

**Table S1** The DNA primers used in this study.

Primer name	Sequence (5'-3')
NPT2U	GCTATTGGCTATGACTGG
NPT2R	ATAGAAGGCGATGCGCTG
PnPGK_LP	GTACTTGATGGTCATGGTG
PnPGK_RP	AGGTCAACCTTCTCCAAGAG
PnFL1_47_C08_LP1	CTATGCTGCCATGGACTGC
PnFL1_47_C08_RP1	CCAGCGTTGAAGTAGAGAGGA
PnFL1-093_N05_LP1	CGGAGAACATTAACCTTGACAA
PnFL1-093_N05_RP1	TCCCACCATTGTTGACCAC
PnFL1_55_M08_LP1	GGAGAACATTAACCTTGACAG
PnFL1_55_M08_RP1	TCCCACCATTGTTGACCAC
PnFL2_32_B04_LP1	GGGTTAGCCAAGGGCTAAA
PnFL2_32_B04_RP1	TGGATGAACAGGCTCAATCTC
PnFL2_46_L21_LP1	TTTGCTAACAGGACTTCCTC
PnFL2_46_L21_RP1	CCAGCAGCACAAATAGTGGAC
PnFL2-078_L13_LP1	AAAACATGGAGTCACACTCTG
PnFL2-078_L13_RP1	TGGCAATACTAGGCTAAACA

**Table S2** Relationship of galactinol synthase (GolS) from *Populus nigra* with those from *P. trichocarpa*.

<i>P. nigra</i>		<i>P. trichocarpa</i>			
Gene name	Locus	Gene name		Predicted protein length	Identity (%)
		Philippe et al. (2010)	Zhou et al. (2014)	(Amino acids)	
<i>PnGolS1</i>	Potri.002G191600	<i>PtGolS6</i>	<i>PtrGolS8</i>	337	96
	Potri.014G116800	<i>PtGolS7</i>	<i>PtrGolS6</i>	336	92
<i>PnGolS2</i>	Potri.005G006800	<i>PtGolS5</i>	<i>PtrGolS3</i>	334	97
<i>PnGolS3.1</i>	Potri.013G005800	<i>PtGolS4</i>	<i>PtrGolS1</i>	334	97
<i>PnGolS3.2</i>	Potri.013G005800	<i>PtGolS4</i>	<i>PtrGolS1</i>	334	97
<i>PnGolS4.1</i>	Potri.013G005900	<i>PtGolS3</i>	<i>PtrGolS2</i>	337	100
<i>PnGolS4.2</i>	Potri.013G005900	<i>PtGolS3</i>	<i>PtrGolS2</i>	337	99

