## S4 Table. Description of differentially methylated regions (DMRs) occurring in annotated promoter and gene body sequences of the *Populus trichocarpa* genome.

DMR gene IDs	annotation	higher methylation site	<sup>m</sup> C context
POPTR_0001s01660g	NBS-LRR resistance gene-like protein ARGH35	Wallstawe	CHG & CHH
POPTR_0001s06110g	dirigent-like protein, regulates coupling of monolignol plant phenols to generate the cell wall polymers lignins and lignans that are involved in structural fortification and defense against pathogens	Wallstawe	СНН
POPTR_0001s17820g	DNA polymerase III	Wallstawe	CHG
POPTR_0001s19460g	zinc knuckle – zinc ion binding, nucleic acid binding	Anderlingen	CHG
POPTR_0001s42010g	catalytic domain of protein kinases	Wallstawe	СНН
POPTR_0002s03260g	horseradish peroxidase and related secretory plant peroxidases	Anderlingen	CHG & CHH
POPTR_0002s07850g	component of the thylakoid-localized Sec system involved in the translocation of cytoplasmic proteins into plastid	Wallstawe	CHG
POPTR_0002s23200g	aconitate hydratase, encodes a aconitase that can catalyze the conversion of citrate to isocitrate through a cis-aconitate intermediate, indicating a role in the response to oxidative stress	Wallstawe	CHG
POPTR_0003s12640g	glycosyl hydrolase family 32, beta- fructosidases	Wallstawe	СНН
POPTR_0005s01450g	PPR repeat family, DYW family of nucleic acid deaminases	Wallstawe	СНН
POPTR_0005s08440g	NAC domain protein	Wallstawe	CHG
POPTR_0005s11730g	AAA+ superfamily represents an ancient group of ATPases belonging to the ASCE division of the P-loop NTPase fold	Anderlingen	СНН
POPTR_0006s02510g	glutathione S-transferases	Anderlingen	СНН

DMR gene IDs	annotation	higher methylation site	<sup>m</sup> C context
POPTR_0006s09150g	aldo-keto reductases, a superfamily of soluble NAD(P)(H) oxidoreductases whose chief purpose is to reduce aldehydes and ketones to primary and secondary alcohols	Anderlingen	CHG & CHH
POPTR_0006s18990g	effector domain of the CAP family of transcription factors, members include CAP (or cAMP receptor protein (CRP)), which binds cAMP, FNR (fumarate and nitrate reduction)	Anderlingen	СНН
POPTR_0006s20500g	elongation factor Tu family protein	Wallstawe	CHG
POPTR_0006s22680g	NAD(P)-binding Rossmann-like domain	Anderlingen	CHG
POPTR_0008s18840g POPTR_0009s09810g POPTR_0009s17130g	chromosome segregation protein UDP-glucosyl transferase lectin L-type, legume lectins	Wallstawe Anderlingen Wallstawe	CHG CHG CpG & CHH
POPTR_0010s05110g	catalytic domain of protein kinases	Anderlingen	CHG
POPTR_0010s20390g POPTR_0011s15760g POPTR_0011s15770g POPTR_0014s01810g	peptidases S8 3 S-locus glycoprotein family galactose mutarotase-like peroxisomal membrane 22 kDa	Wallstawe Wallstawe Wallstawe Anderlingen	CHH CHH CHH CHG &
C	family protein	C	CHH
POPTR_0014s18950g POPTR_0015s09330g	glycosyltransferase like family galactinol-sucrose	Wallstawe Anderlingen	CpG CHH
POPTR_0016s021501g	galactosyltransferase glycosyltransferase family	Wallstawe	CHG
POPTR_0016s04630g	28 C-terminal domain cysteine/histidine-rich C1 domain family protein	Anderlingen	СНН
<i>POPTR</i> _0017s03570g	transcription factor Tfb4, TFIIH (subunit of the general trancriptions factors for promotor recognition and initiation of transcription)	Anderlingen	CHG
POPTR_0017s04440g	leucine-rich repeat receptor-like protein kinase	Wallstawe	CHG & CHH & CpG
POPTR_0017s06300g	AAA+ superfamily represents an ancient group of ATPases belonging to the ASCE division of the P-loop NTPase fold	Wallstawe	СНН
POPTR_0019s00260g	LRR 8, leucine rich repeat	Anderlingen	СНН