYIEMEEIMGKEG
KAHHLSYAKESMKEFIRSYPEAKWANEQYVP-TAEEHMSVAFVSSGYSM
LATTCFVGMGDIV-TDEAFKWALTAKPIIKASCAIARLMDDIHSQKEEK
RIHVASSVESYMKQYDVT-EEHVLKVFNKKIEDAWKDITRESLVRKD--I
PMPLMMRVINLAQVMDVLYKHKGFTNVGEELKDHIKSLLVHPIPI---
--

>ATDTPS 785 weight: 2.07

MSINLRSSGCSSPISATLERRLDSEVQTRANNSFEQTKEKIRKMLEKVE
LSVSAYDTSWAVMVPSPSSQNAFLFPQCVKWLLDNQHEDGSWGLDNHDHQ
SLKKDVLSSTLASILAKKKWIGERQINKGLQFIELNSALVTDETIQKPT
GFDIIIFPGMIKYARDLNLTIPGLSEVVDMMIRKRDLDLKCDSEKFSKGRE
AYLAYVLEGTRNLKDWDLIVKYQRKNGLFDSPATTAAFTQFGNDGCLR
YLCSSLQKFEAAVPSVYPFDQYARLSI---IVTLESLGIDRDFKTEIKSI
LDETYRYWLRGDEEICL----DLATCALAFRLLLALAHGYDVSYDPLKPFA
EESGFSDTLEGYVKNTFSVLELFKAAQSYPH-ESALKKQCWCWTQYLEME
LSSWVKTSVRDKYLKEVEDALAFPSYASLERSDHRRKILNGSAVENTRV
TKTSYRLHNICTSDILKLAVALDDFNFCQSIHREEMERLDRWIVENRLQELK
--FARQKLAYCYFSGAATLFSPELSDARISWAKGGVLTTVVDFFDVGG
KEELENLIHLVEKWDLNGVPEYS-SEHVEIIFSVLRDTILETGDKAFTYQ
GRNVTHHIVKIWL DLLKSMLREAWESSDKSTP-SLEDY MENAYISFALGP
IVLPATYLIBPPL--PEKTVDSHQYNQLYKLVSTMGRLLNDIQGFKRESA
EGKLNAVSLHMKHERDNRSKEVIIESMKGLAERKREELHKLVLEEKGSVV
PRECKE AFLKMSKVLNLFYRKDDGFTSND--LMSLVKSVIYEPVSLQEE
LT

