

Appendix A. Primers used in this study.

Primers	Sequences (5'-3')
<i>For gene cloning</i>	
GhFPS-F	ATGGCGGATCTCAGGTC
GhFPS-R	CTATTTCTGCCTCTTGTAGATCTTC
GhFPS- <i>KpnI</i> -F	TCATCAG <u>GGTACC</u> ATGGCGGATCTCAG
GhFPS- <i>XhoI</i> -R	TCATCA <u>CTCGAG</u> CTATTTCTGCCTCTT
<i>For qPCR</i>	
GhFPS1-F	TGCGTGTGCACTGGTTAIGGC
GhFPS1-R	CACTTTAGCAACATTGGCAGTG
GhFPS2-F	TGCGTGTGCACTGGTTATGTG
GhFPS2-R	CACTTTAGCAACATTGGCAGGG
GhACT4-F	TCGCTCGTTCCTTGTATAATTCATC
GhACT4-R	TGGATTCCAGCAGCTTCCA

Note: the restriction sites used for cloning are underlined and shown in bold.

Appendix B. Protein sequences used in construction of phylogenetic tree.

FPSs	Accession number
<i>Withania somnifera</i> FPS	ADR10437.1
<i>Oryza sativa Japonica Group</i> FPS	BAA19856.1
<i>Lilium longiflorum</i> FPS	ADZ57167.1
<i>Chimonanthus praecox</i> FPS	ACJ38671.1
<i>Betula platyphylla</i> FPS	AKQ62666.1
<i>Ornithogalum longibracteatum</i> FPS	AHA51120.1
<i>Pyrus communis</i> FPS	AHH93009.1
<i>Hedychium coccineum</i> FPS	AER12202.1
<i>Dendrobium catenatum</i> FPS	AFX68799.1
<i>Alisma plantagoaquatica</i> FPS	ADR83704.1
<i>Medicago sativa</i> FPS	ADC32809.1
<i>Chlorophytum borivilianum</i> FPS	AIU80188.1
<i>Salvia miltiorrhiza</i> FPS	ADT70780.1
<i>Magnolia chapensis</i> FPS	ACS74708.1
<i>Cyclocarya paliurus</i> FPS	ACY80695.1
<i>Panax notoginseng</i> FPS	AAAY53905.1
<i>Paeonia lactiflora</i> FPS	AKJ26301.1
<i>Hevea brasiliensis</i> FPS	AKM12344.1
<i>Sonneratia alba</i> FPS	AGJ71757.1
<i>Plagiochasma appendiculatum</i> FPS	AFM78687.1

<i>Cymbidium goeringii</i> FPS	AFP19446.1
<i>Phlegmariurus carinatus</i> FPS	AFO53558.1
<i>Huperzia serrata</i> FPS	AFO53556.1
<i>Helianthus annuus</i> FPS	AAC78557.1
<i>Humulus lupulus</i> FPS	AAK58594.1
<i>Aquilaria sinensis</i> FPS1	AHG54251.1
<i>Triticum aestivum</i> FPSD2	AGC11812.1
<i>Triticum aestivum</i> FPSB2	AGC11811.1
<i>Triticum aestivum</i> FPSA2	AGC11810.1
<i>Triticum aestivum</i> FPSD1	AGC11809.1
<i>Triticum aestivum</i> FPSB1	AGC11808.1
<i>Triticum aestivum</i> FPSA1	AGC11807.1
<i>Gossypium arboreum</i> FPS1	KHG28027.1
<i>Gossypium arboreum</i> FPS2	KHG22761.1
<i>Zostera marina</i> FPS1	KMZ75599.1
<i>Zostera marina</i> FPS2	KMZ59047.1
<i>Zea mays</i> FPS	AAQ14871.1
<i>Chlamydomonas reinhardtii</i> FPS	EDP03194.1
<i>Aralia elata</i> FPS	ADK12004.1
<i>Astragalus membranaceus</i> FPS	AKC92685.1
<i>Vitis vinifera</i> FPS	NP_001267864.1
<i>Bupleurum chinense</i> FPS	ADL74465.1

