|           | 10                                       | 20            |               | 30                   | 40                       | 50 60   |      |
|-----------|--|---------------|---------------|----------------------|--------------------------|---|------|
| GmTLP1    | MALFSNHRHSSEA                            | LLGFLLFL      | VFRGVSV       | .SGATFTFV            | NKCDYT <mark>V</mark> WP | GILGTPELGSTGFELA  | 61   |
| GmTLP2    | MASSTQ                                   | FSTTI VTI CI  | FOFTAGS       | VSTTETIV             | NKCSVTVWP                | CII SCACTSPI PTTCEVIC   | 59   |
| GmTLP3    | MSAGNSVFHLISIYRMAFVVTMATTCNVRMGNSTKMASIV | TMACIEE       | TOTLEGE       | COTDITIT             | MECCATAND                | ATI CATCCCDI CTCCEVI C  | 90   |
|           |  |               |               |                      |                          |   |      |
| GmTLP4    |  | .SLSIFLFIA    | MSCVAAA       | . QAANFNII           | NNCIYIVWA                | AAVPGGGARLIN  | 49   |
| GmTLP5    | MVYLALCS                                 | LLTL          | ALSLATT       | .HAANFEIV            | NNCPYT <mark>v</mark> wa | AASPGGGRR <mark>L</mark> D                                      | 47   |
| GmTLP6    | MSPFFSHHPLSPQR                           |               |               |                      |                          |   |      |
| GmTLP7    | MPIVMTSWK                                | FPSFLAIL      | FTLLSYS       | .FSSIFTIT            | NNCSHT <b>I</b> WP       | <mark>G</mark> TLSGAGSPPLPTT <mark>G</mark> FR <mark>L</mark> D | 60   |
| GmTLP8    |  |               |               |                      |                          |   |      |
| GmTLP9    | MDQLTL                                   | AFPFIITI      | HIIVSGV       | VATTSETIV            | MKCDYTVWP                | CTI SNACTATI PTTCFVI C  | 58   |
| GmTLP10   | MPYR                                     |               |               |                      |                          | O LEDWACIATED TO VEG  | 4    |
|           | MA                                       |               |               | VOATT MITT           | WK CVT                   | TI GAADGI DUUTDGI VI C  |      |
| GmTLP11   |  |               | LGFLSGS       | .YGAILMVI            | NKCSYI WWP               | AILSAARSLPVHIPGLVL  | 52   |
| GmTLP12   |  |               |               |                      | <u></u> <u></u> .        | <u>.</u>  | 6    |
| GmTLP13   | MASPHIALLK                               | .LTSIFIVI     | .ASAPELCG     | .SSKTFTLV            | NYCKET <mark>IW</mark> P | <mark>G</mark> ITRSDNHSGD <mark>G</mark> FA <mark>L</mark> K    | 59   |
| GmTLP14   | MAALRK                                   | AVFFVIA       | QCFTFSA       | .YAARFEIT            | NRCTYT <mark>V</mark> WA | ASVPVGGGVQLN  | 49   |
| GmTLP15   |  | ALFFVTA       | LCCTSAA       | .YAARFDIT            | NRCSYP <mark>V</mark> WA | AAVPGGGRRLN   | 47   |
| GmTLP16   | MPTLRP                                   |               |               |                      |                          |   |      |
| GmTLP17   | MDHLTL.                                  |               |               |                      |                          |   |      |
| GmTLP18   |  | AI I I DD I D | · · IILLADG V | METVI                | ZAMUVIT                  | CTI CCCCCDDI ATTCEDI P  | 30   |
|           |  |               |               | VAAVED TO            | MEGTGEWID                | TUANG TOTTOFRE  | 50   |
| GmTLP19   |  | .SLFIILAIF    | LGFLSGS       | . YAAKFRIS           | NECISI <mark>V</mark> WP | AIVANSISIIGFSLL   | 53   |
| GmTLP20   | MARTS                                    |               |               |                      |                          |   |      |
| GmTLP21   | MAMMGSMLLSLRP                            | FLTTIFFL      | .TLTVLKVS     | .ASSSVIFY            | NKCQHP <mark>V</mark> WP | <mark>G</mark> IQPSAGKPVLARG <mark>G</mark> FK <mark>L</mark> A | 65   |
| GmTLP22   | MANPHIALLK                               |               |               |                      |                          |   |      |
| GmTLP23   | MSQLHG                                   |               |               |                      | <u> </u>                 | LADSGQGLN   | 15   |
| GmTLP24   | MV                                       |               |               |                      |                          |   | 2    |
| GmTLP25   |  | TESTCISE      | SILEVVA       | OCATVTET             | NNCOYT <b>V</b> WP       | CTI TCDONPOI STTCFFI A  | 55   |
| GmTLP26   | MTSNMONSPLHLONENYTYPTMTTT.               | BAVICILE.     | VELEAVV       | ECAKAGEM<br>ECAKAGEM | MKCTVTVWD                | CTI TCDOKDOI STTCEELC   | 76   |
| GmTLP27   |  |               |               |                      |                          |   |      |
|           |  |               |               |                      |                          |   |      |
| GmTLP28   |  | IFSLCLSF      | SLLFYVA       | . QGATVIFI           | NNCQYTVWP                | GTLTGDUNPULSTTGFELA   | . 55 |
| GmTLP29   | MAPTMQNSPLHLQNEKYPTMTTT.                 | RVAICLCF      | AFLYYAA       | .EGAKVSFN            | NKCSYT <mark>V</mark> WP | <mark>G</mark> TLTGDQKPQLSTT <mark>G</mark> FE <mark>L</mark> G | 74   |
| GmTLP30   |  | .SLNLFLPML    | LLCAGVQ       | .SDVTFTFE            | NLCSDT <mark>L</mark> WR | ASNPSIGDLEPD <mark>L</mark> P                                   | 52   |
