

## Supplementary Information

### 1. PLASTID Desaturases

The alanine-phenylalanine dipeptide conserved in the ASAFAF motif of heterokont plastid bipartite targeting signal is highlighted in red. Serine residues known to be enriched in the -12/+12 flanking region of the ASAFAF motif are shown in green.

*PAD* (*Phatrdr\_9316*)

MLSTKLFWT**SSV**LASSAVL**A**FPTSPATRTPR**S**T**S**ILKVAVDPTTVTKKEYEDICGVSFADT  
LEDRLKATNYLYPKHVEVIDDIAPIAGAMVDEILLETGENAWQPQDYLPDLSQDNWHDSIKE  
VRAMAKEIPDELLVVLIGDMVTEEALPTYQTLLNTFEGCDDPTGTSESPWARWSRGWTSEEN  
RHGDLLNKYL**Y**LGGRCMRNIEVTIQHLITNGFNPQARKDPYRGFVYTSFQERATKISHGNVG  
KLARTYGEKNLNKICAKIAGDEGRHEKAYQIFSEEILKRDPDGLIHVFGDMMMRGQIVMPAEQ  
MTDGKDPDLYDNFSMVAQKTGVYTALDYAEIIDHLVKRWDLEHLEGLSPAEEKEREYLCLR  
PERYRKLATRSMNKKKVTEDEDPLKSGWYGRMA

*FAD6* (*Phatrdr\_48423*)

MVRFSTAALF**SL**STLTP**C**IGAFQLSSPAQLPT**S**RLLRRHTNTAPLSAVAVDSGSSDPALVGN  
LPLPNNNDNEDKNRRMPMMDLKGIALSGLKGQALS**VRAEDFPQAKDLRAV**IPKDCFEPDTAK  
SLGYLSVSTM**G**TILCSVVGANLLSVLDPSNPLTWPLWAAYGA**VTGTVAMGLWVLAHECGHG**  
AFSKNRSLQDAVGYIIHSIMLV**PYFSWQRSHAVHHQYTNHMELGETHPDRADKEGEKSLAL**  
RQFMLDSFGKDKGMKAYGGLQSFLH**LIVGWPAYLLIGATGGPDRG**MTNHFPNPLSTPTQPK  
KELFPGNWKEK**VYQSDIGIAAVVGALIAWTVTSG**LPVALYGGPLIVINA**WLVLYTWLQHT**  
DTDVPHFSSDNHNFVKGALHTIDRPYDKLDPWGIIDFLHHKIGTTHVAHHFDSTIPHYKAQIAT  
DAIKAKFPEVYLYDPTPIPQAMWRVAKGCTAVEQRGDAWVWKNEGIEDLVEHRQSKLSSE

*PlastidDelta6FAD* (*Phatrdr\_50443*)

MKFLHSALIVLT**S****A****S****A****F**TATNFFCL**S**QYG**SS****V****S**GC**PED****FIRQW**KAAGSTTNRISKKNVA  
YDCDEDANC**VIV**DACD**E**QCRTSLDV**R**IHG**KWY**DL**S**GR**W**RKAHPAGAH**WIDWY**DGRDATEV  
MDAFH**SEK**GRAMYKRLPASSTESVAMLETTIAPDSSTQIAFRQLR**DD**LEKEGWWKRDMVHE  
FTQLGIWASLV**V**GA**A**VT**A**HS**APPL**AT**F**LL**G**LS**M**TAAGWL**G**HDFI**H**GV**D**SFT**D**RLRN**F**AGVAA  
GLGPTWW**S**DKHNKH**H**ALT**N**EQGV**D**E**I**AT**D**PFL**F**TW**A**P**D**P**K**DD**S**PL**R**K**I**Q**H**L**I**F**V**W**P**FS**A**  
ALWRVDTMQV**V**IE**A**VEN**K**RVG**A**K**G**E**L**Y**G**LL**H**Y**A**VL**F**TF**P**VT**V**W**L**PA**I**FL**S**GL**M**SA**L**IV**T**  
THQ**SEEM**F**E**T**Y**Q**PDW**V**T**A**Q**F**STRNA**V**T**T**NPF****E**WL**W**GG**M**Q**Y**Q**LE**HH**L**F**PSMPRN**R**Y**P**ALR**  
ERLIQFA**ADN**K**I**P**GGY**RES**GE****E****I**LRMN**W**NL**Y**K**S**VA**E****A****D****A****V**PG**AP**TR**G**RL**G**QQ**GAIRET**  
NSPAA**QQ**E**K**A**I**D**Q**TV**A**K**G**NG**P**AE**S**V

