

Accessory Publication

 α Q siRNA (*AtPLD α_1* amino acids 937–957)

At α_1 CCGGATGACGGTGGTA-----GCATA 21
At α_2 CCTGATGATGGTGGAA-----GCATT 17
At α_3 CCGGATGATGGTGGAA-----GCATT 18
Gh α CCAATGATGATGGTGGAA-----GTTTT 15
Ps α CCTGATGATGGAGGCA-----GCTTT 15
Osa α_1 CCTGATGACTCAGGCA-----GCATT 15
Osa α_2 CCCGACGCCGGCCGGA-----GCGCC 9
Zma α_1 CCTGATGATTCTGGCA-----GCTTT 14
At β_1 -----TGCTGGAAA-----GCGAC 4
At β_2 -----TGCTGGAAA-----GCGAC 4
Os β_2 -----TGCTGGAA-----ACGGC 4
At γ_1 -----TGGAGGAAAGGTCACAGCTTC 7
At γ_2 -----TGGGGGAAAGGTCACAGCTTC 8
At γ_3 -----TGGAGGAAAGGTCACAGCTTC 7
At δ -----TATGCCA-----GCAGT 6
Os δ -----TGCCAGCAGTAAGCTGAGCATT 5
At ϵ -----AAGATT-----GC-AC 4
At ζ_1 -----TGGGTGGCTTC--TATTAACGA 6
At ζ_2 -----TGGGTTAAGGCC----- 3

 β Q siRNA (*AtPLD β_1* amino acids 2879–2899)

At α_1 CAAGAGACTCTGAGATAGCAA 17
At α_2 CAAGAGACTCAGAGATAGCAA 18
At α_3 GTCGAGACACAGAGATCGCCA 15
Gh α CTAGAGACTCTGAAATAGCCA 14
Ps α CAAGAGATTCTGAAATTGCCA 17
Osa α_1 CTAGGGACTCTGAGATCGCCA 14
Osa α_2 AAGGAGACTCCGAGATCGCCA 14
Ta α CCCGGGACTCCGAGATTGCCA 14
Zma α_1 CCAGGGACTCCGAGATCGCCA 14
At β_1 CAAGAGATACAGAGATTGCAA 21
At β_2 CAAGAGATACAGAGATTGCAA 21
Ac β_2 CCAGAGATACTGAGATTGCCA 18
Ap β CCAGGGATACTGAGATTGCCA 17
Os β_2 TCAGAGATACTGAGATTGCGA 17
At γ_1 CTAGAGATACTGAAATCGCCA 16
At γ_2 CTAGAGACACTGAAATCGCTA 15
At γ_3 CTAGAGACACTGAAATCGCTA 15
At δ CCAAGATACCTGAAATCGCCA 15
Ap δ CAAGAGATACAGAGATTGCCA 20
Os δ CCAGAGATACTGAAATTGCCA 17
At ϵ GTCGGGACACCGAGATTGCAA 15
At ζ_1 CAAGAGATTCTGAGATTGGAG 17
At ζ_2 CACGAGACTCTGAGATCGGTG 13

γ Q siRNA (*AtPLD γ ₁* amino acids 1658-1678)

At α ₁ TCA-GACGTGCTAAGGATTTCA 14
At α ₂ TTA-GACGAGCTAAAGACTTCA 13
At α ₃ TAA-GGAGAGCCAAAACTTCA 12
Gh α TAC-GACGAGCAAAGAATTTTA 13
Ps α TTC-GAAGAGCAAAGGACTTCA 14
Ac α ₁ TAC-GCAGAGCGAAGAATTTTA 12
Ap α TAC-GCCGAGCGAAGAATTTTA 12
Os α ₁ TCC-GGAGGGCAAAGAACTTCA 13
Os α ₂ TTC-GCCGGCGCAAGAACTTCA 13
Zm α ₁ TAC-GGAGGGCGAAGAACTTCA 14
At β ₁ TTC-GTGCAGCCCAACATTTCA 14
At β ₂ TTC-GTGCAGCTCAACACTTCA 16
Ac β ₂ AACAAAGTTGTACAAAAAAGCA 10
Os β ₂ TTC-GTGGAGCGCAGCACTTCA 15
At γ ₁ TAC-GATCTGCTCAGCACTTCA 21
At γ ₂ TAC-GATCTGCTCAGCACTTCA 21
At γ ₃ TAC-GATCTGCTCAGCACTTCA 21
At δ TCA-GATCTGCTCAGCATTTCA 18
Os δ TCA-GATCCGCCAGCATTTCA 16
At ϵ TTC-GAAAGGCTGAGAGGTTTA 12
At ζ ₁ TTG-ACAAAAGCTGAACAATTTTA 10
At ζ ₂ TCC-AGAACGCTGAACAATTTTA 11

 δ Q1 siRNA (*AtPLD δ* amino acids 1645-1665)

At α ₁ AAAGATAACATCATTGATAGG 9
At α ₂ AAAGATAATATAATTGATAGG 10
At α ₃ AAAGATAACGTGATCGAAAGA 10
Gh α AAGGACAATATCATTGACCGA 9
Ps α AAAGACAACATCATCGACCGC 6
Ac α ₁ AAAGACAATATAATCGATCGA 9
Ap α AAAGACAATATAATCGATCGA 9
Os α ₁ AAGGATCAAATCATTGACAGG 10
Os α ₂ AAGAACAACACCATGACAGG 7
Zm α ₁ AAGGATCAAATCATCGACCGG 9
At β ₁ AAGAACGTGCTGATTGATATG 10
At β ₂ AAGAATGTGCTCATTGACATG 10
Os β ₂ AAGAATGTACTGATTGATATG 10
At γ ₁ AAGAATATTCTCATAGATATG 11
At γ ₂ AAGAACAATTCTCATAGATATG 11
At γ ₃ AAGAACAATTCTCATAGATATG 11
At δ AAGCGTCTTGTTGTAGATAAA 21
Ac δ AAGAATTTAGTAACCGACAAG 11
Os δ AAGGACCTGATAATTGATAAG 11
At ϵ AGAGGTTTGCCGGTAGAGAAG 10
At ζ ₁ GGTACAAGCCAAGTTGAAGAG 5
At ζ ₂ GGGACGAGGCCAGCTGAAGAT 4

δ Q2 siRNA (*AtPLD δ* amino acids 37-57)

<i>Atα₁</i>	CACGGGAC TTT ACATGCT ACC ATC	9
<i>Atα₂</i>	CATGGAA GATT GCACGCG GACA ATC	8
<i>Atα₃</i>	CATGGAA CTCTT-----	4
<i>Ghα</i>	CATGCTAC GAT CTAT -----	5
<i>Psα</i>	CATGGAA CTCT ACATGTT AC GATC	9
<i>Osaα₁</i>	CATGGGAC GCT GCACGCG CACC ATC	8
<i>Osaα₂</i>	CACGGC ACGCT CGAGGC CACC ATC	8
<i>Zmaα₁</i>	CACGGC ACGCT CCACGCG CACC ATC	10
<i>Atβ₁</i>	-----	0
<i>Atβ₂</i>	-----	0
<i>Acβ₂</i>	-----	0
<i>Apβ</i>	-----	0
<i>Osβ₂</i>	-----	0
<i>Atγ₁</i>	-----	0
<i>Atγ₂</i>	-----	0
<i>Atγ₃</i>	-----	0
<i>Atδ</i>	CACGGT---GACCTCGATTTGAAA	21
<i>Osδ</i>	GAGTCTCCGCCTCCGGCGAAGCCC	5
<i>Atϵ</i>	CATGGAA ACTA-----	4
<i>Atζ₁</i>	GGTGGAC GCT ACTTTCAGATGCAG	9
<i>Atζ₂</i>	GACGGAG TCAT CAG -----	5

ζ Q siRNA (*AtPLD ζ ₂* amino acids 298-318)

<i>Atα₁</i>	TATGAG TCG TTTCATATTTAC	7
<i>Atα₂</i>	TTTGAG TC TTTTCATATTTAC	7
<i>Atα₃</i>	-----	0
<i>Ghα</i>	AAAGGAC CGCA ACTGGGCCCCG	5
<i>Psα</i>	TACGAG TC TTTTCACATCTAC	9
<i>Osaα₁</i>	TATGAG TCG TTCCACATCTAT	8
<i>Osaα₂</i>	TACGAG GTG TTCCACATCTAC	8
<i>Taα</i>	-----	0
<i>Zmaα₁</i>	TACGAG TC TTCCACATCTAC	8
<i>Atβ₁</i>	ATGCAAC ATTTT TATGTTCCCT	10
<i>Atβ₂</i>	CAGCAG CA TTTTTATGTTCCCT	8
<i>Osβ₂</i>	ACGCAG AA TTTCCCTTGTCCCT	8
<i>Atγ₁</i>	ATGCA GC ATTTTGATGTACCG	8
<i>Atγ₂</i>	ATGCA GC ATTTTCGATGTACCC	8
<i>Atγ₃</i>	ATGCA GC ATTTTGATGTACCA	9
<i>Atδ</i>	-----	0
<i>Osδ</i>	GACGAGCG TT CGCCGTGCCC	8
<i>Atϵ</i>	-----	0
<i>Atζ₁</i>	AAGAAAC GT GCTTTTATTGAA	16
<i>Atζ₂</i>	AAGAAAC GT TTTGATCATTGAA	21

Fig. S1. Nucleotide sequence alignments of *Arabidopsis* phospholipase D siRNA targets with monocot and eudicot orthologs. Bold letters and numbers on the right indicate nt identities. *Ac*, *Allium cepa*; *Ap*, *Allium porrum*; *At*, *Arabidopsis thaliana*; *Gh*, *Gossypium hirsutum*; *Os*, *Oryza sativa*; *Ps*, *Papaver somniferum*; *Ta*, *Triticum aestivum*; *Zm*, *Zea mays*. Accession numbers are given in Fig. S2.