| GmTLP31   |  | .SLSIFLFIA    | ISCVAVA       | .QAANFNIT            | NNCTYT <mark>V</mark> WA | AAVPGGGSRLN   | 49   |
| GmTLP32   | MASSAQ                                   | FSTTLVVLCL    | FQFLAGS       | .YSTTFTIV            | NKCSYT <mark>V</mark> WP | GILTGAGTSPLSTTGFVLC   | 59   |
| GmTLP33   | MALFSHHRHSYET                            | LLGFLLFL      | VFREV         | .SGATFTFV            | NKCDHT <mark>V</mark> WP | GIIGAPQLGSTGFELV  | 59   |
| GmTLP34   | MGRLLLF                                  | ITTISIVA      | FSFISDV       | OSASEKIV             | NKCBHTTWP                | CI I SCATSPPI PTTCFTI F   | 58   |
| GmTLP35   |  |               |               | . denoi ni v         |                          |   | 0    |
| GmTLP36   | MARSL                                    | EI CCIITEA    | VTTECCDT      | CATTITIV             | MNZNVT <b>V</b>          | TODMACHDVI ACCCITUD   | . 58 |
|           | MATEGO                                   |               | . VIIIFSSPI   | .GALILILV            | NINGIN I I V WP          | A LUPNAGRIP V LAGGGLILA   | 56   |
| GmTLP37   | MATSS                                    |               |               |                      |                          |   |      |
| GmTLP38   | MGRLLLF                                  |               |               |                      |                          |   |      |
| GmTLP39   | MGLLASS.                                 |               |               |                      |                          |   |      |
| GmTLP40   |  |               |               |                      |                          |   |      |
| GmTLP41   | MA                                       | LKQLSLLV      | MLFSLGV       | .DATVLTLQ            | NKCRDI <mark>V</mark> WP | <mark>G</mark> ILTGAGRPELIDD <mark>G</mark> AE <mark>L</mark> T | 53   |
| GmTLP42   | MALHLQT                                  | ILPFILVLAI    | ILSGORTCD     | .SARVFTIV            | NYCKET <mark>L</mark> WP | AVTPGESFNGGGFVLK  | 59   |
| GmTLP43   | MALIPNSKT                                | SALFHLLL      | FTI.GNVA      | YATVETI.E            | NHCSYT <mark>V</mark> WP | GTI.SGNGAATIGDGGFPMA  | 60   |
| GmTLP44   | MALIPNSKT                                |               |               |                      |                          |   | 60   |
| GmTLP45   | MALIPNSKT                                |               |               |                      |                          |   |      |
|           |  |               |               |                      |                          |   |      |
| GmTLP46   | MALHLQT                                  |               |               |                      |                          |   |      |
| GmTLP47   | MPIVMTSWK                                |               |               |                      |                          |   |      |
| GmTLP48   | MDRPTLSS                                 |               |               |                      |                          |   |      |
| GmTLP49   |  |               |               |                      |                          |   |      |
| GmTLP50   | MSPYHGHSSQR                              | FFIFLLIL      | AFRGVRV       | .TATTFTFV            | NKCDHT <mark>VW</mark> P | GVLGKPDIGTT <mark>G</mark> FE <mark>L</mark> K                  | 59   |
| GmTLP51   | MSASTHDQ                                 |               |               |                      |                          |   |      |
| GmTLP52   | MALIPNSKT                                |               |               |                      |                          |   |      |
| GmTLP53   | MDRPTL.S                                 | LESELLTI      | HISTSGV       | ISTTETIV             | NKCEYTUND                | GILSNACVPPI STTCEVIC  | 58   |
| GmTLP54   | MDSMAPHC                                 |               |               |                      |                          |   |      |
| GmTLP54   | MQLPKYI                                  |               |               |                      |                          |   |      |
| GIIITLESS | riųLPKYI                                 | . MADEFULICSS | TTIIGVALA     | ייול אוע בו א עו     | MUCLLLIMA                | TALMOGULATHOGOLA TO   | . 63 |
|           |  |               |               |                      |                          |   |      |

|          | 70                                    | 80   | 90   | 100                                    | 110                  | 12                       | 0                                  | 130 140   | 0   |     |
|----------|---------------------------------------|--|--|--|----------------------|--------------------------|------------------------------------|---|---|-----|
| GmTLP1   | KGGSRSFOAPTG.                         | WSGRFWGRTGCQFD. C  | L. GKGTCATAD                               | CGSGEVNC                               | NGAGA                | SPPATLAEF                | TLGTGSM                            | YYDVSLVDG <mark>Y</mark> NL                               | PMMVVAR                                   | 147 |
| GmTLP2   | PGESNVIAVPAA.                         | WSGRLWGRTFCTQD.A   | T. GKFSCVTGD                               | CGSSAVEC                               | NGGGA                | APPATLAEF                | TINGAG.GL                          | FFDVSLVDGYNL  | PMIVEPQ                                   | 146 |
| GmTLP3   |                                       | WSGRLWGRTLCSLDIT   |  |  |                      |                          |                                    |   |   | 178 |