*FAD4* (*Phatrdr\_41301*)

MILHGKTYT**S**CY**P**ESS**R**LC**S**N**V****S****K****A****F**N**K****M****S****L****P****R****Q****L****A****A****T****V****C****F****L****A****A****K****T****S****S****F****S****L****S****S****ST****R****T****V****H****R**  
SGLKPLHMAVIPDPSSSQGLY**ST****S**LR**A****L****A****S****T****E****T****T****A****S****N****K****E****K****T****K****P****S****W****N****D****G****F****V****G****L****E****G**  
GLERPKGRNAQIVVEGDSLETQPFQVAAVSVTFAAHAGFLVNSFSGMVEASSGNIALTSIHAIV

LTLVSWVIADFGSGVLHWATDNYGNGKTPVMGGIIAAFQGHHSAPWTIAQRGFCNNVYKLCI  
PFGIVPMLAINAIAPPDVFFMATFCVFEIMSQEFKWSQLKSETPGVNVWLQDSGLTIARK  
PHALHHLAPFEGNYCIISGICNPVLDQSGFFRRLERVVYSLNGIESNAWKLDPALERTLSGDY  
KFPKARSSSSKP

*Putative PlastidOmega3FAD (Phatdr\_41570)*

MKLHIAPPLIISAYVFSVSIFHNTVNAFSLRIPSTHRTVFVFLPVTLNAKRWMVATGVETNAA  
VATPENDEIHPRRDWTHDEPPKLSEVKRMLPQEAFHIDTATSLFYFAVDFIAVASTMGLNSV  
VSSDIYLSFPIWGKFLAVAPLQILTGFAMWCWCIGHDAGHTVSKDRRFGALINRVVGEVA  
HSAICLTPFWAKSHLKHHMGHNLTRDYSHQWFIREERESLHPLIQLSHATRNLQLPILYLV  
YLLFGVPDGHHVVFYGRMWEQSTAKEKADAASVIVSLVTAGSLWINMGLANFFVVCMP  
WLVLFWLFMVTYLQHSDDGLLYDETWSFERGAFQTVDRDYGTWINRMSHHMDGHLV  
HHLFFTRVPHYRLEEATKSLYAVMAARGQSHLIKTIDTPDFTQEIAKQFDKNWFFVNENQIVRK

## 2. Cytosol/ER Desaturases

*ADS (Phatdr\_28797)*

MDLSSEMLREIPGTLSQHYKAGNLNYPMIVYTTIVHTVALVGLCTIPQASAETLLWAFVLW  
PISGFGITVGVHRLWSHRSYEALPVRVFLMLCNSIANQGSIYHWSRDHRVHHKFSETDADPH  
NARRGFFFAHMGWLFVKKDPAVIEAGREMDFSDLLEDPVVALQKMVDPWFALYMCYVMPA  
QVASYFWGENFWTAFLVAGGLRYCFVLHCTWLVSAAHYGDHPYDLTSYPAENPFVSWC  
AVGEGWHNWHHKYPFDYAASEFGVSSQYNPSKLVIDVLASVGLVWGRKRGTAAWAMGRA  
RRDRDIAQGKEMPKQPPRPWEVRTVARKIA

*FAD2 (Phatdr\_25769)*

MGKGGQRAVAPKSATSSTGSATLSQSKEQVWTSSYNPLAKDAPELPTKGQIKAVIPKECFQ  
RSAFWSTFYLMRDLAMAAAFCYGTSQVLSTDLPQDATLILPWALGWGVYAFWMGTILTGPW  
VVAHECGHGAYSDSQTNDVVGIVHQALLVPYFAWQYTHAKHHRRTNHLVDGESHPSTA  
KDNGLGPHNERNSFYAAWHEAMGDGAFAVFQVWSHLFVGWPLYLAGLASTGKLAHEGWW  
LEERNAIADHFRPSSPMFPAKIRAKIALSSATELAVLAGLLYVGTQVGHLPVLLWYWGPYTFV  
NAWLVLYTWLQHTDPSIPHGEGETWVKGALSTIDRDYGFDFFHHTIGSTHVVHHLFHEM  
PWYNAGIATQKVKEFLEPQGLYNYDPTPWYKAMWRIARTCHYVESNEGVQYFKSMENVPLT  
KDVRSKAA

