

Supplementary Table 2: SNPs identified through whole genome sequencing

Strain	Genome	Chr.	Position	Systematic and Gene Name	SNP Effect	Reference Nucleotide	SNP Nucleotide
GSY2532 (Vessel A)	<i>S. uvarum</i>	13	567,223	Sbay_13.344 (<i>S. uvarum</i> homolog of <i>YMR164C/MSSI</i>)	Non-synonymous, semi-conservative (G to A)	C	G
GSY2533 (Vessel B)	<i>S. cerevisiae</i>	7	28,807	<i>YGL251C (HFMI)</i> coding sequence	Non-synonymous, conservative (R to K)	C	T
	<i>S. uvarum</i>	15	354,918	Intergenic between Sbay_15.204 and Sbay_15.206 (<i>S. uvarum</i> homologs of <i>YHR020W</i> and <i>YHR021C</i>)	N/A	C	T
GSY2535 (Vessel C)	<i>S. cerevisiae</i>	4	243,445	<i>YDL122W (UBP1)</i> coding sequence	Synonymous (Y to Y)	C	T
	<i>S. cerevisiae</i>	5	405,451	<i>YER123W (YCK3)</i> coding sequence	Non-synonymous, conservative (L to F)	G	T
	<i>S. cerevisiae</i>	7	619,623	<i>YGR065C (VHT1)</i> coding sequence	Non-synonymous, semi-conservative (G to A)	C	G
	<i>S. cerevisiae</i>	14	258,697	<i>YNL204C (SPS18)</i> coding sequence	Non-synonymous, non-conservative (L to W)	A	C
	<i>S. uvarum</i>	4	608,898	Sbay_4.344 (<i>S. uvarum</i> homolog of <i>YBR097W/VPS15</i>)	Synonymous (S to S)	C	A
	<i>S. uvarum</i>	8	695,518	Sbay_8.386 (<i>S. uvarum</i> homolog of <i>YOR336W/KRE5</i>)	Non-synonymous, semi-conservative (D to G)	A	G