| GmTLP4   | PCESWNISVINGT                         | TRGRIWGRTNCTFD. N  | A GRCKCI TCD                               | CE CVIEC                               | NKTC                 | TPPNTVVDE                | ATMOV N NI                         | EADISI ADGENA   | PIOITPT                                   | 135 |
| GmTLP5   | RCOTWNI WVNPCT                        | TAM <mark>ARIWGRT</mark> GCNFD.C                               | S CRCRCOTCD                                | CT GGING                               | OCWC                 | VDDNTI AFE               | VINOE G NO                         | EAD IST ADGENT  | DWDEADI                                   | 133 |
| GmTLP6   | DECTRICATE                            | WSGRFWGRTGCQFD.  | COCTCATCD                                  | CCSCOVIC                               | MCACA                | TDDATIAGE                | TTCEC AD                           | A A D A S T A D C A W T                                   | DMMMDAG                                   | 148 |
| GmTLP7   | COMMETTEELS.                          | WSGRIWARTGCTFD.  | T CICKCOTCD                                |  | DCMCA                | ADDTOLERI                | TICKC N DO                         | VADAGMADGAMI  | DI I WODD                                 | 146 |
|          |                                       |  |  |  |                      |                          |                                    |   |   |     |
| GmTLP8   | SUDEVVLEVPEG.                         | WSGRIWGRQGCCFDT  | II. GKGICEIGD                              | GGGGKTEG                               | GGMGG                | VPPAIL VEM               | GIP USALI                          | IF Y D V S L V D G F N L                                  | PVSMKPV                                   | 144 |
| GmTLP9   | IGESKIVIAPIS.                         | WGGRFWGRTLCTQD.S   | SA.GKFSGLSGD                               | CGSGKLEC                               | AGN <mark>A</mark> A | IPPAILAEF                | DGAG.GL                            | FFDVSLVDGYNV  | PMLVAPQ                                   | 145 |
| GmTLP10  | · · · · · · · · · · · · · · · · · · · | WSGRIWARTGCNFD.  | <u>.</u> S                                 | CA.GGMKC                               | TG.GG                | VPPVTLAEF                | TIGTGS.AQL                         | FYDVSLVDG <mark>Y</mark> NV                               | AMGVRAT                                   | 53  |
| GmTLP11  | P <mark>G</mark> DTNTIQVPPT.          | WSGRLWGRTLCSYN. 1  | T.GSFTCVTGD                                | CISST <mark>V</mark> EC                | GG . <u>.</u> A      | TAPVILAEF                | TLNGTG.GL                          | )   | PLV <mark>V</mark> EPR                    | 137 |
| GmTLP12  | N <mark>A</mark> KQV                  | <u>.</u> . <u></u> . <u></u> <u>.</u>                          | <u>.</u> <u>.</u> . <u></u> .              | T <mark>G</mark> .GRLEC                | EGNGS                | APPT <mark>S</mark> LFEI | TLGAGN.GQL                         | OYYDVS <mark>M</mark> VDG <mark>N</mark> NL               | P <mark>L</mark> LALPR                    | 60  |
| GmTLP13  | S <mark>G</mark> QSTVYTAPDG.          | WSGRIWARTGCNFD. N  | IN.GNGKCQTGG                               | C <mark>G</mark> .TS <mark>I</mark> NC | TVPG                 | SPPATSA <mark>D</mark> F | 'T <mark>L</mark> GEPI             | FYDVSLVDG <mark>F</mark> N <mark>L</mark>                 | PIVVKAI                                   | 141 |
| GmTLP14  | PGQSWSVDVPAGT                         | KG <mark>arvwart</mark> genfd.c                                | S.GRGGCQTGD                                | CG.GVLDC                               | KAYG                 | APPNILAEY                | GLNGFN.NL                          | OFFDISLVDGFNV   | PMDFSPT                                   | 135 |
| GmTLP15  |                                       | KG <mark>arvwart</mark> gcnfd.o                                |  |  |                      |                          |                                    |   |   | 133 |
| GmTLP16  | PNRAYSLQLPAL.                         | WSGRFWGRHGCAFD. V  | G. GRGRCATGD                               | C <mark>G</mark> .GS <mark>L</mark> FC | NGIGG                | SPPATLAEL                | . <mark>TL</mark> GNEQI            | FYDVSLVDG <mark>Y</mark> N <mark>L</mark>                 | PIS <mark>I</mark> TPF                    | 141 |
| GmTLP17  | TAESKTVTAPTS.                         | WGGRFWGRTLCTQD.S   | SA. GKFSCITGD                              | C <mark>G</mark> SGK <b>L</b> EC       | AGSGA                | TPPATLAEF                | TLDGAG.GLI                         | FFDVSLVDG <mark>Y</mark> NV                               | A <mark>M</mark> L <mark>V</mark> APQ     | 145 |
| GmTLP18  |                                       | WSGRIWARTGCTFD. A  |  |  |                      |                          | TLGAGN.GQI                         |   |   | 116 |
| GmTLP19  | PGKSYMVTMPTG.                         | .EVKVWARTFCSYS. ]  | A.GNFSCFTGD                                | CSSSNIDC                               | RGGNANP              | KPPVTLAEF                | TLNGSY.GL                          | YYKVSLREGENL  | PMSTEPG                                   | 141 |
| GmTLP20  |                                       | WSGRIWGRQGCSFD. N  |  |  |                      |                          |                                    |   |   | 143 |