<i>Euphorbia pekinensis</i> FPS	ACN63187.1
<i>Eleutherococcus senticosus</i> FPS	AEY77151.1
<i>Catharanthus roseus</i> FPS	ADO95193.1
<i>Bacopa monnieri</i> FPS	ADV03080.1
<i>Centella asiatica</i> FPS	AAV58896.1
<i>Tripterygium wilfordii</i> FPS	AJY53644.1
<i>Gynostemma pentaphyllum</i> FPS	AII72208.1
<i>Leibnitzia anandria</i> FPS	AFW98445.1
<i>Taraxacum mongolicum</i> FPS	AFW98444.1
<i>Tanacetum coccineum</i> FPS	AFW98443.1
<i>Leucanthemum vulgare</i> FPS	AFW98442.1
<i>Chrysanthemum lavandulifolium</i> FPS	AFW98441.1
<i>Achillea asiatica</i> FPS	AFW98440.1
<i>Bos taurus</i> FPS	NP_803463.1
<i>Drosophila melanogaster</i> FPS	NP_477380.1
<i>Pongo abelii</i> FPS	NP_001125620.1
<i>Equus caballus</i> FPS	NP_001296255.1
<i>Mus musculus</i> FPS	AAL09445.1
<i>Danio rerio</i> FPS	NP_001020642.1
<i>Trichinella pseudospiralis</i> FPS	KRY68475.1
<i>Strongyloides ratti</i> FPS	CEF65900.1
<i>Papilio machaon</i> FPS	KPJ18339.1

<i>Defluviitoga tunisiensis</i> FPS	CEP79083.1
<i>Clostridium bornimense</i> FPS	CDM68610.1
<i>Streptomyces argenteolus</i> FPS	BAC66492.1
<i>Rhodobacter capsulatus</i> FPS	ADE44162.1
<i>Escherichia coli</i> FPS	BAA00599.1
<i>Micrococcus luteus</i> FPS	BAA25265.1
<i>Rhodovulum sulfidophilum</i> FPS	BAA96459.1
<i>Rhodobacter sphaeroides</i> FPS	BAA96457.1
<i>Fusarium langsethiae</i> FPS	KPA39997.1
<i>Phialophora attae</i> FPS	KPI36394.1
<i>Madurella mycetomatis</i> FPS	KOP44214.1
<i>Aspergillus lentulus</i> FPS	GAQ08849.1
<i>Blakeslea trispora</i> FPS	AFC92797.1
<i>Termitomyces</i> sp. J132	KNZ79524.1
<i>Trichoderma harzianum</i> FPS	KKP04355.1
<i>Ceratocystis platani</i> FPS	KKF97128.1
<i>Exophiala xenobiotica</i> FPS	KIW57626.1
<i>Exophiala oligosperma</i> FPS	KIW45882.1
<i>Cladophialophora immunda</i> FPS	KIW30036.1
<i>Exophiala spinifera</i> FPS	KIW18083.1
<i>Verruconis gallopava</i> FPS	KIW07049.1
<i>Exophiala sideris</i> FPS	KIV86042.1

<i>Mucor ambiguus</i> FPS	GAN06747.1
<i>Xanthophyllomyces dendrorhous</i> FPS	AIP94027.1
<i>Scedosporium apiospermum</i> FPS	KEZ39894.1
<i>Colletotrichum higginsianum</i> FPS	CCF34730.1
<i>Capronia semiimmersa</i> FPS	KIW68787.1
<i>Exophiala mesophila</i> FPS	KIV91925.1
<i>Lupinus albus</i> FPS	AAA86687.1
<i>Mangifera indica</i> FPS	AFJ53077.1
<i>Rosa rugosa</i> FPS	AKT74339.1
<i>Triticum urartu</i> FPS	EMS61433.1
<i>Artemisia annua</i> FPS	AIC83778.1
<i>Malus domestica</i> FPS	AAM08927.1

Appendix C. SDS-PAGE analysis of recombinant GhFPS1 and GhFPS2.

M: protein molecular weight marker; Lane 1: supernatant of non-induced GhFPS1; Lane 2: supernatant of induced GhFPS1; Lane 3: pellet of induced GhFPS1; Lane 4: supernatant of non-induced GhFPS2; Lane 5: supernatant of induced GhFPS2; Lane 6: pellet of induced GhFPS2. The arrow in figure indicates the location of the target protein.