*ERDelta6FAD (PtD6) (Phatdr\_2948)*

MGKGGDARASKGSTAARKISWQEVKTHASPEDAWIIHSNKVYDVSNWHEHPGGAIFTH  
AGDDMTDIFAFAFHAPGSQSLMKKFYIGELLPETTGKEPQQIAFEKGYRDLRSKLIMMGMFKSN  
KWFYVYKCLSNMAIWAACALVFYSDRFVHLASAVMLGTFFQQSGWLAHDFLHHQVFTK  
RKHGDLGGLFWGNLMQGYSVQWWKNKHNGHHAVPNLCSSAVAQDGDPIDTMPLLAW  
VQQAQSYRELQADGKDSGLVKFMIRNQSYFYFPILLARLSWLNESFKCAFGLGAASENAAL

ELKAKGLQYPLLEKAGILLHYAWMLTVSSGFGGRFSFAYTAFYFLTATASCGFLLAIVFGLGHN  
GMATYNADARPDFWKLVQTTTRNVGGHGFQAFVDWFCGGLQYQVDHHLFPSLPRHNLA  
KTHALVESFCKEWGVQYHEADLVDGTMEVLHHLGSVAGEFVVDFVRDGPAM

*ERDelta5FAD1 (PtD5) (Phatdr\_46830)*

MAPDADKLQRQTTAVAKHNAATISTQERLCSLSSLNGEEVCIDGIYDLQSFDPGGETIK  
MFGGNDVTVQYKMIHPYHTEKLEMKRVGKVTDFCEYKFDTGFEREIKREVFKIVRRGK  
DFGTLGWFFRAFCYIAIFFYLQYHWVTTGTSWLLAVAYGVSQAMIGMNVQHDANHGATSKR  
PWVNDMLGLGADFIGGSKWLWQEQQHWTHAYTNHAEMDPDSFGAEPMLLFNDYPLDHPA  
RTWLHRFQAVFYMPVLAGYWLSAVFNPQILDLQQRGALSVGIRLDNAFIHSRRKYAVFWRA  
VYIAVNVIAPFYTNSGLEWSWRVFGNIMLMGVAESLALAVALFSLSHNFESADRDPТАPLKKT  
GEPVDWFKTQVETSCTYGGFLSGCFTGGLNFQVEHHLFPRMSSAWYPYIAPKVREICAKHGV  
HYAYYPWIHQNFLSTVRYMHAAGTGANWRQMARENPLTGRA

*ERDelta5FAD2 (Phatdr\_22459)*

MDVSLRNKSLSVDTLAPNHVCIDGKVFDSLSDHPPGDSIHVFGGNDVTVLYKMIHPHHG  
PNQYAQKMKLVGVIDKXRCEYSFDSDFGKEMKREVFQIVRRGQEFGTVGYFFRAFLYIAFFV  
AVVYRWTFTQGPSYALAVVFLAKALIGLNVQHDANHGAAAPPGRKNVWINDLLGWGADL  
IGGCKYLWIQKHWTHAYTNHAEKDPDAFAAEPLIFREYPASHPARQWYHKYQTLLFLPIIA  
GYWLSSVLSLEVAKLQDAGAMSATMKFENNVARQRKFTVFWRIVHLVII LGPPLRQHGLTA  
TALGQALTVGAAAGSLFLGCLFSLSHNFVNAERDPTAILAPPSTDGSDNENTAPVCWYKAQ  
VETSCTYGGFVSGALTGGLNQVEHHLFPRMSSAWYPFIAPTVRRVCAKHNVTYTYPWLW  
QNMASMMRYLHVTGGNTDAITKLE

*Putative ERDelat4FAD (Phatdr\_22510)*

MATTPSQSKTLDPLLLWMIHGNYYDLHTYVSRHPGGKEAILLGRGRDCTALFESYHPFTS  
QHRRVLEKHRVTIDTSFCSKQQDRKVKSKQTSSSEQASDVFYNMLCQRVAHALQSQGVDPIR  
DRGATWTRSIYYVFLFAALLASGYAHCTGSLLGSLLFGVAGWFIGALGHDGGHFAVSRRRAW  
LNDFSVWGISWLCNPIMWQHQHTYAHHSFTNEFDHDPDLHHFTTFLRVHRKFQQNCIYRNQA  
NWVYVFWAYTFVTGACFWIPWGVLRQTLYGLVDWTDKRPSRTAAFVFHLVTYFGLVM  
VLPFWTHGTWYKACLGVLHMTTSGLIFAFFSQINHINELSLDEKVVKLHRSTLDPTVRDSWA  
VAQVEASNNCTDSAFWHLFSNGLNLQIEHHLFPGINHCHLHHIAPIVVRETNEYGVRYKSY  
DSWSDIMRAMLKWLQLSVGLD