| GmTLP21  | PNRAYSLOLPAL                          | WSGRFWGRHGCSFD.  | G GRGRCATGD                                | CC GSTFC                               | NGTGG                |                          | TLGNEQ                             |   |   | 149 |
| GmTLP22  | SCOTTVYTAPDG                          | WSCRIWARTCOFF N  | IN GNGKCKTGG                               | CG TSINC                               | TGPG                 | SPPATSADE                | TIGE P                             | FYDVSLVDGENL  | PTVVKAT                                   | 142 |
| GmTLP23  | AOOTHM                                | WSGRIWARTGCDFD.N   | CNSI VPI                                   | VTVDOVK                                | HASWW                | NPPVSIVE                 | ASN K GO                           | EVWVSI VDGENI   | PVIVTPO                                   | 78  |
| GmTLP24  | Αψφιιιι                               | RHGCIFD.S  | S CSCTCITCD                                | CC CKIKC                               | DCVCC                | ADDASTAFF                | TLDSQEG                            | EADASI ADCANT   | DVGTFDT                                   | 68  |
| GmTLP25  | DECTNOVNIDED                          | WSGRFWARTGCSNN   | CCETCDTCD                                  | CARCOVEC                               | MCACA                | TDDATI VET               | TVAPNG.GQ                          | EADAGMADCEWA  | DWGTTDO                                   | 140 |
| GmTLP26  | LAGINOVNILOL.                         | WSGRFWARTGCSNN   | N CRECOATAD                                | CASCOVAC                               | NCACA                |                          | TVAANG.GQI                         |   |   | 162 |
|          |                                       |  |  |  |                      |                          |                                    |   |   |     |
| GmTLP27  |                                       | WSGRFWARTGCSNN   |  |  |                      |                          |                                    |   |   | 141 |
| GmTLP28  |                                       | WSGRFWARTGCSNN   |  |  |                      |                          |                                    |   |   | 140 |
| GmTLP29  | PGASNSVDLPSP.                         | WSGRFWARTGCSNN   | N. GRESCATAD                               | CASGUVAC                               | NGAGA                | TPPAILVEI                | AANG.GU                            | PYDVSNVDGFNV  | PMSVIPU                                   | 160 |
| GmTLP30  | RDIYEIFNMDDH.                         | YSGSIWVR <mark>T</mark> GCSTN.A                                | S.NYFSCETGD                                | GNGIMEC                                | AGLNP                | SELTIALNE                | NNN                                | SYEVSLKHGQNM  | LVKIKPN                                   | 137 |
| GmTLP31  |                                       | TGGRIW <mark>G</mark> R <mark>T</mark> NCTFD.N                 |  |  |                      |                          |                                    |   |   | 135 |
| GmTLP32  | PGESNVITVPAA.                         | WSGRLWGRTVCTQD.A   | T.GKFSCVTGD                                | CGSSA <mark>V</mark> EC                | NGAGA                | APPATLAEF                | TLNGAG.GL                          | OFFDVSLVDG <mark>Y</mark> NL                              | PMI <mark>V</mark> EPQ                    | 146 |
| GmTLP33  | K <mark>G</mark> GSKSFQAPTG.          | WSGRFWGRTGCQFD.C   | S.GKGTCAT <mark>A</mark> D                 | C <mark>G</mark> SGE <mark>V</mark> NC | NGAGA                | SPPATLAEF                | TLGNGSMI                           | YYDVSLVDG <mark>Y</mark> N <mark>L</mark>                 | PMMVVAR                                   | 145 |
| GmTLP34  | SGKSRIVKIPKS.                         | WSGRIWARTLCGQD.S   | SD.GKFSCA <mark>TA</mark> D                | C <mark>G</mark> SGK <mark>V</mark> QC | VG.GA                | KPPATLAEF                | T <mark>L</mark> NGAD.GL           | FYDVSLVDG <mark>Y</mark> N <mark>L</mark>                 | PMLIVAK                                   | 144 |
| GmTLP35  |                                       | <u>.</u> <u></u>   | <u>.</u> <u>.</u> . <u></u>                | <u></u> <u>.</u> . <u>.</u>            | . <u></u>            | . <u></u> .              | . <u>.</u>                         | . <u>.</u> . <u></u>                                      | <u></u> . <u>.</u>                        | 0   |
| GmTLP36  |                                       | I <mark>W</mark> SGRVW <mark>a</mark> r <mark>t</mark> GCays.( |  |  |                      |                          |                                    |   |   | 145 |
| GmTLP37  | S <mark>G</mark> EEIVIHVPKV.          | WSGRIWGRQGCCFDQN   | IT. <mark>A</mark> KGSCQTGD                | C <mark>G</mark> .GL <mark>L</mark> QC | NGIGG                | VPPATL VEM               | I <mark>TL</mark> GTSQSPLH         | I <mark>F</mark> YDVSLVDG <mark>F</mark> N <mark>L</mark> | P <mark>V</mark> S <mark>V</mark> KPVEGGG | 148 |
| GmTLP38  | S <mark>G</mark> KSRTIKIPKS.          | WSGRIWARTLCGQD.S   | SD.GKFSCAT <mark>A</mark> D                | C <mark>G</mark> SGK <mark>V</mark> EC | VG.GA                | KPPATLAEF                | TLNGAD.GL                          | FYDVSLVDG <mark>Y</mark> N <mark>L</mark>                 | PML <mark>I</mark> VAK                    | 144 |