Ac α , *Allium cepa* *PLD* α ₁-like (TA6879_4679)
Ap α , *Allium porrum* *PLD* α -like (EU982404)
At α ₃, *Arabidopsis thaliana* *PLD* α ₃ (NM_122446)
At α ₁, *Arabidopsis thaliana* *PLD* α ₁ (NM_112443)
At α ₂, *Arabidopsis thaliana* *PLD* α ₂ (NM_104135)
Gh α , *Gossypium hirsutum* *PLD* α (EF378946)
Os α ₁, *Oryza sativa* *PLD* α ₁ (D73411)
Os α ₂, *Oryza sativa* *PLD* α ₂ (AB001919)
Ps α , *Papaver somniferum* *PLD* α (AF451980)
Ta α , *Triticum aestivum* *PLD* α (AJ880008)
Zm α ₁, *Zea mays* *PLD* α ₁ (D73410)

Ac β ₂, *Allium cepa* *PLD* β ₂-like (CF435023)
Ap β , *Allium porrum* *PLD* β -like (EU982403)
At β ₁, *Arabidopsis thaliana* *PLD* β ₁ (U84568)
At β ₂, *Arabidopsis thaliana* *PLD* β ₂ (NM_116245)
Os β ₂, *Oryza sativa* *PLD* β ₂ (AC078894)

At γ ₁, *Arabidopsis thaliana* *PLD* γ ₁ (NM_117255)
At γ ₂, *Arabidopsis thaliana* *PLD* γ ₂ (NM_179208)
At γ ₃, *Arabidopsis thaliana* *PLD* γ ₃ (NM_117254)

Ac δ , *Allium cepa* *PLD* δ -like (CF648854)
At δ , *Arabidopsis thaliana* *PLD* δ (AF322228)
Os δ *Oryza sativa* *PLD* δ (NM_001070361)

At ϵ , *Arabidopsis thaliana* *PLD* ϵ (NM_104391)

At ζ ₁, *Arabidopsis thaliana* *PLD* ζ ₁ (NM_112553)
At ζ ₂, *Arabidopsis thaliana* *PLD* ζ ₂ (NM_111436)

Fig. S2. Codes, names and accession numbers for species used for nucleotide sequence alignments of phospholipase D isotypes in Figs 2–5.

Allium cepa

Ac α , *Allium cepa* PLD α (TA6879_4679); Ac β , *Allium cepa* PLD β (CF435023); Ac δ , *Allium cepa* PLD δ (CF448854);

Allium porrum

Ap α , *Allium porrum* PLD α (ACM16811); Ap β , *Allium porrum* PLD β (ACM16810); Ap δ , *Allium porrum* PLD δ (ACM16809);

Arabidopsis thaliana

At α 1, *Arabidopsis thaliana* PLD α 1 (NP_188194); At α 2, *Arabidopsis thaliana* PLD α 2 (NP_175666); At α 3, *Arabidopsis thaliana* PLD α 3 (NP_197919); At ϵ , *Arabidopsis thaliana* PLD ϵ (NP_175914); At β 1, *Arabidopsis thaliana* PLD β 1 (NP_565963); At β 2, *Arabidopsis thaliana* PLD β 2 (NP_567160); At γ 1, *Arabidopsis thaliana* PLD γ 1 (NP_192922); At γ 2, *Arabidopsis thaliana* PLD γ 2 (NP_849539); At γ 3, *Arabidopsis thaliana* PLD γ 3 (NP_192921); At δ , *Arabidopsis thaliana* PLD δ (NP_567989); At ζ 1, *Arabidopsis thaliana* PLD ζ 1 (NP_188302); At ζ 2, *Arabidopsis thaliana* PLD ζ 2 (NP_187214);

Medicago truncatula

Mt α 1, *Medicago truncatula* PLD α 1 (ABD28731); Mt α 2, *Medicago truncatula* PLD α 2 (ABE88474); Mt α 3, *Medicago truncatula* PLD α 3 (ABE88470); Mt ϵ , *Medicago truncatula* PLD ϵ (AC167330); Mt β 1, *Medicago truncatula* PLD β 1 (AC169173); Mt β 2, *Medicago truncatula* PLD β 2 (CU326392); Mt β 3, *Medicago truncatula* PLD β 3 (CU326392); Mt δ 1, *Medicago truncatula* PLD δ 1 (CR962130); Mt δ 3, *Medicago truncatula* PLD δ 3 (CU013531); Mtv, *Medicago truncatula* PLDv (AC153000); Mt ζ 1, *Medicago truncatula* PLD ζ 1 (AC166898);

Oryza sativa

Os α 1, *Oryza sativa* PLD α 1 (NP_001042153); Os α 2, *Oryza sativa* PLD α 2 (NP_001054773); Os β 1, *Oryza sativa* PLD β 1 (NP_001065108); Os β 2, *Oryza sativa* PLD β 2 (NP_001048777); Os δ , *Oryza sativa* PLD δ (NP_001063826); Os η 1, *Oryza sativa* PLD η 1 (NP_001058017); Os η 2, *Oryza sativa* PLD η 2 (NP_001058015); Os η 3, *Oryza sativa* PLD η 3 (NP_001058016); Os κ , *Oryza sativa* PLD κ (NP_001045707); Os λ , *Oryza sativa* PLD λ (NP_001063203);

Os μ *Oryza sativa* PLD μ (NP_001050281), Os θ *Oryza sativa* PLD θ (NP_001061758); Osv1 *Oryza sativa* PLDv1 (NP_001051849); Osv2 *Oryza sativa* PLDv2 (NP_001059324); Os ζ 1 *Oryza sativa* PLD ζ 1 (NP_001055304); Os ζ 2 *Oryza sativa* PLD ζ 2 (NP_001042854);

Populus trichocarpa

Pt α 1, *Populus trichocarpa* PLD α 1 (XP_002299756); Pt α 2, *Populus trichocarpa* PLD α 2 (XP_002308663); Pt α 3, *Populus trichocarpa* PLD α 3 (XP_002325225); Pt α 4, *Populus trichocarpa* PLD α 4 (XP_002327529); Pt ϵ , *Populus trichocarpa* PLD ϵ (XP_002327429); Pt β 1, *Populus trichocarpa* PLD β 1 (XP_002301317); Pt β 2, *Populus trichocarpa* PLD β 2 (XP_00232008); Pt β 3, *Populus trichocarpa* PLD β 3 (XP_002299568); Pt δ 1, *Populus trichocarpa* PLD δ 1 (XP_002306960); Pt δ 2, *Populus trichocarpa* PLD δ 2 (XP_002331748); Pt δ 3, *Populus trichocarpa* PLD δ 3 (XP_002327624); Pt δ 4, *Populus trichocarpa* PLD δ 4 (XP_002301946); Ptv, *Populus trichocarpa* PLDv (XP_002304050); Pt ζ 1, *Populus trichocarpa* PLD ζ 1 (XP_002319499); Pt ζ 2, *Populus trichocarpa* PLD ζ 2 (XP_002328619);

Sorghum bicolor

Sb α 1, *Sorghum bicolor* PLD α 1 (142971); Sb α 2, *Sorghum bicolor* PLD α 2 (4893392); Sb α 3, *Sorghum bicolor* PLD α 3 (129832); Sb β 1, *Sorghum bicolor* PLD β 1 (123654); Sb β 2, *Sorghum bicolor* PLD β 2 (4780883); Sb δ , *Sorghum bicolor* PLD δ (5051569); Sb η 1, *Sorghum bicolor* PLD η 1 (130625); Sb η 2, *Sorghum bicolor* PLD η 2 (4886858); Sb κ , *Sorghum bicolor* PLD κ (5037418); Sb λ , *Sorghum bicolor* PLD λ (4970832); Sb μ , *Sorghum bicolor* PLD μ (4743947); Sb ν , *Sorghum bicolor* PLD ν (5033008); Sb ζ 1, *Sorghum bicolor* PLD ζ 1 (4853720); Sb ζ 2, *Sorghum bicolor* PLD ζ 2 (125502).

Fig. S3. Codes, names and accession numbers for species used for molecular phylogenetic analysis of phospholipase D protein sequences in Fig. 6.