**Figure S1**

PnGols1	1	MAPEIIVQS--ALKPAGEF-TKLASLPS-----RAYVTFLAGDGDYVKGVVGLAKGLRKVKT	52
PnGols2	1	MAPDITATL-ANNANSL-VKQASISS-----CAYVTFLAGDGDYVKGVVGLAKGLRKAKS	53
PnGols3.1	1	MAPDITTAI-ANNATTI-VKQASISS-----CAYVTFLAGDGDYVKGVVGLAKGLKKAES	53
PnGols3.2	1	MAPDITTAI-ANNATTI-VKQASISS-----CAYVTFLAGDGDYVKGVVGLAKGLRKAES	53
PnGols4.1	1	MAPHITTAL-ANSTNSL-VKQASISS-----CAYVTFLAGDGDYVKGVVGLAKGLRKAKS	53
PnGols4.2	1	MAPHITTAL-ANSTNSL-VKQASISS-----CAYVTFLAGDGDYVKGVVGLAKGLRKAKS	53
PtrGols1	1	MAPDITTPL-ANNATTI-VKQASISS-----CAYVTFLAGDGDYVKGVVGLAKGLRKAKS	53
PtrGols2	1	MAPHITTAL-ANSTNSL-VKQASISS-----CAYVTFLAGDGDYVKGVVGLAKGLRKAKS	53
PtrGols3	1	MAPDITATL-ANNTNSL-VKQASISS-----CAYVTFLAGDGDYVKGVVGLAKGLRKAKC	53
PtrGols4	1	MAPGV-PMDVIS-C-TGKVSTASTGYSKRAFVTFLAGNGDYVKGVVGLAKGLRKVKS	54
PtrGols5	1	MSPNS---IEPTTD-LQK-----RAYVTFLAGNGDYVKGVVGLAKGLRKVKS	44
PtrGols6	1	MAPELVQA-ALKPAGEF-TKPASLPS-----RAYVTFLAGNGDYVKGVVGLAKGLRKVKT	52
PtrGols7	1	MSPNA---IEPTFN-SHK-----RAYVTFLAGNGDYVKGVVGLAKGLRKAKS	44
PtrGols8	1	MAPELVRS-ALKPAGEF-TKLANLPS-----RAYVTFLAGDGDYVKGVVGLAKGLRKVKT	52
PtrGols9	1	MAPGV-PIDGNILGTGKVSTVNTGYSKRAYVTFLAGNGDYVKGVVGLAKGLRKVKS	55
PaxgGols1	1	MAPGV-PMDVIS-C-TGKVSTASHGYSKRAFVTFLAGNGDYVKGVVGLAKGLRRVKS	54
PaxgGols2	1	MAPHITTTL-ANTTNSL-VKQASISS-----CAYVTFLAGDGDYVKGVVGLAKGLRKAKS	53
AtGols1	1	MAPGLTQTADAMSTVTI-TRF-SLESVQDSL-RAYVTFLAGNGDYVKGVVGLAKGLRKVKS	58
AtGols2	1	MAPEINTKL-TVFPVHSA-TGGEK-----RAYVTFLAGTGDYVKGVVGLAKGLRKAKS	50
AtGols3	1	MAPEI-NNKLSYG-EKK-----RAYVTFLAGTGDYVKGVVGLAKGLRKTKS	44
AtGols4	1	MAPEI-SVNPMLY-SEKAHQAPP-----RAYVTFLAGNGDYVKGVVGLAKGLRKVKS	51
AtGols5	1	MTMTVEK---RIEADVT-VSHEGVE-----RAYVTFLAGNEDWMLVVVGLAKGLRKVKS	50
AtGols6	1	MAQMSMTIVEKSIAKADVT-VSHDRV-----RAYVTFLAGNEDWWMGVVGLAKGLRKVKS	53
AtGols7	1	MTPETHVDM-INASEKA-FKE-----RAYVTFLAGNGDYVKGVVGLAKGLRKVKS	48
PnGols1	53	AYPLIVAVLPDVPEEHFQIILESQGCMVREIEPVYPFENQTFAMAYYVINYSKLRWEFV	112