| GmTLP39  |                                       | WSGRFWGRTGCTFD. (  |  |  |                      |                          |                                    |   |   | 144 |
| GmTLP40  | SLTHQSIPVPDAH                         | I <mark>W</mark> SGRVW <mark>A</mark> RTGCSYS.(                | IT. <mark>A</mark> .FSCA <mark>S</mark> GD | C <mark>G</mark> .GR <mark>L</mark> QC | NGAGG                | AAPATLAQI                | .E <mark>V</mark> HHGSNDY <i>I</i> | SY <mark>G</mark> VSLVDG <mark>F</mark> N <mark>V</mark>  | PMTFTPH                                   | 145 |
| GmTLP41  | PGKAINITAPKG.                         | WSGRIWGRRGCTFH.S   | SY. GSGTCITGD                              | C <mark>G</mark> .GK <mark>L</mark> KC | DGVGG                | APPASLAEF                | <mark>TL</mark> DSQEG(             | FYDVSLV <mark>NGY</mark> NL                               | PV <mark>S</mark> IFPT                    | 138 |
| GmTLP42  | PGESAVFTAPVS.                         | WSGRIWARTGCKFD.Q   | N. GNGNCQTGS                               | C <mark>G</mark> .TT <mark>L</mark> KC | S <mark>A</mark> SG  | KTPASLAEF                | TLAQPI                             | FYDVSLVDGFN <mark>V</mark>                                | PMA <mark>V</mark> KPI                    | 141 |
| GmTLP43  | PGSSVQLTAPSG.                         | WSGRLWPRTGCNFD. A  | S. GNGKCLTGY                               | CA.GGMRC                               | TG.GG                | <b>VPPATLAEF</b>         | T <mark>I</mark> GŜG.GKI           | FYDVSLVDG <mark>Y</mark> N <mark>V</mark>                 | G <mark>V</mark> GVRAT                    | 144 |
| GmTLP44  |                                       | WSGRLWARTGCSFD.A   |  |  |                      |                          |                                    |   |   | 148 |
| GmTLP45  | PGSAVRLIAPAG.                         | WSGRFWARTGCRFD. C  | S.GAGKCTTGD                                | CA.GGLRC                               | TG.SG                | <b>VPPVTLAEF</b>         | TIGSASN.G.NK                       | FYDVSLVDG <mark>Y</mark> N <mark>V</mark>                 | GMGVRAA                                   | 147 |
| GmTLP46  | PGOSTVFTAPVS.                         | WSGRIWARTGCKFN. C  | N. GNGNCOTGS                               | CG.TTLKC                               | SASG                 | KTPASLAEF                | TLAQV                              | YYDVSLVDG <mark>F</mark> N <mark>V</mark>                 | PLFVKPI                                   | 141 |
| GmTLP47  | SGOMIKLTSVPG.                         | WSGRIWARTGCTFD. A  | T.GIGKCQTGD                                | CG.GKLEC                               | DGNGA                | APPTSLFEI                | TIGKGN.DQ                          | YYDVS <mark>M</mark> VDG <mark>Y</mark> NL                | PLLAQPR                                   | 146 |
| GmTLP48  | TGASTTIAAPAS.                         | WGGRFWGRTLCSED.S<br>WSGRFWAKSQCSTD.T                           | ST. GKFSCVTGD                              | CGSGKLEC                               | SGNGA                | APPATLAEF                | TLDGAG.GL                          | FFDVSLVDGYNV  | PMLVVPQ                                   | 146 |
| GmTLP49  | SKASSTIDVTAP.                         | WSGREWAKSOCSTD. T  | Y.GKFTCATGD                                | CGSGQVPC                               | NGNGG                | TPPVSI.VEF               | TLASNK.GO                          | FYDVSLVDGENI.   | PVIVIAQ                                   | 142 |
| GmTLP50  | RGSTRSFDAPPG                          | WSGRFWGRTGCEFD.I   | S. GHATCATCD                               | CGSGOVNO                               | NGAGA                | TPPATTAEF                | TIGEG. AP                          | YYDVSI.VDG <mark>Y</mark> NI                              | PMMVDAS                                   | 145 |
| GmTI P51 | PGOSNAVAVPTA                          | WSGRLWGRTLCSTD.S   | SA CKESCUTCD                               | CCSSAVE                                | GGGGA                | VPPATI AEF               | TINGA G CI                         | FYDVSVVDGVNI  | PMIVEAR                                   | 148 |
| CmTI P52 | PCSAVOITAPAC                          | WSGRFWARTGCSFD.  | S CSCKCVTCD                                | CC SCIK                                | SC CC                | VPPATI AFF               | GSASN G NK                         | FYDVSI VDGVNU   | GMGVBAT                                   | 147 |
| GmTLP53  | TCASTTIANDAS                          | WGGRFWGRTLCSQD.S   | TSCKESCVICD                                | CCSCKIF                                | SCNCA                | ΔΡΡΔΤΙΛΕΕ                | DCA C CI                           | FEDVSLVDGVNV  | DMI WVDU                                  | 146 |
|          | DUCCOUCERIDDO                         | WSGRFWGRTGCNFDPN   | T OOKT BOVIGD                              |  | NCCC A               | SDDVLLAR                 | TTCCC TO                           | EVDVSLVDGINV  | DI IMDDM                                  | 147 |
| GmTLP54  | DCOTORNIALLO.                         | WNGRIWARTGCNFA.S   | IN WKDACETCD                               | CD GBIVO                               | NCI TC               | TDDATIVE                 | CINCR ND                           | IFYDVSLVDG <mark>V</mark> NL                              | PVSVTPV                                   | 148 |