Amino acids 1-50

AtPLD α 1 SQR----QGCKVSLYQDAWEDIFDAISNAKHLIYITGWSVIALVRTIGEL
 AtPLD α 2 SQR----RGCKVSLYQDAWEDIFDAITNAKHLIYITGWSVISLVRTVGEL
 AtPLD β 1 PLR----KGGTVRLYQDAWHDMFDAIRQARRLIYITGWSVVKLIRTLGEL
 AtPLD β 2 PLR----EGGSVTLYQDAWHDMFHAICQARRLIYITGWSVVRVLRRLGEL
 AtPLD γ 1 PLR----KGGRTVLYQDAWEDMADAIRQARRLIYITGWSVVRVLRVTLGEL
 AtPLD γ 2 PLR----KGGRTVLYQDAWEDMADAIRRARRLIYITGWSVVRVLRVTLGEL
 AtPLD γ 3 PLR----KGGRTVLYQDAWEDMADAIRRARRLIYITGWSVVRVLRVTLGEL
 AtPLD δ PVR----KGSQVRLYQDAWEDICYAISEAHHMIYIVGWSI IKLVRTLIGEL
 AtPLD ϵ PQR----SNCRVLYQDAWEDVYKAIESARHLVYIAGWALLVLRVTVGEL
 AtPLD α 3 NQR----EGCKVTLYQDAWEEIFDAIWEAKHLIYIAGWSVVTLVKRKLGEL
 OsPLD α 1 SQR----QGCKVTLYQDAWEDIFDAISNAQHLYITGWSVITLVRTLIGEL
 OsPLD α 2 SQR----QGCKVTLYQDAWEDIFDAINGARHFIYITGWSVIALIRTLGEL
 OsPLD η 1 SQR----PGCRVTLYQDAWEDVFDASNAKHLIYLTGWSVITLIRTLGEL
 OsPLD η 2 KQH----TGCGVTLYQDAWEDVFDASNAKHLIYITGWSVITLIRTLGEL
 OsPLD η 3 KQH----AGCRVTLYQDAWEDVFDASNAKHLIYITGWSVITLIRTLGEL
 OsPLD ν 1 PLR----RGGGVTLYQDAWEDICHSIVEAHHLYVMVGWSIVKLVRTLIGEL
 OsPLD ν 2 PLR----KGGMVTLYQDAWEDICHAIAEAHHLIYIIGWSLVKLVRTLGGEL
 OsPLD β 1 PMR----RGNRVTLYQDAWRDIYDAICQARRLIYIVGWSVIHLIRSLGEL
 OsPLD β 2 PLR----RGGRTVLYQDAWHDVYDAISQARRLIYITGWSVIQLVRS LGDL
 OsPLD λ PER----SGCRVTLYQNSWEDMYVAIRDARRFVYVAGWSVITLVRTLIGEL
 OsPLD μ PQR----RGCRVRLYQDAWEDVFDAISSARRMVYVAGWSVVVLRVTLGEL
 OsPLD δ PLR----HGGRTVLYQDAWEDICHAILEAHHMIYIVGWSVVRVLRNLGEL
 OsPLD κ --A----SGVQ-KLYQDAWEDMCLAVLGAQHLYVYVAGWSVVRVLRSLGAL
 OsPLD θ GQH----RGCKVTLYQDAWEDVFDAPANARSLVYIAGWSVVALVRTLGHL
 PtPLD α 1 SQR----QGCRVSLYQDAWEDVFDAITNAKHLIYITGWSVISLVRTLGEL
 PtPLD α 2 NQR----QGCRVTLYQDAWEDIFDAISDAKHLIYITGWSVITLIRKLGEL
 PtPLD α 3 SQR----SGCRVTLYQDAWEDIFDAIFNAKHLIYITGWSVITLVRI LGEL
 PtPLD α 4 PQR----QGCKVSLYQDAWEDVFD SITNAKHFIYITGWSVISLVRTLGEL
 PtPLD β 1 PLR----KGGTVTLYQDAWQDIFDAIRQARRLIYITGWSVVALVRPLGDL
 PtPLD β 2 PLR----KGGTVTLYQDAWQDIFDAIRQARRLIYITGWSVVTLVRTLGD L
 PtPLD β 3 PIR----RGGKVTLYQDAWDDIFNAISQARRLIYITGWSVVKLVRTLGD L
 PtPLD δ 1 PVR----HGGNVTLYQDAWEDICHAIVEAHHLYIVGWSIVRVRNLGEL
 PtPLD δ 2 PVR----KGCHVTLYQDAWEDICYAISEAHHMIYIVGWSVIKLVRTLIGEL
 PtPLD δ 3 PVR----KGGHVTLYQDAWEDICYAISEAHHMIYIVGWSVIKLVRTLIGEL
 PtPLD δ 4 PVR----HGGNVTLYQDAWEDICHAIVEAHHLYIAGWSIVKLVVRNLGEL
 PtPLD ν PLR----KGSVTLYQDAWEDICHAILEAHNLIYIVGWSVVKLVRTLIGEL
 PtPLD ϵ PQR----SNCHVTLYQDAWEDVYKAVEGAKHLIYIAGWSFMVLRMLGEL

MtPLD α 1 TQR----QGCKVSLYQDSWEDVFDAITKARHLIYITGWSVISLVRTLGEEL
 MtPLD α 2 DQR----QGCSVTLYQDAWEDIYNAIMNAKVFIYITGWSVITLIRTLGEM
 MtPLD α 3 KQN----NGCQITLYQDAWEDTCNAINDAKHFICITGWSVITLIRTLGEL
 MtPLD β 1 PLR----KGGNVTLYQDAWHDIFDAISQARRLIYITGWSVVRILIRTLGDL
 MtPLD β 2 PLR----KGGKVTLYQDAWQDIFDAISQARRLVYIVGWSVVSLIRTLGDL
 MtPLD β 3 PLR----KGGKVTLYQDAWHDIFDAISQARRLVYIVGWSVVSLIRTLGDL
 MtPLD δ 1 PVR----KGSVRLYQDAWEDICYAITEAHMVLVLAGWSIVKLVRTLGDGL
 MtPLD ν PLR----RGGNVTLYQDAWEDICHAILEAHMIYIIGWSIVKLVRSLGEL
 MtPLD δ 3 PVR----RGSVTLYQDAWEDICHAILEAHHLVYIVGWSIVKLVRLGEL
 MtPLD ϵ PQR----SNCHVKLYHDAWEDVYKAIEGAKYLVYIAGWSFMVLVRKLGEL
 SbPLD α 1 SQR----QGCKVTLYQDAWEDIFDAISKAQHLYITGWSVITLVRTLGEEL
 SbPLD α 2 SQR----QGCKVTLYQDAWEDIFDAISKAQHLYITGWSVITLVRTLGEEL
 SbPLD α 3 SQR----QGCRVTLYQDAWEDIFDAISGAKHLYITGWSVITLLRTLGEEL
 SbPLD η 1 SQR----PGCRVTLYQDAWEDVFDASINARHLIYITGWSVITLLRTLGEEL
 SbPLD η 2 KQR----QGCRVTLYQDAWEDVYDAIDGARHFVYVITGWSVTTLVRTLGEEL
 SbPLD μ KQR----RGCRVRLYEDAVDVFDAINRARRMVYVAGWSVVVLVRLGEL
 SbPLD λ PER----TNCVTLYQNAWEDLFAAIRDARRFVYVAGWSVITLVRTLGEEL
 SbPLD ν PLR----KGGRTLYQDAWEDICHAIEAHHLIYIVGWSLVKLVRTIGEEL
 SbPLD β 1 PLR----RGMVTLYQDAWRDIYDAICQARKLIYIVGWSVIHLVRSLGDL
 SbPLD β 2 PLR----RGGKVTLYQDAWRDVYDAISQARRLIYITGWSVIHLVRLGDL
 SbPLD δ PLR----HGGQVTLYQDAWEDICHAILEAHMIYIVGWSVVRILIRTLGDL
 SbPLD κ PLR----RGCDVRLYQDAWEDLCMAVLCAQRLVYVAGWSVVQLLRSLGEL
 SbPLD ζ 1 PPRGLEDGSMVQWFVDGFKAIASSIEEAKSEIFITGWLLYLRRRLDAL
 SbPLD ζ 2 PPRGLTEDGSVVQWFIDGFDAIASSIEEAKSEIFITDWLLYLRRRLDSL
 AtPLD ζ 2 PPRGLTSDGSQAQWFVDGFEAIAFAIQNATSEIFMTGWLLYLKRRLDAL
 AtPLD ζ 1 PPRGLTDDGSQAQWFVDGFAAIAAAIENAKSEIFICGWVLYLRRRLDNL
 OsPLD ζ 2 PPRGLTEDGSVVQWFIDGFNAIASSIEEAKSEIFITDWLLYLRRRLDIL
 OsPLD ζ 1 PPRGLMPDGSVQWFIDGFQAIASSIEQAKSEIFITGWLLFLRRRLDAL
 PtPLD ζ 1 PPRGLAEDGSQAQWFVDGFEAIASAIENARSEIFITGWLLYLRRRLDSL
 PtPLD ζ 2 PPRGLSEDGSQAQWFVDGFEAIALSIEDAKSEIFICGWLLYLRRRLDSL
 MtPLD ζ 1 PIRGLTGDGSQAQWFVDGFEAIASIQDAKSEIFITGWLLYLRRRLDSL
 ApPLD α ???
 ApPLD β ???
 ApPLD δ ???
 AcPLD δ ???
 AcPLD β ???
 AcPLD α ???

AtPLD α 1 LKKKASEGVRVLLLVWDDCILCPMFTHHQKIVVVDVSFVGGIDLCDGRYD
 AtPLD α 2 LKKKASEGVKIVLLLVWDDCILCPMFTHHQKIVVVDVSFVGGDLCDGRYD
 AtPLD β 1 LRSKSQEGVRVLLLIWDDVLLCPIYTHHQKNVIVDIAFVGGDLCDGRYD
 AtPLD β 2 LRSKSQEGVRVLLLVWDDVLLCPIYTHHQKNLIVDVAFVGGDLCDGRYD
 AtPLD γ 1 LKVKSQEGVRVLLLVWDDVLLCPIYTHHQKTIVVDVAFVGGDLCDGRYD
 AtPLD γ 2 LKVKSQEGVRVLLLVWDDVLLCPIYTHHQKTMIVDVAFVGGDLCDGRYD
 AtPLD γ 3 LKVKSQEGVRVLLLVWDDVLLCPIYTHHQKTMIVDVAFVGGDLCDGRYD
 AtPLD δ LKYKSQEGVRVLLLVWDDCVLSPFTHHQKCVLVDTAFIGGLDLCDGRYD
 AtPLD ϵ LKRKSEEGVAVRMLWDDCRLCPAFAHHQKTITLDMSFLGGFDLCDGRYD
 AtPLD α 3 LKKKAENVTVLMVWDDCVLCPMFTHHQKTIVVDVSFLGGIDLCDGRYD
 OsPLD α 1 LKKKASEGVRVLMVWDDCVLCPMFTHHQKIVVVDVSFVGGDLCDGRYD
 OsPLD α 2 LKKKASEGVRVLMVWDDCVLCPMFTHHQKIVVVDVSFVGGDLCDGRYD
 OsPLD η 1 LKRKASEGVRVLLLVWDDCVLCPMITHHQKTIVVDVSFVGGDLCDGRYD
 OsPLD η 2 LKRKASEGVRVLMVWDDCVCAASTHHQKTIVVDVSFVGGDLCDGRYD
 OsPLD η 3 LKRKASEGVRVLMVWDDCVLCPATHHQKTIVVDVSFVGGDLCDGRYD
 OsPLD ν 1 LKKKAREGVRIVILLWDDCVLPLFTHHQKCVLVDTAFIGGLDLCDGRYD
 OsPLD ν 2 LKTKVQEGVRVIVLLWDDCVLAPLTHHQKCVLVDTAFIGGLDLCDGRYD
 OsPLD β 1 LKMKSQEGVRVLLLVWDDVLLCPIFTHHQKTIVILDIAFVGGDLCDGRYD
 OsPLD β 2 LKRKSQEGVRVLLLVWDDVLLCPIFTHHQKTIVILDVAFVGGDLCDGRYD
 OsPLD λ LRRKADEGVAVLMPWQDCFLCPEFTHHQKTIVTDVDFIGGIDLCDGRYD
 OsPLD μ LKRKAQEGVMVLLLVWDDCVLCPMFTHHQKTIVVDVSFLGGIDLCDGRYD
 OsPLD δ LKFKSQEGVRVCLLVWDDCVLSPFTHHQKCVLVDTAFIGGLDLCDGRYD
 OsPLD κ LKYKSQEGVRVCLLVWDDCVLSPMYTQHQQKCVLVDTAFIGGLDLAAGRYD
 OsPLD θ LKSKAGERVAVLLLVWDDCVVCPVFTTHHQKAVVADVAFLGGIDLCDGRYD
 PtPLD α 1 LKKKASEGVRVLMVWDDCILCPMFTHHQKIVVVDVSFVGGIDLCDGRYD
 PtPLD α 2 LKKKAEEGVTVLMVWDDCILCPMFTHHQKTIVVDVSFVGGIDLCDGRYD
 PtPLD α 3 LKKKANEGVRVLMVWDDCVLCPMFTHHQKTIVVDVSFVGGIDLCDGRYD
 PtPLD α 4 LKKKASEGVRVLLIIVWDDCVLCPMFTHHQKIVVVDVSFVGGIDLCDGRYD
 PtPLD β 1 LRSKSQEGVRVLLLVWDDVLLCPIYTHHQKTIVDIAFVGGDLCDGRYD
 PtPLD β 2 LRSKSQEGVRVLLLVWDDVLLCPIYTHHQKTIVDIAFVGGDLCDGRYD
 PtPLD β 3 LKTKSQEGVRVLLLVWDDVLLCPIYTHHQKTIVDIAFVGGDLCKGRYD
 PtPLD δ 1 LKYKSQEGVRVLLLVWDDCVLSPYTHHQKCVLVDTAFIGGLDLCDGRYD
 PtPLD δ 2 LRYKSEEGVRVLLLVWDDCVLAPIFTHHQKCVLVDTAFIGGIDLCDGRYD
 PtPLD δ 3 LKYKSEEGVRVLLLVWDDCVLAPMFTHHQKCVLVDTAFIGGIDLCDGRYD
 PtPLD δ 4 LKYKSQEGVRVLLLVWDDCVLSPYTHHQKCVLVDTAFIGGLDLCDGRYD
 PtPLD ν LKYKSEEGVRVMLLVWDDCVLAPLFSHHQKCVILDTSFIGGLDLCDGRYD
 PtPLD ϵ LKRKSEEGVAVRMLWDDCKLCPFAHHQKTITVDMSFVGGDLCDGRYD
 MtPLD α 1 LKKKAQEGVRVLMVWDDCVLCPMFTHHQKILVVDVSFVGGIDLCDGRYD