PnGols2	54	NYPLVVAILPDVPEEHHRKILASQGCIVREIEPVNPFFENQTFAMAYYVINYSKLRWEFV	113
PnGols3.1	54	KYPLVVAILPDVPEEHHRKILVSQGCIVREIEPVHPPFENQTEFAMAYYVINYSKLRWEFV	113
PnGols3.2	54	KYPLVVAILPDVPEEHHRKILVSQGCIVREIEPVHPPFENQTEFAMAYYVINYSKLRWEFV	113
PnGols4.1	54	KYPLVVAILPDVPEEHHRMILVSQGCIVREIEPVHPPFENQTRFAMPYYVINYSKLRWEFV	113
PnGols4.2	54	KYPLVVAILPDVPEEHHRMILVSQGCIVREIEPVHPPFENQTRFAMPYYVINYSKLRWEFV	113
PtrGols1	54	KYPLVVAILPDVPEEHHRKILVSQGCIVREIEPVHPPFENQTRFAMPYYVINYSKLRWEFV	113
PtrGols2	54	KYPLVVAILPDVPEEHHRMILVSQGCIVREIEPVHPPFENQTRFAMPYYVINYSKLRWEFV	113
PtrGols3	54	NYPLVVAILPDVPEEHHRKILASQGCIVREIEPVNPFFENQTFAMAYYVINYSKLRWEFV	113
PtrGols4	55	AYPLVVAMLPDVPEEHHRDILRSQGCIVREIEPIYPFENQIQFAMAYYVINYSKLRWNFE	114
PtrGols5	45	AYPLVVAILPDVPEEHHRREIILESQGCIIREIEPIYPFENQTOFSMAYYVINYSKLRWEFV	104
PtrGols6	53	AYPLIVAVLPDVPEEHRRKILESQGCIVREIEPVYPFENQTFAMAYYVINYSKLRWEFV	112
PtrGols7	45	SYPLMVAILPDVPEEHHRKILESQGCIVREIEPVYPFDNQTQFAMAYYVINYSKLRWEFV	104
PtrGols8	53	AYPLIVAVLPDVPEEHHRQILESQGCIVREIEPVYPFENQTFAMAYYVINYSKLRWEFV	112
PtrGols9	56	AYPLVVAILPDVPEEHHRDILRSQGCIVREIEPIYPFENQIQFAMAYYVINYSKLRWNFE	115
PaxgGols1	55	AYPLVVAMLPDVPEEHHRDILRSQGCIVREIEPIYPFENQIQFAMAYYVINYSKLRWNFE	114
PaxgGols2	54	KYPLVVAILPDVPEEHHRMILVSQGCIVREIEPVHPPFENQTRFAMPYYVINYSKLRWEFV	113
AtGols1	59	AYPLVVAMLPDVPEEHRRILVDQGCIVREIEPVYPFENQTFAMAYYVINYSKLRIWKFV	118
AtGols2	51	KYPLVVAVLPDVPEEHHRQLVDQGCIVREIEPVYPFENQTEFAMAYYVINYSKLRWEFV	110
AtGols3	45	KYPLVVAVLPDVPAHDHRQLLDQGCIVREIEPIYPFENQIQFAMAYYVINYSKLRIWKFV	104
AtGols4	52	AYPLVVAMLPDVPEEHREIILRSQGCIVREIEPVYPFEDNQTFAMAYYVINYSKLRWNFE	111
AtGols5	51	AYPLVVAVLPDVPEEHHRQIILVDQGCIIIRDIEPVYPFENITGYSMAYYVINYSKLRWEFV	110
AtGols6	54	AYPLVVAVLPDVPEEHHRQILLAQGCIIREIEPVYPFENITGYSMAYYVINYSKLRWEFV	113
AtGols7	49	AYPLVVAMLPDVPEEHREIILRSQGCIVREIEPVHPPDSQDAYARAYYIINYSKLRWNFE	108