| GILLTE   | Tadicature.                           | *  | WILL HOETGD                                | D. GILL H                              | иопта                | *                        | Marr                               | * *   | DV II IV                                  | 140 |
|          |                                       | T  |  |  |                      | *                        |                                    | <b>Υ Υ</b>  |   |     |

|         | 150 160  | 170               | 180                               | 190 200  | 210                              | 220 230  |     |
|---------|--|-------------------|-----------------------------------|--|----------------------------------|--|-----|
| GmTLP1  | GGSGSCGATGCGED   | NRRCPSELR.        | . VEGGD ACQS                      | CGAFGKPEYCCNGAFSNPS                                | . TCKPSMYSQMFKST                 | CPKAYSYAYDDA <mark>T</mark> STF                | 230 |
| GmTLP2  | GGTGAGNCTATGCVVD   | N. TPCPAELM.      | .VMSSGEGVACKS                     | CEAFDDPQYCCSGAYATPD                                | .TCKPSSYSOFFKSA                  | CPRAYSYAYDDG <mark>S</mark> STF                | 234 |
| GmTLP3  | GGRNRRATGCEMD  | N LSCPTELK        | VI BOGD AVACKSS                   | SCO A  | EPCLTSOFFKTA                     | CPGAHVH  | 236 |
| GmTLP4  | AMGGGAKCVVD  | T CECPTOIO        | VDC CONNE                         | CTVFNTTQYCCSTGAA                                   | CCCPTDVCKEEKER                   | CDD AVSVDMDD ATSME                             | 210 |
|         | NGGCHKISCSAD   | T. GEOFIGE,       | ADC CONNE                         | CTURNITUTE COSTGAR                                 | CCCPTNYCEEVED                    | CUDGVCVDODDDDTCTE                              | 209 |
| GmTLP5  | NGGCHKISCSAD   | NGUCPGPLR.        | APGGUNNI                          | OT VEKINET COINGUG                                 | . SCGPINYSRFFKDR                 | CHUSYSYPUDDPISIF                               |     |
| GmTLP6  | GGSGA <mark>C</mark> AATG <mark>C</mark> GA <mark>D</mark> I | N RRCPAELR.       | . VDGGDACQS                       | CRAFGKPEFCCSGAFNSPA                                | . ACAPSMYSQIFKAA                 | CPKSYSYAFDDA <mark>T</mark> STF                | 231 |
| GmTLP7  | GVYGTGVCNATGCVTD   |                   |                                   |  |                                  |  | 238 |
| GmTLP8  | GGGIGCGVAACEA <mark>N</mark> I                               | NVCCPFALV.        | .MK.RQGKVVGCKS/                   | CL <mark>A</mark> AKSDR <mark>YCC</mark> TGEFATPK  | . SCKPSAFSRLFKTI                 | CPKAYSYAYDDS <mark>TG</mark> LK                | 230 |
| GmTLP9  | GGSGDNCTSTGCVGDI   | N. GACPSELR.      | .VTSVDGKOSVACKS/                  | CEAFGSPOYCCSGAYGSPT                                | .TCKPSPYSQIFKNA                  | CPRAYSYAYDDK <mark>T</mark> STF                | 234 |
| GmTLP10 |  | N GKCPTELO        | VTAEGGS TVACKS                    | CAAFNAAEECCTGEYSTPO                                | TCKPTOYSETEKSA                   | CPSAYSYAYDDA <mark>T</mark> STC                | 140 |
| GmTLP11 | SAIGAENCRATGCRR  |                   |                                   |  |                                  |  | 208 |
|         |  | N DCCDDELO        | . VIKEGEG VVGCKSK                 | CEATURDING CD N                                    | TOWN ON LING IN LINE IN LANGUAGE |  | 144 |
| GmTLP12 | GVIGGVCNAIGCIID.   | N RGCPRELQ.       | LLGGHQNQQ.VVGCKSI                 | CEAFKIDKICCKN                                      | . ICKPS YYSTIFKKA                | CPRSYSYAFDDGISTF                               |     |
| GmTLP13 |  | R. QNCPSELA.      | .SK.DNDKVIACRS                    | ICDVFNIDEYCCRGIYGNPA                               | . ICLASNYSKIFKUV                 | SPAAYS <mark>H</mark> ALDDPISLI                | 227 |
| GmTLP14 |  |                   |                                   |  |                                  |  | 210 |
| GmTLP15 |  |                   |                                   |  |                                  |  | 208 |
| GmTLP16 | KGSGKCSYAGCVSD   | NTMCPVGLQ.        | .VRSRDNKRVVACKS/                  | CSAFNSPK <mark>YCC</mark> TGSYGSPQ                 | . ACKPTVYSKIFKTA                 | CPKAYSYAYDDP <mark>T</mark> SIA                | 229 |
| GmTLP17 |  |                   |                                   |  |                                  |  | 234 |
| GmTLP18 |  | N BCCPKELO        | LIGGDONOB VVGCBS/                 | CEAEKTDKYCCSGAFANPN                                | TCOPSYVSTIFKKA                   | CPRAYSYAFDDG <mark>T</mark> STF                | 206 |
| GmTLP19 |  | N OKCDTKI K       | WIIDCCD VVCCOSI                   | COAHE  | TCKSTI ESEESKTK                  | CPEAYQHGSF                                     | 205 |
|         |  | N VCCDCALE        | WE DACK WACKSI                    | CLAMQSAK <mark>YCC</mark> TGSYSDPK                 | TCKDTI FALLERAT                  | CDKAVCVAVDDCCCI M                              | 229 |
| GmTLP20 | GGGVGGGVASCEVD   | NVUCPSALE.        | . VA . RNGA VVGCASI               | CLAMUSAK ICCIGS ISDPK                              | . ICKPILFAHLFKAI                 | SPKAYSYAYDDS SLN                               |     |
| GmTLP21 |  |                   |                                   |  |                                  |  | 237 |