Amino acids 101-150

AtPLD α 1 TPFHSLFRTLDTPREPWH-DIHSRLEGPIAWDVLMYNFQRWSVQLFRSID
 AtPLD α 2 TPFHSLFRTLDTPREPWH-DIHCRLGPIAWDVLNFEQRWSVQLFRSID
 AtPLD β 1 TPQHPLFRTLQTPREPWH-DLHSKIDGPAAYDVLNFEERWLVIQFRSID
 AtPLD β 2 TPQHPLFRTLQTPREPWH-DLHSKIDGPAAYDVLNFEERWLVIQFRSID
 AtPLD γ 1 TPKHPLFRTLKTPREPWH-DLHSKIDGPAAYDVLNFEERWMVQVFRSID
 AtPLD γ 2 TPKHSLFRTLKTPREPWH-DLHSKIDGPAAYDVLNFEERWMVQVFRSID
 AtPLD γ 3 TPKHPLFRTLKTPREPWH-DLHSKIDGPAAYDVLNFEERWMVQVFRSID
 AtPLD δ TPEHRLHDLDTPRQPWH-DLHCRIDGPAAYDVLNFEQRWRVQIFRSID
 AtPLD ϵ TEEHSLFRTLGTREPWH-DCHVSVVGGAAWDVLKNFEQRWTVQVLRSID
 AtPLD α 3 TVEHPLFGLTNSPREPWH-DIHCKLDGPAAWDVLNFEQRWMVQVFRSID
 OsPLD α 1 TQYHSLFRTLDSPREPWH-DIHSRLEGPIAWDVLNFEQRWRVQLFRSID
 OsPLD α 2 TPFHSLFRTLGTREPWH-DIHCRLGPAAWDVLNFEQRWRVQLFRSID
 OsPLD η 1 TQFHSLFRTLDTPREPWH-DIHSKIEGPAAWDVLNFEQRWRVQLFRSID
 OsPLD η 2 TQSHSLFRTLDTAPREPWH-DIHSRLEGPAAWDVLNFEQRWRVQVFRSID
 OsPLD η 3 TQSHSLFRTLDTAPREPWH-DIHSKIEGPAAWDVLNFEQRWRVQLFRSID
 OsPLD ν 1 TPEHRLFKDLDTPRQPWH-DLHCKIEGPAAFDILTNTFEQRWRVQVFRSID
 OsPLD ν 2 TPEHRLFKDLDTPRQPWH-DLHCKIEGPAAYDILTNTFEQRWRVQIFRSID
 OsPLD β 1 TPSHPLFRSLQTPREPWH-DLHSKIDGPAAYDVLQNFQERWLVIQFRSID
 OsPLD β 2 TPTHPLFRSLQTPREPWH-DLHSKIDGPAAYDVLNFEERWLVIQFRSID
 OsPLD λ DENHTLFRDLDTPREPWH-DVHCRLEGRAAWDVLNFEQRWRVQVFRSID
 OsPLD μ TQEHPLFRTLDTPREPWH-DIHCVEGPAAWDVLNFEQRWRVQVFRSID
 OsPLD δ TPEHRLFKDLDTPRQPWH-DLHCRIDGPAAYDVLKNFEQRWRVQVFRSID
 OsPLD κ TPSHRLFADLGTPRQPWH-DMHCRVDGPAAYDVLNFEQRWRAQVFRSVD
 OsPLD θ TQEHPLFRTLDTAPREPWH-DVHCRIEGPAAWDVLNFEQRWRVQVFRSID
 PtPLD α 1 TPFHSLFRTLDTPREPWH-DIHSRLEGPIAWDVLNFEQRWKVQLFRSID
 PtPLD α 2 TQDHPLFKTLDSPREPWH-DIHCLEGPAAWDVLNFEQRWTVQVFRSID
 PtPLD α 3 TPSHPIFRTLDTPREPWH-DVHCRLEGSIAWDVLNFEQRWRVQLFRSID
 PtPLD α 4 TPFHSLFRTLDTPREPWH-DIHSRLEGPIAWDVLNFEQRWKVQLFRSID
 PtPLD β 1 NPDHSLFRTLQTPREPWH-DLHSRIDGPAAYDVLNFEEDRWMVQIFRSID
 PtPLD β 2 TPDHPLFRTLQNPREPWH-DLHSRIDGPAAYDVLNFEERWMVQIFRSID
 PtPLD β 3 TPQHPLFRTLQTPREPWH-DLHCQIDGPAAYDILTNTFEERWLVIQFRSID
 PtPLD δ 1 TPEHRLFQDLDTPRQPWH-DLHCKIEGPAAYDVLNFEQRWRVQVFRSID
 PtPLD δ 2 TPEHRLFHDLNTPRQPWH-DLHCKIDGPAAYDVLNFEQRWRVQIFRSID
 PtPLD δ 3 TPGHRLFHDLNTPRQPWH-DLHCKIDGPAAYDALINFEQRWRVQIFRSID
 PtPLD δ 4 TPEHRLFRLDTPRQPWH-DLHCKIEGPAAYDVLNFEQRWRVQVFRSID

AcPLD α ???

Amino acids 151-200

AtPLD α 1 GGAAAGDRSIQDAYIHAIRRAKDFIYVENQYFLGALHLIPKELSLKIVSK
 AtPLD α 2 GGAAAGDRSIQDAYIHAIRRAKDFIYIENQYFLGALHLIPKELSLKIVSK
 AtPLD β 1 SNSVKGDMSIHTAYVKAIRAAQHFIYIENQYFIGANNLIPMEIALKIAEK
 AtPLD β 2 SNSVKGDMSIHTAYVKAIRAAQHFIYIENQYFIGANNLIPMEIALKIADK
 AtPLD γ 1 SSSVKGDMSIHAAYVKAIRSAQHFIYIENQYFLGANNLIPMEIALKIANK
 AtPLD γ 2 STSVKGDMSIHAAYVKAIRSAQHFIYIENQYFLGANNLIPMEIALKIANK
 AtPLD γ 3 SSSVKGDMSIHAAYVKAIRSAQHFIYIENQYFLGANNLIPMEIALKIANK
 AtPLD δ SGSVKGDKSIQTAYIQTIRSAQHFIYIENQYFLGADNLIPMELALKIVSK
 AtPLD ϵ HISATEEKSVHDGYVAAIRKAERFIYIENQYFMGCTNLIPVEIALKIAAK
 AtPLD α 3 DGAVEGERSIQDAYVNAIRRAKNFIYIENQYFLGALQLIPKEISLKIVSK
 OsPLD α 1 GGAAFGRDSIQDAYIHAIRRAKNFIYIENQYFLGALHLIPKELALKVVSK
 OsPLD α 2 GGAAFGRDSIQDAYIAAIRRARSFIYIENQYFLGALHLIPKELSMKVVSK
 OsPLD η 1 GGACFGDRSIQDAYIHAIRRAKNFIYIENQYFLGALHLIPREISLKIVNK
 OsPLD η 2 GGACDGRERSIQDAYIHAIRRARDFIYIENQYFIGAVNLIPRELSLKIMSK
 OsPLD η 3 GGACDGRERSIQDAYIHAIRRARDFIYIENQYFIGAVNLIPRELSLKIVSK
 OsPLD ν 1 SGSVKGDKSIHSAYVKAIRSAQHFIYIENQYFIGADNLIPVELALKIASK
 OsPLD ν 2 SGSVKGDKSIHNAYVKAIRSAQHFIYIENQYFIGAENLIPIELAIKIARK
 OsPLD β 1 SNSAKGDMSIHTAYVNAIRGAQHFIYIENQYFIGANNLIPIEIALKIANK
 OsPLD β 2 SNSVKGDMSVQTAYVNAIRGAQHFIYIENQYFLGANNLIPIEIALKIANK
 OsPLD λ DASVVGDRSIQAGYVEAIRRARRFIYVENQYFLGCLNLVPVEIALKVAAK
 OsPLD μ GGAAAGERSIQDAYIHAIRRARDFIYVENQYFLGALHLVPRELSLKIASK
 OsPLD δ SGSLKGDKSIHTAYVRAIRSAQHFIYIENQYFLGADNLVPIELALKIASK
 OsPLD κ SGSVKGEQSIHTAYVRAIRSAKRFIYIENQYFIGAGNLVPMEIALKVASK
 OsPLD θ SRAVDRERSIQDGYIHAIRRAKYFIYIESQCFLGAPHTIPKELSLKLASK
 PtPLD α 1 GGAAFGRDSIQDAYINAIRRAKNFIYIENQYFLGALHLIPKELSLKIVSK
 PtPLD α 2 DGAVVGRDSIQDAYINAIRRAKNFIYIENQYFLGALHLIPKELSLKIVSK
 PtPLD α 3 GGAAFGRDSIQDAYINAIRRAKSFYIENQYFLGALHLIPKELSLKIVSK
 PtPLD α 4 GGAAFGRDSIQDAYVNAIRRAKNFIYIENQYFLGALHLIPKELSLKIVSK
 PtPLD β 1 SNSVKDDMSIHTAYVKAIRAAQHFIYIENQYFIGANNLIPMEIALKIANK
 PtPLD β 2 SNSVKDDMSIHTAYVMAIRAAQHFIYIENQYFIGANNLIPMEIALKIANK
 PtPLD β 3 SNSVKGDMSIHTAYVNAIRAAQHFIYIENQYFLGANNLIPMEIALKIADK