Figure S1 (continued)

PnGols1	113	EYSKMIYLDGDIQVYDNIDHLDLFDGRFYAVMDCFCEKTWSHTLQYKIGYCQQCPDKVN	172
PnGols2	114	EYSKMIYLDGDIQVFDNIDHLDMDFDGYFYAAMDCCFCEKTWSNSPQYKIGYCQQCPDKVH	173
PnGols3.1	114	EYSKMIYLDGDIQVFDNIDHLDMDFDGYFYAAMDCCFCEKTWSNSPQYKIGYCQQCPDKVQ	173
PnGols3.2	114	EYSKMIYLDGDIQVFDNIDHLDMDFDGYFYAAMDCCFCEKTWSNSPQYKIGYCQQCPDKVQ	173
PnGols4.1	114	EYSKMIYLDGDIQVFDNIDHLDMDFDGYFYAAMDCCFCEKTWSNSPQYKIGYCQQCPDKVQ	173
PnGols4.2	114	EYSKMIYLDGDIQVFDNIDHLDMDFDGYFYAAMDCCFCEKTWSNSPQYKIGYCQQCPDKVQ	173
PtrGols1	114	EYSKMIYLDGDIQVFDNIDHLDMDFDGCFYAVMDCCFCEKTWSNSPQYKIGYCQQCPDKVQ	173
PtrGols2	114	EYSKMIYLDGDIQVFDNIDHLDMDFDGYFYAVMDCCFCEKTWSNSPQYKIGYCQQCPDKVQ	173
PtrGols3	114	EYSKMIYLDGDIQVFDNIDHLDMDFDGYFYAAMDCCFCEKTWSNSPQYKIGYCQQCPDKVH	173
PtrGols4	115	EYSKMMYLDADIQVFENIDHLDFTQDGYFYAAMDCCFCEKTWSHSPQHSIGYCYCQQCPPEKVT	174
PtrGols5	105	DYGKMIYLDGDIQVFDNIDHLDFKPTGYFYAAMDCCFCEKTWSHSPQHSIGYCYCQQCPPEKVT	164
PtrGols6	113	EYSKMIYLDGDIQVYDNIDHLDLFDGHTYAVMDCCFCEKTWSHSPQYKIGYCQQCPDKVN	172
PtrGols7	105	DYEKMIYLDGDIQVFDNIDHLDDEPNGYFYAAMDCCFCEKTWSSTTPQYQIGYCQQCPPEKVR	164
PtrGols8	113	EYSKMIYLDGDIQVYDNIDHLDLFDGRFYAAMDCCFCEKTWSHSPQYKIGYCQQCPDKVN	172
PtrGols9	116	EYSKMIYLDADIQVFENIDHLDFTQDGYFYAAMDCCFCEKTWSHSPQYSVGYCQQCPPEKIT	175
PaxgGols1	115	DYSKMMYLDADIQVFENIDHLDFTQDGYFYAAMDCCFCEKTWSHSPQYSIGYCYCQQCPPEKVT	174
PaxgGols2	114	EYSKMIYLDGDIQVFDNIDHLDMDFDGYFYAAMDCCFCEKTWSNSPQYKIGYCQQCPDKVQ	173
AtGols1	119	EYSKMIYLDGDIQVYENIDHLDLFDGYLYAVMDCCFCEKTWSHSPQYKIRYCYCQQCPDKVQ	178
AtGols2	111	EYNKMIYLDGDIQVFDNIDHLDLDPNGQFYAAMDCCFCEKTWSHSPQYKIGYCQQCPDKVT	170
AtGols3	105	EYSKLIYLDGDIQVFENIDHLDLFDLFDGNFYAVKDCFCCEKTWSHSPQYKIGYCQQCPDKVT	164
AtGols4	112	EYSKMIYLDADIQVFENIDHLDLFDLSDAYFYAAMDCCFCEKTWSHSLQYSIGYCYCQQCPPEKVT	171
AtGols5	111	EYEKMIYLDGDIQVFDNIDHLDFTFRGYLYAVKDCFCCEVSWSKTPQYKIGYCQQSPEKVT	170
AtGols6	114	EYEKMIYLDGDIQVFSNIDHLDFTFRGYLYAVKDCFCCEISWSKTPQYKIGYCQQCPPEKVT	173
AtGols7	109	EYNKMIYLDADIQVFGNIDDLFDMDQDGYLHGVISCFCEKIWSYTPLYSIGYCYCQQCPPEKVV	168