Supplemental_Dataset_2.txt

>"TPS23 Zea mays diploperennis.pro" Translate DNA Sequence TPS23 Zea mays diploperennis.seq(1,1644)
MAADEARSVSLHSEEDMHGKHHSTLWGDFFLHHVPCRPGQYLIMKDKVKIMKEEVKKMLLDVGSSDLSHKLDCTLE
RLGLDYHYTKEIDELMCNVFEARDQDLDLTTSQLFYLLRKHGTVSSDVFLKFRDDKGDIVTDDARCLLRMYEAHVR
VNGEEILDNLIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRLNRVQARQFISTYEKSTTRNNMLLEFSKLDFNILL
TLYCEELKDLTLWWEFQAQANTAIYARDRMVEMHFWMGVFFEPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNT
FTAALQRWDEEGVEQCPTYLRTLYTNIRATVKAIIEEDLNQNNKHAKLVKGLIIDMAMCYNAETEWRDKYYVPATVDEH
LKISARSSGCMHLVSQGFISMGDVATSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTT
TVEQAIKIRELIEEAWMDITHECLRQPQPKALLERAVNLARTMDFLYKDVVDGYTDSSRSIKGILDSLYVHLID.
>"TPS23 Zea mays huehuetenangensis.pro" Translate DNA Sequence TPS23 Zea mays huehuetenangensis.seq(1,1644)
MAADEARSVSLHSEEDMHGKHHSTLWGDFFLHHVPCRPGQYSIMKDNVKIMKEEVKKMLLDVGSSDLSHKLECIDTLE
RLGLDYHYTKEIDELMCNVFEARDQDLDLTTSQLFYLLRKHGTVSSDVFLKFRDDKGDIVTDDARCLLRMYEAHVR
VNGEEILDNLIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRLNRVQARQFISTYEKSTTRNNMLLEFSKLDFNILL
TLYCEELKDLTMWWWEFQAQANTTIYARDRMVEMHFWMGVFFEPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNA
FTAALQRWDEEGVEQCPTYLRTLYTNIRATVKAIIEEDLNQNNKHAKLVKGLIIDMAMCYNAETEWRDKYYVPATVDEH
LKISARSSGCMHLVSQGFISMGDVATSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTT
TVEQAIKIRELIEEAWMDITHECLRQPQPMALLERAVNLARTMDFLYKDVVDGYTDSSRSIKGILDSLYVHLID.
>"TPS23 Zea mays luxurians.pro" Translate DNA Sequence TPS23 Zea mays luxurians.seq(1,1644)
MAADEARSVSLHGEEDMHGKHHSTLWGDFFLHHVPCRPGQYLIMKDNVEIMKEEVKKMLLDVGSSDLSHKLDCTLG
RLGLDYHYTKEIDELMCNVFEAMDQDLDLTTSQLFYLLRKHGVISSDVFLKFRDDKGDIVTNDARCLLRMYEAHVR
VNGEEILDNLIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRLNRVQARQFISTYEKSTTRNNMLLEFSKLDFNILL
TLYCEELKDLTLWWEFQAQANMTIYARDRMVEMHFWMGVFFEPQYSYSRKMLTRLFMIVSVLDDLYDSHCTTEEGNA
FTAALQRWDEEGVEQCPTYLRTLYTNIRATVKAIIEEDLNQNNKHAKLVKGLIIDMAMCYNAETEWRDKYYVPATVDEH
LKISARSSGCMHLVSQGFISMGDVATSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTT
TVEQAIKIRELIEEAWMDITHECLRQPQPKALLERAVNLARTMDFLYKDVVDGYTDSSRSIKGILDSLYVHLID.
>"TPS23 Zea mays mays Delprim 1.pro" Translate DNA Sequence TPS23 Zea mays mays Delprim 1.seq(1,1644)
MAADEARSVSLHSEEDMHGKHHSTLWGDFFLHHVPCRPGQYLIMKDNVEIMKEEVKKMLLDVGSSDLSHKLDCTLE
RLGLDYHYTKEIDELMCNVFEARDQDLDLTTSQLFYLLRKHGVISSDVFLKFRDDKGDIVTNDARCLLRMYEAHVR
VNGEEILDNLIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRLNRVQARQFISTYEKSTTRNNMLLEFSKLDFNILL
TLYCEELKDLTLWWEFQAQANTTIYARDRMVEMHFWMGVFFEPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNA
FTAALQRWDEEGVEQCPTYLRTLYTNIRATIKAIIEEDLNQNNKHAKLVKGLIIDMAMCYNAETEWRDKYYVPATVDEH
LKISARSSGCMHLVSQGFISMGDVATSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTT
TVEQAIKIRELIEEAWMDITHECLRQPQPKALLERAVNLARTMDFLYKDVVDGYTDSSRSIKGILDSLYVHLID.
>"TPS23 Zea mays mexicana.pro" Translate DNA Sequence TPS23 Zea mays mexicana.seq(1,1644)
MAADEARSVSLHSEEDMHGKHHSTLWGDFFLHHVPCRPGQYLIMKDNVEIMKEEVKKMLLDVGSSDLSHKLDCTLE
RLGLDYHYTKEIDELMCNVFEARDQDLDLTTSQLFYLLRKHGVISSDVFLKFRDDKGDIVTNDARCLLRMYEAHVR
VNGEEILDNLIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRLNRVQARQFISTYEKSTTRNNMLLEFSKLDFNILL
TLYCEELKDLTLWWEFQAQANTTIYARDRMVEMHFWMGVFFEPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNA
FTAALQRWDEEGVEQCPTYLRTLYTNIRATVKAIIEEDLNQNNKHAKLVKGLIIDMAMCYNAETEWRDKYYVPATVDEH
LKISARSSGCMHLVSQGFISMGDVATSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTT
TVEQAIKIRELIEEAWMDITHECLRQPQPKALLERAVNLARTMDFLYKDVVDGYTDSSRSIKGILDSLYVHLID.
>"TPS23 Zea mays parviglumis.pro" Translate DNA Sequence TPS23 Zea mays parviglumis.seq(1,1644)
MAADEARSVSLHSEEDMHGKHHSTLWGDFFLHHVPCRPGQYLIMKDNVEIMKEEVKKMLLDVGSSDLSHKLDCTLE
RLGLDYHYTKEIDELMCNVFEARDQDLDLTTSQLFYLLRKHGVISSDVFLKFRDDKGDIVTNDARCLLRMYEAHVR
VNGEEILDNLIHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRLNRVQARQFISTYEKSTTRNNMLLEFSKLDFNILL
TLYCEELKDLTLWWEFQAQANTTIYARDRMVEMHFWMGVFFEPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNA
FTAALQRWDEEGVEQCPTYLRTLYTNIRATIKAIIEEDLNQNNKHAKLVKGLIIDMAMCYNAETEWRDKYYVPATVDEH
LKISARSSGCMHLVSQGFISMGDVATSEALEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTT
TVEQAIKIRELIEEAWMDITHECLRQPQPKALLERAVNLARTMDFLYKDVVDGYTDSSRSIKGILDSLYVHLID.
>"TPS23 Zea mays perennis.pro" Translate DNA Sequence TPS23 Zea mays perennis.seq(1,1644)
MAADEARSVSLHSEEDMHGKHHSTLWGDFFLHHVPCRPGQYLIMKDNVEIMKEEVKKMLLDVGSSDLSHKLDCTLE
RLGLGYHYTKEIDELMCNVFEARDQDLDLTTSQLFYLLRKHGVISSDVFLKFRDDKGDIVTDDARCLLRMYEAHVR
VNGEEILDNLSHTKRQLQCIVDDLEPTLQEEVRYALETPLFRRLNRVQARQFISTYEKSTTRNNMLLEFSKLDFNILL
TLYCEELKDLTLWWEFQAQANTAIYARDRMVEMHFWMGVFFEPQYSYSRKMLTQLFMIVSVLDDLYDSHCTTEEGNT
FTAALQRWDEEGVEQCPTYLRTLYTNIRATVKAIIEEDLNQNNKHAKLVKGLIIDMAMCYNAETEWRDKYYVPATVDEH
LKISARSSGCMHLVSQGFISMGDVATSETLEWASTYPKIVRAVCIIARLANDIMSYKREASNNTMVSTVQTCAKEYGTT
TVEQAIKIRELIEEAWMDVTHECLRQPQPKALLERAVNLARTMDFLYKDVVDGYTDSSRSIKGILDSLYVHLID.