PtPLD δ 1 SGSLKGDKSIQTAYVQAIRLAQHFIYIENQYFLGAENLIPMELALKIASK
 PtPLD δ 2 SGSLKGDKSIQTAYIQAIRSAQHFIYIENQYFLGADNLIPMELALKIASK
 PtPLD δ 3 SGSLKGDKSIQTAYIQAIRSAQHFIYIENQYFLGADNLIPMELALKIVSK
 PtPLD δ 4 SGSLKGDKSIQTAYIQAIRSAQHFIYIENQYFLGAENLIPMELALKIASK
 PtPLD ν SGSVKGDKSIHTAYVKAIRSAQHFIYIENQYFLGADNLVPMELALKIASK
 PtPLD ϵ HVSAIHRSIHEAYVEAIRRAERFIYIENQYFIGCTNLPIEIALKVVNK
 MtPLD α 1 GGAAFGRDSIQDAYINAIRRAKNFIYIENQYFLGALHLIPKELSLKIVSK
 MtPLD α 2 GGAVSNDRSIQDAYINAIRRAKNFIYIENQYFLGALHLIPKELSLKIISK
 MtPLD α 3 AGAASGDRSIQDAYINAIRRAKNFIYIENQYFLGALHLIPKELSLKIVSK
 MtPLD β 1 SGSVKGDMSIHTAYVKAIRAAQHFIYIENQYFIGANNLIPMEIALKIAEK
 MtPLD β 2 SNSVKGDMSIHSAYVKAIRAAQKFIYIENQYFLGANNLIPMEIALKIANK
 MtPLD β 3 SNSVKGDMSIHSAYVKAIRAAQKFIYIENQYFLGANNLPIEIALKIANK
 MtPLD δ 1 SGSLKGEKSIQTGYIQAIRSAQHFIYIENQYFIGADNLIPMELALKIASK
 MtPLD ν SGSVKGDQSIHAAYVKAIRSAQRFVYIENQYFLGANHLIPMELALKIASK
 MtPLD δ 3 SGSLKGDKSIQTAYIHAIRSAQHFIYIENQYFIGADNLIPMELALKIVSK
 MtPLD ϵ HVSASQESSIHEAYVEAIRRADRFVYIENQYFIGCTNLPIEIALKVVNK
 SbPLD α 1 GGAAFGRDSIQDAYINAIRRAKNFIYIENQYFLGALHLIPKELSLKIVSK
 SbPLD α 2 GGAAFGRDSIQDAYINAIRRAKNFIYIENQYFLGALHLIPKELSLKIVSK
 SbPLD α 3 GGAAFGRDSIQDAYIHAIRRARFIYIENQYFLGALHLIPKELSMKVVSK
 SbPLD η 1 GGACYGDRSIQDAYIHAIRRAKHFYIENQYFLGALHLIPRELSLKIVSK
 SbPLD η 2 GSACSGDRSIQDAYIHAIRRAKRFYIENQYFLGALQLVPRELSLKIVSK
 SbPLD μ GGAAAGERSIQDAYIHAIRRARDFVYIENQYFLGALHLIPKELSLKIVSK
 SbPLD λ DASVVGDRSIQIGYVEAVRRARFIYIENQYFLGCLNLVPEIALKVAAK
 SbPLD ν SGSVKGDKSIHNAYVKAIRSAQHFIYIENQYFIGAENLPIELAIKIARK
 SbPLD β 1 SNSAKGDMSIHTAYVHAIRAAQHFIYIENQYFIGANNLPIEIALKIANK
 SbPLD β 2 SNSVKGDMSVHTAYVNAIRGAQHFIYIENQYFLGANNLPIEIALKIANK
 SbPLD δ SGSLKGDKSIHTAYVRAIRSAQHFIYIENQYFLGADNLPIELALKIASK
 SbPLD κ AGSVKREQSIHAAYVAAIRSAERFVYVENQYFIGACNLVPMELALKVASK
 SbPLD ζ 1 QWSA-GEFSIHNAFYSLIEKAEHFVYIENQFFISIKNRVLEALYRRILRA
 SbPLD ζ 2 PWSA-GEFSIHNAFYSLIEKAEHFVYIENQFFISIKNRVLEALYRRILRA
 AtPLD ζ 2 QWSA-GEDSIHRAYCSLIQNAEHFIYIENQFFISILNRVLEALYRRILKA
 AtPLD ζ 1 QWSA-GEESIHSAYRSLIDKAEHFYIENQFFISVKNRVLEALYKRILRA
 OsPLD ζ 2 PWSA-GEFSIHNAFYSLIEKAEHFVYIENQFFISIKNRVLEALYRRILRA
 OsPLD ζ 1 QWSA-GEFSIHNAFYSLIEKAEHFVYIENQFFISIKNRVLEALYRRILRA
 PtPLD ζ 1 QWST-GEESIHKAYCSLIEKAEHFYIENQFFISIQNRVLDIYKRVIQA
 PtPLD ζ 2 QWSA-GEESIHCAYCSLIEKAEHFVYIENQFLISIRNRVLEALYRRIMRA
 MtPLD ζ 1 QWSV-GEESIHTAYCSLIEKAKHFYIENQFFISIQNRILEAIYRRILKA
 ApPLD α ?????DRSIQDAYIHAIRRAKNFIYIENQSFLGALHLIPKELSLKTVSK

ApPLD β ?????????????????????????????????GANNLIPMEIALKIANK
ApPLD δ ?????????????????????????????????EADNLIPMEIALKVASK
AcPLD δ SGSVKGDKSIHAAYVNAIRAAEHFIYVENQYFIEADNLIPMEIALKVASK
AcPLD β ?????????????????????????????????GANNLIPMEIALKIANK
AcPLD α ??????DRSIQDAYIHAIRRAKNFIYIENQYFLGALHLIPKELSLKIVSK

Amino acids 201-250

AtPLD α 1 IEKGEKFRVYVVVPMWP--EGSGSVQAILDWQRRTMEMMYKDVIQALRA-
AtPLD α 2 IKAGEKFKVYVVVPMWP--EGSGSVQAILDWQRRTMEMMYKDVIKALRE-
AtPLD β 1 IRANERFAAYIVIPMWP--EGGAATQRILYWQHKTIQMMYETIYKALVE-
AtPLD β 2 IRAKERFAAYIVIPMWP--EGGAATQRILYWQHKTQMMYGTIYNALVE-
AtPLD γ 1 IRAREKFAAYIVIPMWP--EGSNPIQRILYWQHKTQMMYQTIYKALVE-
AtPLD γ 2 IRARENFAAYIVIPMWP--EGSKPIQRILYWQHKTQMMYQTIYKALLE-
AtPLD γ 3 IRAREKFAAYIVIPMWP--EGSNPIQRILYWQHKTQMMYQTIYKALVE-
AtPLD δ IRAKERFAVYVVIPLWP--EGSGPVQEILYWQSQTMQMMYDVIKELKA-
AtPLD ϵ IRARERFAVYIVIPMWP--EGSETVEEILHWTRETMSMMYQIIGEAIWE-
AtPLD α 3 IEAGERFSVYIVIPMWP--EGSASVQAILDWQRRTMEMMYTDIIIALRK-
OsPLD α 1 IEAGERFTVYVVVPMWP--EGSGSVQAILDWQRRTMEMMYTDITEALQA-
OsPLD α 2 IEAGERFTVYVVVPMWP--EGSGSVQAILDWQRRTMEMMYTDIAHAIQA-
OsPLD η 1 IEAGERFAVYVVLPMWP--EGSGSVQAILDWQRRTMEMMYDIAVALEA-
OsPLD η 2 IAAGERFTVYVVVPMWP--EGSQAMQAILDWQRRTMEMMYADIAGALKA-
OsPLD η 3 IAAGERFAVYVVVPMWP--EGNEAMQAILDWQRRTMEMMYDIAVALKA-
OsPLD ν 1 IKANEQFAVYIVLPMWP--EGAAPMQQILFWQQTMSMMYKIIADALRM-
OsPLD ν 2 IKARERFAAYIVIPMWP--EGTAAMQEILFWQQTMSMMYKIVAEALQK-
OsPLD β 1 IKAKERFSAYIVIPMWP--EGGAPTQRILYWQHKTQMMYETIYRALKE-
OsPLD β 2 IYANERFSAYIVIPMWP--EGGAPTQRILYWQKKTQMMYEVIIHKALKE-
OsPLD λ IRRGERFAAYVVTMWP--EGGDSVQAILRWNRLTVEMMYGIVTKAIDD-
OsPLD μ IAAGERFAVYVVVPMWP--EGSDSVQAILDWQRRTMEMMYRDVDAAIQA-
OsPLD δ IRAGERFAVYVVIPLWP--EGAASVQEILFFQ--TMEEMMYRIIAQELKA-
OsPLD κ IAAGERFAVYIVIPMWP--EGSGPIQEILFWQRTMQAMYEVIAAAIRA-
OsPLD θ IRSGDSFRVYVVLPMWP--EGSATVQAVLDWQRRTMEMMYKDVAALAA-
PtPLD α 1 IEAGERFTVYVVVPMWP--EGSASVQAILDWQRRTMDMMYKDVIQALRA-
PtPLD α 2 IEAGERFTVYIVIPMWP--EGSGSVQAILDWQRRTMDMMYSDITEALVK-
PtPLD α 3 IKAGERFSVYVVIPLWP--EGSGSVQAILNWQKRTMEMMYSDIAEALQD-