PnGols1	173	WP-AEMGQ-PPSPYFNAGMFVFEESIATYHDLIKTLKVTPPTPFAEQDFLNMYFRDIYTP	230
PnGols2	174	WP-AEMGP-KPPLYFNAGMFVYEPNLSTYHDLIELTTLKITSPLTLFAEQDFLNIFFRDVYKP	231
PnGols3.1	174	WP-AEMGP-KPPLYFNAGMFVYEPNLSTYHDLIELTTLKITSPLTLFAEQDFLNMFFRDVYKP	231
PnGols3.2	174	WP-AVMGP-KPPLYFNAGMFVYEPNLSTYHDLIELTTLKITSPLTLFAEQDFLNMFFRDVYKP	231
PnGols4.1	174	WP-AEMGP-KPPLYFNAGMFVYEPNLSTYHDLIELTTLKITSPLTLFAEQDFLNMFFRDVYKP	231
PnGols4.2	174	WP-AEMGP-KPPLYFNAGMFVYEPNLSTYHDLIELTTLKITSPLTLFAEQDFLNMFFRDVYKP	231
PtrGols1	174	WP-AEMGP-KPPLYFNAGMFVYEPNLSTYHDLIELTTLKITSPLTLFAEQDFLNMFFRDVYKP	231
PtrGols2	174	WP-AEMGP-KPPLYFNAGMFVYEPNLSTYHDLIELTVKTSPLTLFAEQDFLNMFFRDVYKP	231
PtrGols3	174	WP-AEMGP-KPPLYFNAGMFVYEPNLSTYHDLIELTVKTSPLTLFAEQDFLNMFFRDVYKP	231
PtrGols4	175	WP-AEMGS-PPPLYFNAGMFVFEPSRLTYESLIELTQLITPPTPFAEQDFLNMFQKTYKP	232
PtrGols5	165	WP-LEMGS-PPPLYFNAGMCLFEPKLETYEDLIELTLKVTTPTSFAEQDFLNMFFRDVYQP	222
PtrGols6	173	WP-AEMGQ-PPSLYFNAGMFVFEESIISTYHDLIKTLKVTPPTPFAEQDFLNMYFKDIYKP	230
PtrGols7	165	WP-KEMGS-PPPLYFNAGMFVFEKLLTLYEDLIELTLKVTPPTSFAEQDFLNMFFRDVYKP	222
PtrGols8	173	WP-AEMGQ-PPSPYFNAGMCLFEESIATYHDLIKTLKVTPPTPFAEQDFLNMYFKDIYTP	230
PtrGols9	176	WP-AEMGS-PPPLYFNAGMFVFEPSRLTYESLIELERLQITPPTPFAEQDFLNMFQKTYKP	233
PaxgGols1	175	WP-AEMGS-PPPLYFNAGMFVFEPSRLTYESLIELERLQITPPTPFAEQDFLNMFQKTYKP	232
PaxgGols2	174	WP-AEMGP-KPPLYFNAGMFVFEPSRLTYHDLIELTVKTSPLTLFAEQDFLNMFFRDVYKP	231
AtGols1	179	WPKAELGE-PPALYFNAGMFLYEPNLETYEDLIRTLKITPPTPFAEQDFLNMYFKDIYKP	237
AtGols2	171	WPEAKLGP-KPPLYFNAGMFVYEPNLSTYHNLIELTVKIVPPTLFAEQDFLNMYFKDIYKP	229
AtGols3	165	WPESELGP-KPPLYFNAGMFVYEPNLSTYHNLIELTVKIVPPTLFAEQDFLNMYFKDIYKP	223
AtGols4	172	WP-EDMESPPPPLYFNAGMFVFEPSPLTYESLILQTEITPPSPFAEQDFLNMFKEKVKYKP	230
AtGols5	171	WPVESLGA-PPPPLYFNAGMLVFGPNLVTYEDLIRVVQITTPTYFAEQDFLNIFYFRDIYKP	229
AtGols6	174	WPVESLGS-PPPPLYFNAGMLVFEPNLLTYEDLIRVVQITTPTYFAEQDFLINEYFTDIYKP	232
AtGols7	169	WP-AEMESAPPSPYFNAGMFVFEPNPLTYESLLQTLQVTPPTPFAEQDFLNMFVFKVFP	227