| GmTLP22 |  | RQNCPSELA.        | .SK.DNDKVIACRS                    | CDVFNTDDYCCRGTYGNPA                                | .TCLASNYSKIFKQV                  | CPAAYS <mark>F</mark> ALDDP <mark>T</mark> SLI | 228 |
| GmTLP23 | GGLRGCNTTSSPSD   | /NKVYPPSFS.       | .VKGSDGSVIACKSA                   | CLALNQPEYCCTGPFASAD                                | .KCPPTPYSIIFKNM                  | CPQAYSYAYDDR <mark>T</mark> STF                | 165 |
| GmTLP24 | GGSRQCKAVTCRSD   | NRNCPSRLE.        | .VRNQSHIVGCKS/                    | CMAFNKPEYCCTGYFNSPK                                | .KCQPTSYSKVFKAS                  | CPQAYSYAYDDA <mark>T</mark> STF                | 154 |
| GmTLP25 |  | NDVNVCPSELQ.      | . VKGŠDGN VT <mark>AC</mark> NSA  | CVAENEDQYCCRGDYDTEE                                | TCPPTNYSQIFEEQ                   | CPDAYSYAYDDK <mark>S</mark> STF                | 229 |
| GmTLP26 |  | N SVCPAFIO        | VKGSDGN VIACKSA                   | CFAFKEDBYCCTGPNNTAF                                | TCPPTNYSOIFFFO                   | CPDAYSYAYDDK <mark>S</mark> STE                | 249 |
| GmTLP27 |  | N GVCDAELQ.       | WKCGDCN VIACKS                    | CEAEKEDDVCCTCDNNTAE                                | TCDDTMVCOTEEEO                   | CDDAVSVAVDDKGSTE                               | 228 |
|         |  | NOUNICO CELO.     | . V KGSDGN VIACKS                 | CHAPKEDRICCIGPINIAE                                | TOPPINIOQIFEEQ                   | CDD AVGVAVDD KOGTE                             |     |
| GmTLP28 |  |                   |                                   |  |                                  |  | 229 |
| GmTLP29 |  |                   |                                   |  |                                  |  | 247 |
| GmTLP30 |  | SSVCPLDLI.        | . AGNKNDQYV <mark>GC</mark> NSF   | PCDGYKDPKYCCNGN                                    | .GCQPDEISVK <mark>Y</mark> KGL   | CPFAH <mark>TY</mark> PGD <mark>N</mark> QPPIY | 224 |
| GmTLP31 | YNCSSVKCAAD  | IIGECPTQLQ.       | . VPGGCNNF                        | CTVFNTTQYCCSTGAA                                   | . GCGPTDYSKFFKER                 | CPDA <mark>F</mark> SYPMDDA <mark>S</mark> SMF | 210 |
| GmTLP32 |  | NTPCPTELK.        | .VMSSGDGVACKS/                    | CEAFDDPQ <mark>YCC</mark> SGAYATPD                 | .TCKP <mark>S</mark> SYSQFFKSA   | CPRAYSYAYDDG <mark>S</mark> STF                | 234 |
| GmTLP33 |  | N ORCPSELR.       | .VEGGDACQS/                       | CGAFGKPE <mark>YCC</mark> NGAFSNPS                 | . TCKP <mark>S</mark> MYSQMFKSV  | CPKAYSYAYDDA <mark>T</mark> STF                | 228 |
| GmTLP34 |  | N GGCPAFIB        | VARSNGSRG SVACRS                  | CFAFGDPRECCSFAYSTPD                                | MCGPSPYSI FEKHA                  | CPRAYSYAYDDKTSTY                               | 234 |
| GmTLP35 |  | IN G G OT ALLEIT. | . Whitehabita . B v Holton        | CDHI GDI IL CODERIBII D                            |                                  | CPRSYSYAYDDA <mark>T</mark> STF                | 21  |
|         |  | T ATCDUVIOU       | DWDAVIICD VVACVO                  | ACENTUTE I CODMIEMNDO                              | TOVOCTVOCERVIIA                  | VDATETE AUDTOOL MU                             | 234 |
| GmTLP36 |  |                   |                                   |  |                                  |  | 234 |
| GmTLP37 | GGGAGCGVAACEANI  | NVYCPSSLV.        | . VE. RNGK VVGCKSI                | ICLAAKSDRYCCIGEFAGGR                               | YYCKPIVFARLFKIV                  | SPNAYSYAYDDD <mark>SA</mark> VL                | 235 |
| GmTLP38 |  | NGGCPAELR.        | .VARSNGSRG.SVACRS                 | CEAFGDPRFCCSEAYSTPD                                | . TCGPSPYSLFFKHA                 | CPRAYSYAYDDK <mark>T</mark> STY                | 234 |
| GmTLP39 |  |                   |                                   |  |                                  |  | 227 |
| GmTLP40 |  |                   |                                   |  |                                  |  | 234 |
| GmTLP41 | GGSGQCKAVTCQSDI  | NRNCPNGLE.        | .VRNQSHIVGCKSA                    | ACM <mark>AF</mark> NKPE <mark>Y</mark> CCTGDFSNPR | . KCQP <mark>T</mark> SYSKVFKAS  | CPQAYSYAYDDA <mark>T</mark> STF                | 224 |
| GmTLP42 |  |                   |                                   |  |                                  |  | 227 |
| GmTLP43 |  |                   |                                   | ACAAFNTAEFCCTGDHSSPQ                               |                                  |  | 231 |
| GmTLP44 |  |                   | VKDGGGE VVACKS                    | CAAENTAFECCTCDHGGDO                                | TCSPTIVSKIEKMA                   | CPA AVSVAVDDPSTC                               | 235 |
| GmTLP45 |  | IN CYCDARIO       | WDDCACC WWACKS                    | CAAFNTAEFCCTGDHSTPQ                                |                                  |  | 234 |
|         |  |                   |                                   |  |                                  |  |     |
| GmTLP46 |  | K. AIGPNELA.      | . V N . V N G K I V G C K S I     | CONTROL OF CORGINY GNPS                            | . I CNPIFYSKKFKEA                | SPISYSYAYDDP <mark>I</mark> SIF                | 227 |
| GmTLP47 | GVYGTGV <mark>C</mark> NATG <mark>C</mark> VT <mark>D</mark> | NRGCPKELQ.        | . VVGGDGYQGGVV <mark>GC</mark> KS | ACE <mark>AF</mark> GSDQ <mark>Y</mark> CCSGEFANPT | .TCQP <mark>S</mark> YYSTLFKQA   | CPKAYSYAFDDA <mark>T</mark> STF                | 238 |