PtPLD ζ 1 YKENKCFRVIIVIP LSPGFQGAATVRAIMHWQYRTISRKKT S ILYNLNT-
 PtPLD ζ 2 FNDKKCFRVIIVIP L L P G F Q G A A S V R A I M H W Q Y R T I C R G Q N S I L H N L Y D -
 MtPLD ζ 1 HKEQEDFRVIVVLP L L P G F Q G A A T V R A L T H W Q Y R T I S R E R H S I L H N L D A -
 ApPLD α IEAGERFTVYVVVPMWP--EGSGSVQAILNWPRRTMEMMYTDIAEALQAK
 ApPLD β IKANERFSVYIVIPMWP--EGGAATQRILYWQKKTMMYETIYKALKEE
 ApPLD δ IRARKRFAYVYIIIPMWP--EGSAAQEIILFWQNQTMRMYKIIGQELRSM
 AcPLD δ IRARERFAYVYIIIPMWP--EGSPAVQEIILFWQNQTMMYKIVGQELRSM
 AcPLD β IKANERFSVYIVIPMWP--EGGAATQRILYWQKKTMMYETIYKALKEE
 AcPLD α IEAGERFTVYVVVPMWP--EGSGSVQAILDQWRRTMEMMYTDIAEALQAK

Amino acids 251-300

AtPLD α 1 GLE--EDPRNYLTF FCLGNIYVHTKMMIVDDEYIIIGSANINQRSMDGAR
 AtPLD α 2 GLEG-EDPRDYLTFFCLGNIYVHTKMMIVDDEYIIIGSANINQRSMDGAR
 AtPLD β 1 GLEGAFSPQDYLNFFCLGNVYVHSGMIVDDEYVIGSANINQRSMEGTR
 AtPLD β 2 GLEDEYSPQDYLNFFCLGNIYVHSGMIVDDEYVIGSANINQRSMEGTR
 AtPLD γ 1 GLDSQFEPQDFLNFFCLGTIYVHSGMIVDDEYVIGSANINQRSLEGTR
 AtPLD γ 2 GLDGQLEPQDFLNFFCLGNIYVHSGMIVDDEYVIGSANINQRSLEGTR
 AtPLD γ 3 GLDGQLEPQDFLNFFCLGTIYVHSGMIVDDEYVIGSANINQRSLEGTR
 AtPLD δ QSD--AHPDYLNFFYCLGKIYVHAKGMIVDDEYVLMGSANINQRSMAGTK
 AtPLD ϵ GDK--SHPRDYLNFFCLANVYVHSKLMIVDDTYILIGSANINQRSMDGCR
 AtPLD α 3 GLD--ANPRDYLTFFCLGNIYVHSGMIVDDEYIIIGSANINQRSMDGGR
 OsPLD α 1 GIE--ANPKDYLTFFCLGNIYVHTKMMIVDDEYIIIGSANINQRSMDGAR
 OsPLD α 2 GID--ADPKDYLTFFCLGNIYVHTKMMIVDDEYIIIGSANINQRSMDGAR
 OsPLD η 1 RIN--ADPRDYLTFFCLGNIYVHSGMIVDDEYIIIGSANINQRSMDGGR
 OsPLD η 2 RMD--ADPRDYLTFFCLGNIYVHSGMIVDDEYIIIGSANINQRSMDGGR
 OsPLD η 3 HSD--ADPRDYLTFFCLGNIYVHSGMIVDDEYIIIGSANINQRSMDGGR
 OsPLD ν 1 GLVE-AHPQDYLNFFYCLGKIYVHSGMIVDDEYVIGSANINQRSMEGCR
 OsPLD ν 2 GLDD-THPQDYLNFFYCLGKIYVHSGMIVDDEYVIGSANINQRSMDGSR
 OsPLD β 1 GLDDLYEPQDYLNFFCLGNVYVHSGMIVDDEYVIGSANINQRSMEGIR
 OsPLD β 2 GLDNITYEPQDYLNFFCLGNVYVHSGMIVDDEYVIGSANINQRSMEGTR
 OsPLD λ GLRGQAHPDYLNFFCLGNIYVHAKLMIVDDEYVMVGSANLNERSLAGNR
 OsPLD μ GIR--ADPTDYLNFFCLGNIYVHAKTMIVDDEYIIIGSANINQRSMDGGR
 OsPLD δ NIKN-AHPQDYLNFFYCLGKIYVHAKGMIVDDEYVILGSANINQRSLAGSR
 OsPLD κ GMEGAHPDYLNFFYCLGKIYVHSGMIVDDEYVIGSANINQRSLAGSR

OsPLD θ GST--QNPREYLSFFCLGNINVNANIMIVDDEYIIIVGSANVNQRSMDGGR
 PtPLD α 1 GLE--EDPRNYLTFFFCLGNIYVHAKMMIVDDEYIIIGSANINQRSMDGAR
 PtPLD α 2 GLN--TDPREYLAFFFCLGNIYVHAKMMIVDDEYIIIGSANINQRSMDGSR
 PtPLD α 3 GVE--ANPKDYLTFFCLGKIYIHAKMMIVDDEYIITGSANINQRSMDGGR
 PtPLD α 4 GLE--EDPRNYLTFFFCLGNIYVHAKMMIVDDEYIIIGSANINQRSMDGAR
 PtPLD β 1 GLEDAFSPQDFLNFCLGNIYVHAKMMIVDDEYVILGSANINQRSMEGTR
 PtPLD β 2 GLEEAFFSPQDFLNFCLGNIYVHAKMMIVDDEYVILGSANINQRSMEGTR
 PtPLD β 3 GLENTYEPQDYLNFFCLGNIYHSGKGIIVDDEYVILGSANINQRSMEGTR
 PtPLD δ 1 NLEN-SHPQDYLNIFYCLGNVYVHAKMMIVDDEYIILGSANINERSMAGSR
 PtPLD δ 2 DLVD-SHPQDYLNIFYCIGKIYVHAKMMIVDDEYVIVGSANINQRSMAGSK
 PtPLD δ 3 NLVD-SHPQDYLNIFYCIGKIYVHAKMMIVDDEYVIMGSANINQRSMAGSK
 PtPLD δ 4 NLEN-SHPQDYLNIFYCLGNIYVHAKMMIVDDEYVILGSANINQRSMAGSR
 PtPLD ν GLSYQYHPQYLNIFYCLGKIYVHAKMMIVDDEYVIMGSANINQRSLDGSR
 PtPLD ϵ GEP--GHPRDYLNFFCLANVYVHAKMMIVDDAYMLIGSANVNQRSMDGRR
 MtPLD α 1 GLE--ENPRDYLTFFCLGNIYVHAKMMIVDDEYIIIVGSANINQRSMDGAR
 MtPLD α 2 NIE--AHPRDYLTFFCLGKIYVHAKMMIVDDEYIIVGSANINQRSMDGGR
 MtPLD α 3 GIR--ANPRDYLSFYCLGNIYVHAKMMIVDDEYIIIGSANINQRSMDGAR
 MtPLD β 1 GLEAAFSVQDYLNFFCLGNIYVHAKMMIVDDEYVIVGSANINQRSMEGTR
 MtPLD β 2 GLENEYEPQDYLNFFCLGNIYVHAKMMIVDDEYVLMGSANINQRSMEGTR
 MtPLD β 3 GLANEYEPQDYLNFFCLGNIYVHAKMMIVDDEYVLMGSANINQRSMEGTR
 MtPLD δ 1 QLSD-VHPQDYLNIFYCLGNIYVHAKMMIVDDEYVILGSANINQRSMAGTK
 MtPLD ν GLSECYHPQDYLNIFYCLGKIYVHAKMMIVDDGYVIIGSANINQRSMDGSR
 MtPLD δ 3 KLND-SHPQDYLNIFYCLGNIYVHAKMMIVDDEYAMVGSANINQRSLAGSR
 MtPLD ϵ GEP--GHPRDYLNFFCLANVYVHAKMMIVDDLILIGSANVNQRSMDGQR
 SbPLD α 1 GIE--ANPKEYLTFFFCLGNIYVHTKMMIVDDEYIIIGSANINQRSMDGAR
 SbPLD α 2 GIE--ANPKDYLTFFCLGNIYVHAKMMIVDDEYIIIVGSANINQRSMDGAR
 SbPLD α 3 GID--ANPRDYLTFFCLGNIYVHTKMMIVDDEYIIIVGSANINQRSMDGAR
 SbPLD η 1 RID--RNPRDYLTFFCLGNIYVHAKMMIVDDEYIIIVGSANINQRSMDGGR
 SbPLD η 2 NID--ANPKDYLSFFCLGNIYVHAKMMIVDDEYIIIGSANINQRSMDGAR
 SbPLD μ GLQ--DDPRDYLTFFCLGNIYVHAKMMIVDDEYIIIVGSANINQRSMDGGR
 SbPLD λ GLRGQAHPDYLNFFCLGNIYVHAKMMIVDDEYVMVGSANLNERSLAGNR
 SbPLD ν GLHE-SHPQEYLNIFYCLGKIYVHAKMMIVDDEYVILGSANINQRSMDGSR
 SbPLD β 1 GLDDMYEPQDYLNFFCLGNVYVHAKMMIVDDEYVIVGSANINQRSMEGIR
 SbPLD β 2 GLDGYEPQDYLNFFCLGNVYVHAKMMIVDDEYVIIGSANINQRSMEGTR
 SbPLD δ DIKD-MQLEDYLNFFCLGNIYVHAKMMIVDDEYVILGSANINQRSLAGSR
 SbPLD κ GKSG--HPTDYLNIFYCLGNVYVHAKMMIVDDEYVIIGSANINQRSLAGSR
 SbPLD ζ 1 GPK----AHDYISFYGLRAIYVHAKMMIVDDRITLIGSANINDRSLGSR
 SbPLD ζ 2 GSK----AHDYISFYGLRAIYVHAKMMIIDDRMALIGSANINDRSLGSR