Figure S1 (continued).

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PnGols1	231	IPLVYNLVLAMLWRHFENVELDKVKVVHYCAAGSKPWRYTGTKEEKMQREDIEMLVKKWWD	290
PnGols2	232	IPSDYNLVLALLWRHFENINVDKVKVVHYCAAGSKPWRYTGTKEEDNMREDINMLVNKWWD	291
PnGols3.1	232	IPSDYNLVLAMLWRHFENINVLDKVKKVVHYCAAGSKPWRYTGTKEENMDREDIKMLVQKWWD	291
PnGols3.2	232	IPSDYNLVLAMLWRHFENINVLDKVKKVVHYCAAGSKPWRFTGKEENMDREDIKMLVQKWWD	291
PnGols4.1	232	IPSDYNLVLAMLWRHFENINVLDKVKKVVHYCAAGSKPWRFTGKEENMDREDIKMVVNKWWD	291
PnGols4.2	232	IPSDYNLVLAMLWRHFENINVLDKVKKVVHYCAAGSKPWRFTGKEENMDREDIKMVVNKWWD	291
PtrGols1	232	IPSDYNLVLAMLWRHFENINVLDKVKKVVHYCAAGSKPWRYTGTKEENMDREDIKMLVQKWWD	291
PtrGols2	232	IPSDYNLVLAMLWRHFENINVLDKVKKVVHYCAAGSKPWRFTGKEENMDREDIKMVVNKWWD	291
PtrGols3	232	IPSDYNLVLALLWRHFENINVLDKVKKVVHYCAAGSKPWRYTGTKEEDNMREDIKMLVQKWWD	291
PtrGols4	233	IPLMYNLVLAMLWRHFENVEVEKVVKVVHYCAAGSKPWRYTGTKEANMDREDIKMLVARWWD	292
PtrGols5	223	IPPVYNLVSAMLWRHPFDLDKVKKVVHYCAAGSKPWRYTGTKEENMDREDIKVLVKKWWE	282
PtrGols6	231	IPLVYNLVLAMLWRHFDNVELDKVKVVHYCAAGSKPWRYTGTKEENMQREDIKMLVEKWWG	290
PtrGols7	223	IPAVVYNLVSAMLWRHFENFEELDKVKVVHYCAAGAKPWRYTGTKEENMDREDIKVLVKKWWE	282
PtrGols8	231	IPLVYNLVLAMLWRHFENVELDRVVKVVHYCAAGSKPWRFTGKEENMDREDIKMLVKKWWD	290
PtrGols9	234	IPLLYNLVLAMLWRHFENVEVEKVVKVVHYCAAGSKPWRYTGTGEEEANMDREDIKMLVAKWWD	293
PaxgGols1	233	IPLMYNLVLAMLWRHFENVEVEKVVKVVHYCAAGSKPWRYTGTKEANMDREDIKMLVARWWD	292
PaxgGols2	232	IPSDYNLVLAMLWRHFENINVLDKVKKVVHYCAAGSKPWRFTGKEENMDREEIKMVVNKWWD	291
AtGols1	238	IPLVYNLVLAMLWRHFENVELGKVKKVVHYCAAGAKPWRYTGTKEANMEREDIKMLVKKWWD	297
AtGols2	230	IPPVYNLVLAMLWRHFENIELDQVKVVHYCAAGAKPWRFTGQEENMDREDIKMLVKKWWD	289
AtGols3	224	IPPVYNLVLAMLWRHFENIELNEAKVVHYCAAGAKPWRFTGQEENMDREDIKMLVKKWWD	283
AtGols4	231	IPLVYNLVLAMLWRHFENVELEKVVKVVHYCAAGSKPWRYTGTGEEEANMDREDIKMLVDKWWD	290
AtGols5	230	IPSTYNLVMAMLWRHPEHIDLDQISVVHYCANGSKPKWFDEAEEANMDREDIKMLVKKWWE	289
AtGols6	233	IPSTYNLVMAMLWRHPEHIDLDQISVIHYCANGSKPWRFDETEEHMDREDIKMLVKKWWD	292
AtGols7	228	VSPVYNLILSVLWRHPGKVDSLSEVKVVHYCEPGSKPWRYTGTGEEPNMDREDVKMLIKKWWD	287
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PnGols1	291	IYSDESLSQKLVADC-----TTDAEPVNL-QPFIAALSEAGAVQYVT-APSAA	337
PnGols2	292	IYHDESLDYKNTV-----VAAAGAEV-QPFLAVLSEAGIAHYIT-APSAA	334
PnGols3.1	292	IYNDESLDHNNTV-----VASSGSEL-QPIIIEALYEAGVDFHFT-APSAA	334
PnGols3.2	292	IYNDESLDHNNTV-----VASSGSEL-QPIIIEALYEAGVDFHFT-APSAA	334
PnGols4.1	292	IYQDESLDYKNTVAA-----AAASAGAEL-HPFLAALSEAGVVHYVT-APSAA	337
PnGols4.2	292	IYQDESLDYKNTVAAA-----AAASAGAEL-HPFLAALSEAGVVHYVT-APSAA	338
PtrGols1	292	IYNDESLDHKNTV-----VASSGSEL-QPIIIEALYEAGVDFHFT-APSAA	334
PtrGols2	292	IYQDESLDYKNTVAA-----AAASAGAEL-HPFLAALSEAGVVHYVT-APSAA	337
PtrGols3	292	IYHDESLDYKNTV-----VAAAGAEV-QPFLAALSEAGIAHYIT-APSAA	334
PtrGols4	293	IYNNESLDFKAEN-----SVFEEETLS-RSSIISSKPEPAISYIS-APSAA	336
PtrGols5	283	IYEDESLDFKNAT-----VPVAQEKL-GPLIAAI TEDGVGNHMN-LPSAA	325
PtrGols6	291	IYNDESLDYMKFVAD-----GIDAEPVNL-QSFIAALYEAGAVQYVT-APSAA	336
PtrGols7	283	IYEDESLDYKNIT-----VPVDQEKL-GPLIAAI TDDGVINHRN-LPSAA	325
PtrGols8	291	IYSDESLDSKKLVADC-----TTDAEPVNL-QPFIAALSEAGAVQYVT-APSAA	337
PtrGols9	294	IYNDESLLDFNGEN-----SVFEEETIFSRSSIISSMPEPAISYVP-APTA	338
PaxgGols1	293	IYNNESLDFKAEN-----SVFDEETILSPSSIISSKPGPAISYIS-APSAA	337
PaxgGols2	292	IYQDESLDYKNTV-AA-----ATASAGAEL-HPFLAALSEAGVVHYVT-APSAA	337
AtGols1	298	IYDDESLDYKKPVTVV-----DTEVDLVNL-KPFITALTEAGRLNYVT-APSAA	344
AtGols2	290	IYNDESLDYKNVVG-----DSHKKQQTL-QQFIEALSEAGALQYVK-APSAA	335
AtGols3	284	IYNDESLDYKNFNVHCGQKEDVHRKFKTL-BQFFTDLSEADVIQCAY-APSAA	334
AtGols4	291	VYNDESLDFKNSKI-----FADAEETVT-KSSILASVIEPEMTYFP-APSAA	334
AtGols5	290	IYEDSSLDDYKNFV-----ETESKINPV-TATLASKKLIVGDVITSI-APSAA	333
AtGols6	293	IYEDSSLDDYKNFV-----ETESKLSPNATLASKEESVGDVLISI-APSAA	336
AtGols7	288	IYNDESLLDKEPKS-----PADIEATVLI-ESTIILASVTEAPLSYSPAAPSAA	332