| GmTLP48 | GGSGEK <mark>C</mark> TATG <mark>C</mark> VGD <mark>I</mark> | NGACPSELK.        | . VMGADGREG. VVACKS/              | ACE <mark>AF</mark> NSPQ <mark>Y</mark> CCSGAYGTPD | .TCKP <mark>S</mark> AYSAIFKSA   | CPRAYSYAYDDK <mark>T</mark> STF                | 236 |
| GmTLP49 | GGLRG <mark>C</mark> NTTS <mark>C</mark> PSD                 | NKVCPPNFA.        | . VKGSDGSVIACKS!                  | RL <mark>A</mark> LDQPE <mark>YCC</mark> TGPFASAD  | .KCP <mark>PT</mark> PYSVIFKNQ   | CPQAYSYAYDDR <mark>T</mark> STF                | 229 |
| GmTLP50 | GGSGSCAATGCGAD   | NRRCPEELR.        | . VDGGDACQS                       | CR <mark>AF</mark> GKPE <mark>FCC</mark> SDAFNSPA  | . ACAPSMYSQIFKAA                 | CPKSYSYAYDDA <mark>T</mark> STF                | 228 |
| GmTLP51 |  | N. GGCPAFIK       | VKTVASGE GVACKS                   | CEAFGDPRYCCSGRYATPE                                | . TCKASSYSOLEKSA                 | CPRAYSYAYDDA <mark>T</mark> STY                | 235 |
| GmTLP52 |  | N GVCPAFIO        | VRDGSGA VVACKS                    | CLALNTAEY CCTGDHNTPQ                               | TCPPTHYSFIERMA                   | CPTAYSYAYDDAGSTC                               | 234 |
| GmTLP53 |  |                   | MMCCDCDEC WWACK                   | CEAFNSPQYCCSGAYGTPD                                | TCKGGGAGETEKGA                   | CDD AVGVAVDDVTQTE                              | 234 |
|         |  |                   |                                   |  |                                  |  |     |
| GmTLP54 |  | W. QUOPNELK.      | . VAUKSACKSI                      | ACEAFQSPEYCCSGAYASPA                               | . ICKPOVYSUIFKSA                 | SPASYSYAYDDA <mark>I</mark> SIF                | 230 |
| GmTLP55 | ITNPKCNIPGCLKD   | KGLCPPELE.        | .VLNSKGEIVACKS                    | CLAFDNDKFCCRNQYGSPG                                | . KCKP <mark>S</mark> VYSK1FKDA  | SPNY FSYAFDTPTPLV                              | 235 |
|         |  |                   |                                   |  |                                  | .1.  |     |

|         | 240   |     |
|---------|---|-----|
| GmTLP1  | .TCSG.AD.YT <mark>I</mark> TFCP                               | 243 |
| GmTLP2  | .TCAT.AD.YT <mark>I</mark> TFCP                               | 247 |
| GmTLP3  | .TCSS.HD.YT <mark>I</mark> TFCP                               | 249 |
| GmTLP4  | .TCMG.SD.YR <mark>V</mark> VFCP                               | 223 |
| GmTLP5  | .TCPAGSN.YK <mark>V</mark> VFCP                               | 223 |
| GmTLP6  | .TCTA.AD.YT <mark>V</mark> TFCP                               | 244 |
| GmTLP7  | .ICKA.FE.YDIIFCP  | 251 |
| GmTLP8  | .ICKA.HR.YV <mark>I</mark> TFCP                               | 243 |
| GmTLP9  | .TCASAAA.YTITFCP  | 248 |
| GmTLP10 | .TCSA.SD.YVITFCP  | 153 |
| GmTLP11 | .TCVS.AD.FTITFCP  | 221 |
| GmTLP12 | .TCKA.YE.YDIVFCP  | 157 |
| GmTLP13 | .TCSK.AN.FVVIFCG  | 240 |
| GmTLP14 | .TCNGGTD.YRVVFCP  | 224 |
| GmTLP15 | .TCKGGTN.YRVVFCP  | 222 |
| GmTLP16 | .TCTK.AN.YFLTFCP  | 242 |
| GmTLP17 | .TCASAAA.YTITFCP  | 248 |
| GmTLP18 | .TCKA.NE.YDIVFCP  | 219 |
| GmTLP19 | .TCAS.AQYYTVTFCP  | 219 |
| GmTLP20 | .RCRA.PR.YVITFCP  | 242 |
| GmTLP21 | .TCTK.AN.YLVTFCP  | 250 |
| GmTLP22 | .TCSK.AD.FVVIFCG  | 241 |
| GmTLP23 | .TCSGGAN.YSITFCP  | 179 |
| GmTLP24 | .TCQG.AN.YLIRFC.  | 166 |
| GmTLP25 | .TCFNGPD.YAIIFCP  | 243 |
| GmTLP26 | .TCSNRPD.YAITFCP  | 263 |
| GmTLP27 | .TCSNRPD.YA <mark>I</mark> TFCP                               | 242 |
| GmTLP28 | .TCFNGPD.YA <mark>I</mark> IFCP                               | 243 |
| GmTLP29 | .TCSNRPD.YA <mark>I</mark> TFCP                               | 261 |
| GmTLP30 | . QCKG. ADSYD <mark>I</mark> TFCP                             | 238 |
| GmTLP31 | .TCMGGSD.YR <mark>L</mark> VFCP                               | 224 |
| GmTLP32 | .TCAS.AD.YT <mark>I</mark> TFCP                               | 247 |
| GmTLP33 | .TCSG.AD.YT <mark>I</mark> TFCP                               | 241 |
| GmTLP34 | .TCAN.AN.YL <mark>I</mark> IFCP                               | 247 |
| GmTLP35 | .TCSG.VD.YT <mark>V</mark> TFCP                               | 34  |
| GmTLP36 | .QCSSPRE.LK <mark>V</mark> IFCH                               | 248 |
| GmTLP37 | K <mark>TC</mark> VA.PR.YV <mark>I</mark> TFCP                | 249 |
| GmTLP38 | .TCAN.AN.YL <mark>I</mark> IFCP                               | 247 |
| GmTLP39 | .TCSG.AD.YT <mark>V</mark> TFCP                               | 240 |
| GmTLP40 | .QCSSPRE.LK <mark>V</mark> IFCH                               | 248 |
| GmTLP41 | .TCQG.AN.YL <mark>I</mark> RFC.                               | 236 |
| GmTLP42 | .TCTG.TD.YV <mark>I</mark> AFCS                               | 240 |
| GmTLP43 | .TCSG.SD.YV <mark>I</mark> TFCP                               | 244 |
| GmTLP44 |   | 248 |
| GmTLP45 | .TCSG.SD.YR <mark>I</mark> TFCP                               | 247 |
| GmTLP46 | . <mark>TC</mark> SG.TD. YV <mark>V</mark> AFC <mark>A</mark> | 240 |
| GmTLP47 | . ICKA.FE.YD <mark>I</mark> VFCP                              | 251 |
| GmTLP48 | .TCAS.AD.YT <mark>I</mark> TFCP                               | 249 |
| GmTLP49 | .TCSGEQT.IPSHFAP  | 243 |
| GmTLP50 | .TCTA.AD.YTVTFCP  | 241 |
| GmTLP51 | .TCAS.AD.YRITFCP  | 248 |
| GmTLP52 | .TCSG.SD.YRITFCS  | 247 |
| GmTLP53 | .TCAS.AD.YT <mark>I</mark> TFCP                               | 249 |
| GmTLP54 | .TCTG.AD.YTTTFCP  | 243 |
| GmTLP55 | .SCAS.SE.FIITFCP  | 248 |

X non-conserved x similar x ≥ 80% conserved