AtPLD ζ 2 GPK----TQDYISFYGLRSIYVHVKLMIVDDRIAVIGSSNINDRSLGSR
 AtPLD ζ 1 GVK----AHDYISFYGLRAVYVHVKIMIVDDRAALIGSANINDRSLGSR
 OsPLD ζ 2 GSK----AHDYISFYGLRAIYVHVKLMIIDDRMTLIGSANINDRSLGSR
 OsPLD ζ 1 GPK----AHDYISFYGLRAIYVHVKLMIIDDRITLIGSANINDRSLGSR
 PtPLD ζ 1 GPK----THDYISFCGLRTVYVHVKVMIVDDRIAYIGSSNINDRSLGSR
 PtPLD ζ 2 GPK----THDYISFYGLRSVYVHVKIMIIDDRITLIGSANINDRSLGSR
 MtPLD ζ 1 GRK----THDYISFYGLRSVYVHVKLMIIDDRVALIGSSNINDRSLGSR
 ApPLD α GLE--DNPQDYLTFFCLGN????????????????????????????????
 ApPLD β GLEDIYGPQDYLNFFCLGNIYVHVKGMIVDDEYVIVGSANINQRSLEGTR
 ApPLD δ NMEE-AHPQDYLNFFCLGNIYVHVKGMIVDDEFVLLGSANINQRSMDGSR
 AcPLD δ NMEG-AHPHEYLNFYCLGNIYVHVKGMIVDDEVLLGSANINQRSMDGIR
 AcPLD β GLEDIYGPQDYLNFFCLGNIYVHVKGMIVDDEYVIVGSANINQRSLEGTR
 AcPLD α GLE--DNPQDYLTFFCLGNIYVHAKMMIVDDEKIVGSANINQR??????

Amino acids 301-350

AtPLD α 1 DSEIAMGGYQPHH-LSHR----QPARGQI-HGFRMSLWYEHL-GFLDPSS
 AtPLD α 2 DSEIAMGGYQPYH-LSTR----QPARGQI-HGFRMSLWYEHL-GFLDPSS
 AtPLD β 1 DTEIAMGAYQPQH-TWAR--KHSGPRGQI-YGYRMSLWAEHM-AFTQPES
 AtPLD β 2 DTEIAMGAYQPQH-TWAR--RQSGPRGQI-YGYRMSLWAEHM-AFVEPES
 AtPLD γ 1 DTEIAMGGYQPHY-SWAM--KGSRPHGQI-FGYRMSLWAEHL-GFEOPEN
 AtPLD γ 2 DTEIAMGGYQPHH-SWAK--KGSRPRGQI-FGYRMSLWAEHL-GFEOPEN
 AtPLD γ 3 DTEIAMGGYQPHH-SWAK--KGSRPRGQI-FGYRMSLWAEHL-GFEOPEN
 AtPLD δ DTEIAMGAYQPNH-TWAH--KGRHPRGQV-YGYRMSLWAEHL-GFVEPSD
 AtPLD ϵ DTEIAIGCYQTNT-NNTN-----EI-QAYRLSLWYEHT-GSSEPES
 AtPLD α 3 DTEIAMGAYQPSHLLSTN--NMRPVGQI-FSFRISLWLEHL-RFQCPES
 OsPLD α 1 DSEIAMGGYQPYH-LATR----QPARGQI-HGFRMALWYEHL-GFQRPES
 OsPLD α 2 DSEIAMGAYQPHH-LAAA---GRPARGQV-HGFRMALWYEHL-GFQRPES
 OsPLD η 1 DSEIAMGAFQPCH-LNTK--GLVARGQI-HGFRMSLWYEHL-GFLNPES
 OsPLD η 2 DSEIAMGAFQPHH-LNIG--GQLARGQI-HGFRMSLWYEHLGGFLHPGS
 OsPLD η 3 DSEIAMGAFQPHH-LNVN--GQAARGQI-HGFRMSLWYEHL-GFVHPGS
 OsPLD ν 1 DTEIAMGAYQPHY-KWSADHDQGPPRGQV-YGYRMSLWAEHL-GFGRPET
 OsPLD ν 2 DTEIAMGAYQPHY-SWAG--RKKAPRGQV-YGYRMSLWAEHL-GFRWPHS
 OsPLD β 1 DTEIAMGAYQPQY-TWAS--KVSAPRGQI-YGYRMSLWAEHI-GFNYPET
 OsPLD β 2 DTEIAMGAYQPQY-TWAN--MLSAPRGQI-YGYRMSLWAEHI-GFSCPES

OsPLD λ DSEIAQGSYQPAH-LNGG--PSGRARGLV-HAFRMSLWHEHL-MFLEPES
 OsPLD μ DTEIAMGAYQPSH-LASV---NRPARGQV-HGFRLALWHEHL-GLLRPSS
 OsPLD δ DTEIAMGAYQPHH-TWST--KGGHPRGQV-YGYRTSLWAEHL-GFKDPSS
 OsPLD κ DTEIAVGAYQPNL-RAGA---G-AGDGQV-FGFRMLLWEEHL-GLRSPES
 OsPLD θ DTEMAMGAYQPRH-LDTP---NSWPRGQV-HQFRLALWREHL-GMIYPSR
 PtPLD α 1 DSEIAMGGYQPYH-LATR----QPARGQI-HGFRLGLWYEHL-GFLKPEN
 PtPLD α 2 DSEIAMGGYQPHH-LATS----QPARGQI-YGFRMALWYEHL-GFQHPES
 PtPLD α 3 DTEIAMGAYQPYH-LATN----QPARGQI-HGFRMSLWYEHL-GFCHPES
 PtPLD α 4 DSEIAMGGYQPYH-LATR----QPARGQI-HGFRLGLWYEHL-GFLHPES
 PtPLD β 1 DTEIAMGAYQPQH-TWAK--KQSNPLGQI-HGYRMSLWAEHT-GFTKPES
 PtPLD β 2 DTEIAMGAYQPQH-TWAR--KQSNPLGQI-HGYRMSLWAEHT-GFTKPES
 PtPLD β 3 DTEIAMGAYQPNH-TLAR--KNSRSHGQV-YGYRMSLWAEHI-GFEQPET
 PtPLD δ 1 DTEIAMGAYQPHH-TWSN--KKRHPLGQV-YGYRMSLWAEHL-GFKEPGS
 PtPLD δ 2 DTEIAMGSYQPHH-TWVT--KKKHPRGQV-YGYRMSLWREHL-GFMEDPN
 PtPLD δ 3 DTEIAMGSYQPRH-TWAA--RKKHPRGQI-YGYRMSLWVEHL-GFMEDPED
 PtPLD δ 4 DTEIAMGAYQPHH-TWSN--KKRHPLGQV-YGYRMSLWAEHL-GFKEPES
 PtPLD ν DTEIAMGAYQPTY-TWAR--KNSHPHGQV-YGYRMSLWAEHL-GFDEPQS
 PtPLD ϵ DTEIAIGCYQPKNGENTR-----NPRDIL--AYRMSLWYEHT-GFLEPES
 MtPLD α 1 DSEIAMGAYQPYH-LANR----QPARGQI-HGFRMSLWYEHL-GFLHPES
 MtPLD α 2 DTEIAMGAFQPHH-LASTSNGSQRPQGQV-YGFRRALWYEHI-GFDEPER
 MtPLD α 3 DSEIAIGAFQPNH-IATN---NRPPKGQI-YGFRRALWHEHL-GYENPES
 MtPLD β 1 DSEIAMGAYQPHH-TWAR--KHSNPLGQI-HGYRMSLWAEHT-GFLQPES
 MtPLD β 2 DTEIAMAAAYQPNH-TWAT--KKSHPHGQV-HGYRMSLWSEHI-GFKQPES
 MtPLD β 3 DTEIAMGAYQPNH-TWAT--KKSHPHGQV-HGYRMSLWSEHI-GFKQPES
 MtPLD δ 1 DTEIAMGSYQPHY-TWSA--RKKHHPGQI-YGYRMSLWAEHL-GFKEPER
 MtPLD ν DTEIAMGAYQPKH-TLAE--KNSLPRGQV-YGYRMSLWAEHL-GFTEPHT
 MtPLD δ 3 DTEIAMGAYQPHH-TWSK--KNGHPRGQV-YGYRMSLWAEHL-GFKEPES
 MtPLD ϵ DTEIAIGGYQSHQ-DGVD--HPISKGDI-HEYRMSMWYEHT-GFLEPES
 SbPLD α 1 DSEIAMGAYQPYH-LATR----QPARGQI-HGFRMSLWYEHL-GFQRPES
 SbPLD α 2 DSEIAMGAYQPYH-LATK----QPARGQI-HGFRMSLWYEHL-GFQHPES
 SbPLD α 3 DSEIAMGAYQPHH-LAAA---SRPARGQV-HGFRMSLWYEHL-GFTRPDS
 SbPLD η 1 DSEIAMGAYQPCH-LNTK--GQVARGQV-HGFRMSLWYEHL-GFLNPGS
 SbPLD η 2 DSEIAMGAYQPSH-LNTN--GDVARGQV-HGLRMSLWYEHL-GFKDPGS
 SbPLD μ DTEIAMGAYQPGY-LATR--NRPARGQV-HGFRVSLWQEHL-ALLRPSS
 SbPLD λ DSEIAQGSYQPAH-LNGP---CRRARGQV-HGFRMSLWHEHFIMFLEPES
 SbPLD ν DTEIAMGAYQPHY-SWAG--SGSPPKGQV-YGYRMSLWAEHL-GFRRPES
 SbPLD β 1 DTEIAMGAYQPQY-TWAN--KLSAPRGQI-YGYRMSLWAEHI-GFHYPES
 SbPLD β 2 DTEIAMAAAYQPQH-TWAN--TLSAPRGQI-FGYRMSLWAEHI-GFTMPES