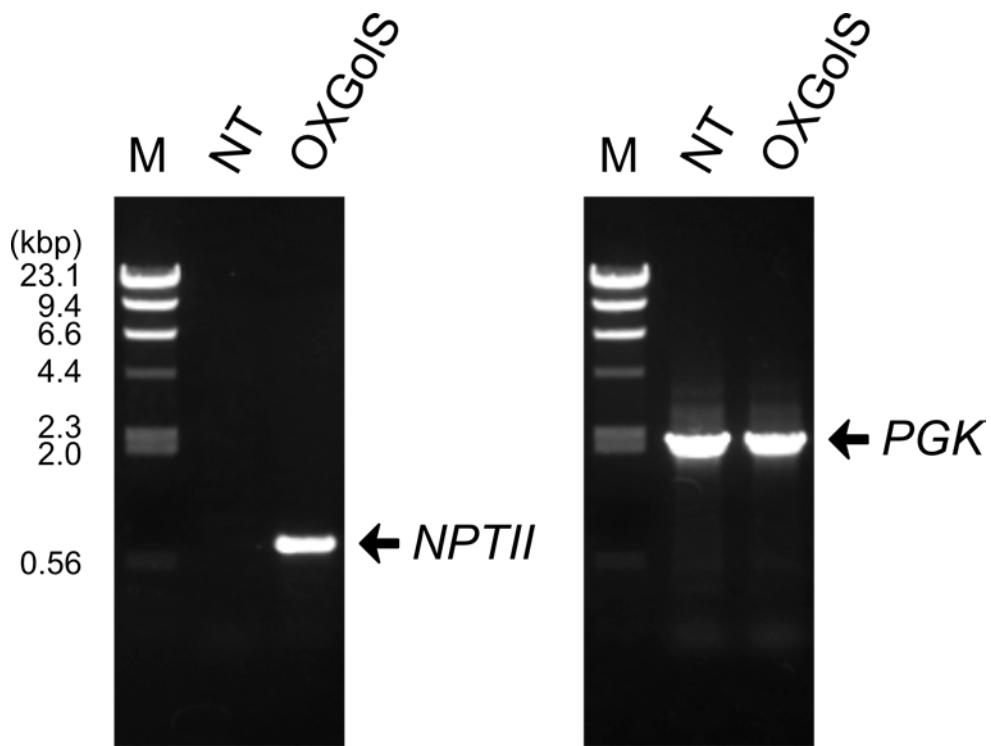
**Fig. S1**

Comparison of the predicted amino acid sequence of galactinol synthase (Gols) from *Populus nigra*

with those of other Golss from *P. trichocarpa*, *P. alba* × *grandidentata*, and *Arabidopsis thaliana*. A

putative serine phosphorylation site is indicated by an asterisk and the characteristic hydrophobic pentapeptide (APSAA) at the C-terminus is shown by a black bar.

