

**Table S1.** Starch synthesis related genes in rice

Classification		cDNA Acc. NO.		BAC or Scaffold Acc. NO.		Chromosome
				<i>japonica</i>	<i>indica</i>	
AGPase	Large subunit ( <i>AGPlar</i> )	<b>D50317*</b>	AK100910	AC120988	AAAA01004522	Chr. 5
	Large subunit isoform ( <i>AGPiso</i> )	<b>AY028314</b>	AK071497	AP004317 AP003688	AAAA01003476	Chr. 1
	small subunit ( <i>AGP<sub>sma</sub></i> )	<b>AY028315</b>	AK060270 AK073146	AP004756 AP004011	AAAA01000836	Chr. 9
SS	GBSS GBSSI ( <i>W<sub>x</sub></i> )	<b>X53694</b>	AF515480	AP002542	AAAA01010252	Chr. 6
			AF515481 AF515482 AF515483 AF031162 AF141954 AF141955 X65183 AB066093 X62134			
				AL607102		
	GBSSII ( <i>GBSSII</i> )	<b>AF109395</b>	AY069940	AP005325 AP003889	AAAA01001091	Chr. 7
	SSS SSI( <i>SSI</i> )	<b>AY299404</b>	D38221 AF165890 D16202 E06904 NM_185620	AB026295	AAAA01000263	Chr. 6
	SSII <i>SSII-1</i>	<b>AF383878</b>	NM_196673	AC087550 AC087547 AE017098	AAAA01005292	Chr.10
	<i>SSII-2</i>	<b>AF395537</b>	AJ308110	AP005297 AP004114	AAAA01001861	Chr. 2

		<i>SSII-3</i>	<b>AY423717</b> AF419099	AP003509	AAAA01002444	Chr. 6
					AAAA01011906	
	SSIII	<i>SSIII-1</i>	<b>AF432915</b>	AL606645	AAAA01000521	Chr. 4
		<i>SSIII-2</i>	<b>AY100469</b>	AP005441	AAAA01000600	Chr. 8
				AP004660		
	SSIV	<i>SSIV-1</i>	<b>AY373257</b> AY100470	AP003292	AAAA01000144	Chr. 1
		<i>SSIV-2</i>	<b>AY373258</b> AY100471	AC121365	AAAA01002233	Chr. 5
SBE	SBE1( <i>Sbe1</i> )		<b>D10838</b> AF136268 AY302112 D10752 E07175	AP003685 AP004685	AAAA01002467	Chr. 6
	SBE3 ( <i>Sbe3</i> )		<b>D16201</b> E08183	AP004879	AAAA01009319	Chr. 2
				AP004777	AAAA01010110	
				AP004790		
	SBE4 ( <i>Sbe4</i> )		<b>AB023498</b> E14723	AL731641	AAAA02013735	Chr. 4
DBE	Isoamylase( <i>ISA</i> )		<b>AB093426</b> AB015615	AP005509	AAAA01000810	Chr. 8
	Pullulanase ( <i>PUL</i> )		<b>D50602</b> AB012915	AL662959	AAAA01001658	Chr. 4

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\* Letters marked in bold imply the cDNA used as bait in search for genome sequence by Blastn