SbPLD δ DTEIAMGAYQPHY-AWST--KNGHPHGQV-YGYRTSLWAEHL-GFKDPSS
 SbPLD κ DTEIAVGAYQPDH-TGAD---GDAPRGKV-HAYRMSLWEEHL-GVQRPES
 SbPLD ζ 1 DSEIADVIEDKEV-VNSKMDGRPWEAGKFSLSLRLSLWAEHL-GIMDPID
 SbPLD ζ 2 DSEIGVIEDKEV-VSSIMDGRPWEAGKFSLSLRLSLWAEHL-GIMDPVD
 AtPLD ζ 2 DSEIGVIEDKEF-VESSMNGMKWMAGKFSYSLRCSLWSEHL-GIEDPIK
 AtPLD ζ 1 DSEIGVLIEDTEL-VDSRMAGKPKWAGKFSSSLRSLWSEHL-GIIDPVS
 OsPLD ζ 2 DSEIGMIEDKEV-VSSIMDGRHWEAGKFSLSLRLSLWAEHL-GIMDPVD
 OsPLD ζ 1 DSEIADVIEDKEV-VSSKMNGKPWEAGKFSLSLRLSLWAEHL-GIMDPID
 PtPLD ζ 1 DSEIGIVTEDKEF-VESSMNGETWKAGKFAYSLRRSLWSEHL-GISDPVA
 PtPLD ζ 2 DSEIGVLIEDKEL-VDSLMGGKPRKAGKFTLSLRLSLWSEHL-GVIDPVI
 MtPLD ζ 1 DSEIGVIEDKEY-VESLMNGKPKWAGKFSHSLRCSLWSEHL-GIMDPVA
 ApPLD α ETKQQTGEYQPEQ??
 ApPLD β DTEIAMGA??
 ApPLD δ DTEIAMGA??
 AcPLD δ DTEIAMGAYPPHH-TWAG????????????????????????????????
 AcPLD β DTEIAMGAYQPEY-TWAR--KHSRPYQGI-YSYRMSLWAEHL-GFLEPEK
 AcPLD α ???

Amino acids 3561-380

AtPLD α 1 LECIEKV-NRISDKYWDFYLPG-HLLRYPI
 AtPLD α 2 QECIQKV-NRVADKYWDLYLPG-HLLRYPI
 AtPLD β 1 IECVRKV-RTMGERNWKQFMRG-HLLKYPV
 AtPLD β 2 LGCVRKV-RTVAEENWEQFMRG-HLMKYPV
 AtPLD γ 1 MECVRRV-RQLSELNWRQYMSG-HLLKYPV
 AtPLD γ 2 MECVRRV-RQLSELNWQYMSG-HLLKYPV
 AtPLD γ 3 MECVRRV-RQLSELNWRQYMPG-HLLKYPV
 AtPLD δ LECLKKV-NTISEENWKRFLQG-HLIKYPV
 AtPLD ϵ LECVRGL-RTIGEQMWEIYMLGIHLVAYPI
 AtPLD α 3 EECIRMV-NATADELWGLYLPG-HLLSYPI
 OsPLD α 1 LECVQKV-NRIAEKYWDMYLPG-HLLSYPI
 OsPLD α 2 LDCVRKV-NAMADRCWDLYLPG-HLLTYPV
 OsPLD η 1 LECVQRV-NKMADKYWDLYLPG-HLLTYPV
 OsPLD η 2 LECVRRV-NEMANKHWELYLPG-HLLTYPI
 OsPLD η 3 LECVRRV-NAMADRHQWLYLPG-HLLTYPV

OsPLD ν 1 GECVRRV-REMAEENWRAYTKG-HLMCYPL
 OsPLD ν 2 VECVRQV-NEMAEENWARYMRG-HLMRYPI
 OsPLD β 1 MECMRRV-RQIGEQNWERFMRG-HLMKYPV
 OsPLD β 2 LECTRQV-RHIGEQNWRQFMRG-HLVKYPV
 OsPLD λ AECVRAV-RRAAEATWDAYLLG-HLLPFPI
 OsPLD μ LASVRLV-NQAARRHWD AFLPG-HLMAYPV
 OsPLD δ LECVNYV-NEIAEENWRRFLQG-HLLKYPV
 OsPLD κ PECVKRV-NEIAEENWRRYMQG-HLMRYPV
 OsPLD θ HGCMSRV-NQAARQHWD MYLPG-HLMAYPV
 PtPLD α 1 EDCIRKV-NQIADKYWDLYLPG-HLLRYPI
 PtPLD α 2 VQCIQLV-NQVANENWEKYLMS-HLLRYPI
 PtPLD α 3 LECVRKV-NHIAEKNWQLY LPG-HLLAYPI
 PtPLD α 4 EECVTKV-NQITDKYWDLYLPG-HLLRYPI
 PtPLD β 1 LECVRRI-RTMGEMNWKQFMRG-HLLKYPV
 PtPLD β 2 LECVRRI-KAMGEMNWKQFMTG-HLLKYPV
 PtPLD β 3 IECVRRV-RSLGEQNWRQYMKG-HLLKYPV
 PtPLD δ 1 LDCVKSV-NKIAEDNWKFLQG-HLLKYPV
 PtPLD δ 2 LLCVKRV-NHTAEENWKKFLKG-HLLKYPL
 PtPLD δ 3 LHCVKKV-NKIAEDNWRKFLQG-HLLKYPL
 PtPLD δ 4 LDCVKSV-NKIAEDNWKFLQG-HLLKYPV
 PtPLD ν LECMKLV-NKTSRHNWKAYMRG-HLMQYPI
 PtPLD ϵ LACAQRM-RLVGEQMWNVYMEGVHLVNYPL
 MtPLD α 1 EECIRKV-NQIADKYWDLYLPG-HLLRYPI
 MtPLD α 2 LECVKLL-NRVAERNWKLYVNT-HLLRYPV
 MtPLD α 3 LDCINLV-NGFAKTNWDIYRSFHMLMQYPI
 MtPLD β 1 LECVRKV-RAIGEMNWKQFMRG-HLLKYPV
 MtPLD β 2 IECVRRI-RSLSEYNWRQYMNG-HLLKYPL
 MtPLD β 3 IECVRRI-RSLSEYNWRQYMNG-HLLKYPL
 MtPLD δ 1 LECVRKV-NEIADDNWRKYLQG-HLLKYPV
 MtPLD ν LECVRRV-NRIARKNWSIYMKG-NLMHYPV
 MtPLD δ 3 LECVESV-NKIAEDNWRKFLQG-HIMKYPI
 MtPLD ϵ LKCVQRM-CSIGDKMWKIYMEGVHLVTYPM
 SbPLD α 1 VECVQKV-NRIAEKYWDLYLPG-HLLMYPI
 SbPLD α 2 MECVQKV-NRIADKYWDLYLPG-HLLSYPI
 SbPLD α 3 VECIRKV-NAMADRYWDLYLPG-HLLTYPV
 SbPLD η 1 LECVQRV-NQMADKYWDLYLPG-HLLRYPV
 SbPLD η 2 LKCVRKV-NKRAEEFWKMYLHG-HLLSYPI
 SbPLD μ LACVRRM-NQVAKQHWD MF LPG-HLMAYPV

SbPLD λ LECVRRAV-RRAAERLWDAYLPG-HLLPFPI
 SbPLD ν EECVQLV-NQMADDNWSYMKG-HLMKYPV
 SbPLD β 1 LECMRRV-RHLGEENWKQFMRG-HLMKYPV
 SbPLD β 2 LECTRQV-RHIGQQNWEKFMKG-HLLKYPV
 SbPLD δ LECVRRV-NQIAVENWQRFLQG-HLLKYPV
 SbPLD κ PGCVKLV-NGI?????????G-HLMRYPV
 SbPLD ζ 1 DSTFKNIWMATAKTNTMIYVVG-HLVSFPL
 SbPLD ζ 2 DSVYKSIWMATAKENTMIYVVG-HIVCFPL
 AtPLD ζ 2 DATYKDLWMATAKNTDIYTRG-NLVCFPL
 AtPLD ζ 1 DSTYKEIWMATAKTNTMIYIKG-HLVSFPL
 OsPLD ζ 2 DLTYNNIWMGTAKANTKIYVVG-HLVSFPL
 OsPLD ζ 1 DSTFKNIWMATAKTNTMIYIKG-HLVSFPL
 PtPLD ζ 1 ETTYRDLWLATAKENSKIYVVG-HLVSFPL
 PtPLD ζ 2 DSTYKDIWMSTAKTNTMIYTRG-HLVSFPL
 MtPLD ζ 1 DSTYKELWAATAKENTRIYIKG-LLVSFPL
 ApPLD α ?????????????????????????????????
 ApPLD β ?????????????????????????????????
 ApPLD δ ?????????????????????????????????
 AcPLD δ ?????????????????????????????????
 AcPLD β PQCMRRI-RELGEYNWKQF????????????
 AcPLD α ?????????????????????????????????

Fig. S4. Alignment of phospholipase D amino acid sequences of species used for molecular phylogenetic analysis in Fig. 6. At, *Arabidopsis thaliana*; Os, *Oryza sativa*; Pt, *Populus trichocarpa*; Mt, *Medicago truncatula*; Sb, *Sorghum bicolor*; Ap, *Allium porrum*; Ac, *Allium cepa*.