**Figure S2**



**Fig. S2**

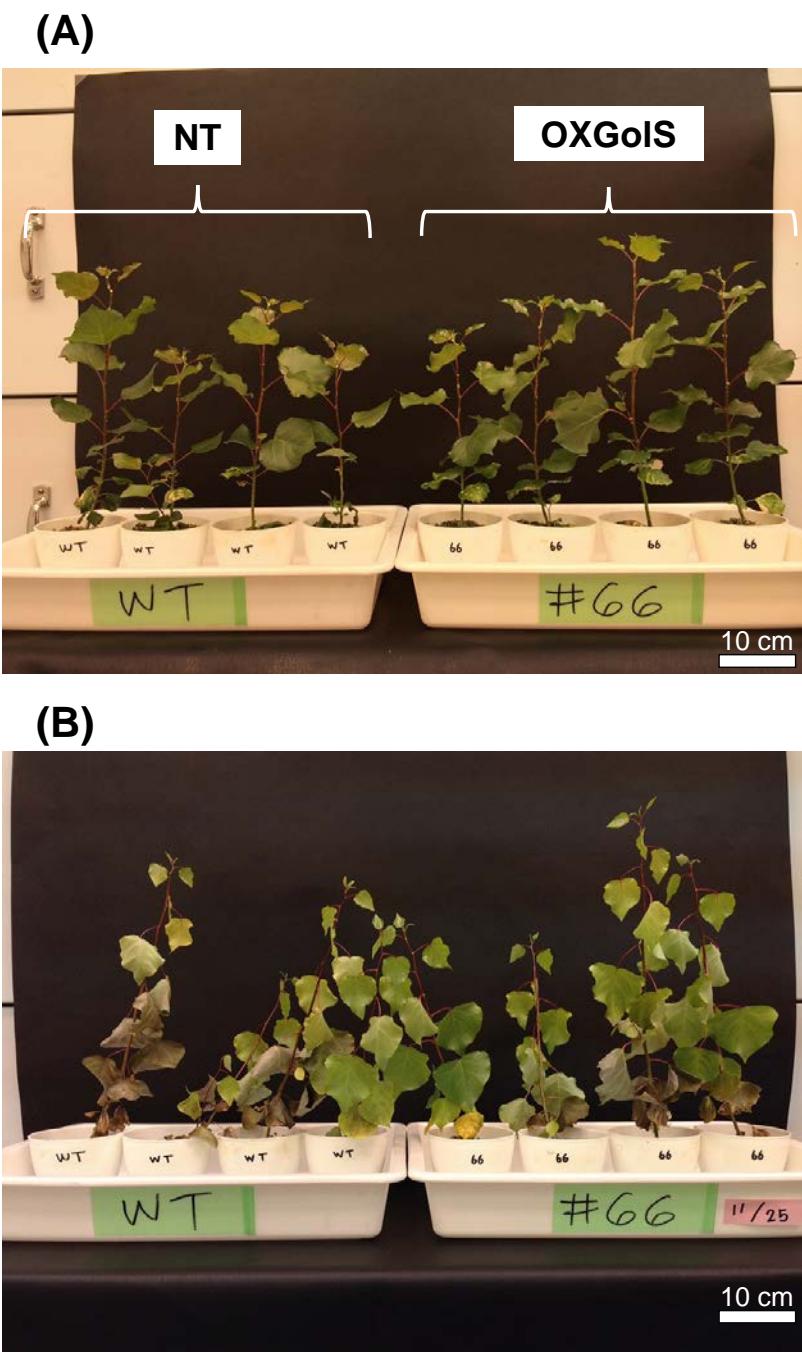
PCR confirmation of transgene integration. The neomycin phosphotransferase gene (*NPTII*) was amplified from only the genomic DNA of a *PnGolS2*-overexpressed poplar (OXGolS), whereas a non-transformant (NT) did not have the *NPTII* gene. A phosphoglycerate kinase gene (*PGK*, accession number AB018410) was used as a positive control for PCR. M, Lambda DNA/HindIII marker.



**Fig. S3**

Photographs of (A) non-transformants and (B) transformants overexpressing *PnGolS2* in *Populus nigra*. The photographs were taken 8 weeks after transfer to the Wagner-pots. The scale bar on the right bottom corner equals 10 cm in length.

**Figure S4**



**Fig. S4**

Photographs of the transformants overexpressing *PnGolS2* in *Populus nigra* (OXGoIS) and the

non-transformants (NT) (A) before and (B) on 14th day after withholding water supply. Aseptically cultivated clones were transplanted in a 1/10000 Wagner pot filled with 40 g vermiculite (in dry weight per pot). The plants were grown during about two months under an environmentally-controlled growth chamber before withholding water supply. The photosynthetically active photon flux density over the plants was 300–500  $\mu\text{mol m}^{-2} \text{s}^{-1}$ . The scale bar on the right bottom corner equals 10 cm in length.