

59 A	R	D	R	E	T
S. 288C	EN TGE	KKNIK	EFKDI	SVQEY	YNK
RM11-1a	ENRGE	KKDIK	EFRDI	SVEEY	YTK
VIN13	ENRGE	KKDIK	EFRDI	SVEEY	YTK
YJM789	ENRGE	KKDIK	EFRDI	SVEEY	YTK
Y12	ENRGE	KKDIK	EFKDI	SVEEY	YTK
UWOPS87.2421	ENRGE	KKDIK	EFKDI	SVEEY	YTK
UWOPS05.217	ENRGE	KKDIK	EFKDI	SVEEY	YTK
<i>S. bayanus</i>	ENRGE	KKDVE	EFNDI	SVEEY	YTK
<i>S. mikatae</i>	ENRGE	KKDVR	EFNDI	SVEEY	YTK
<i>S. paradoxus</i>	ENRGE	KEDVR	EFNDI	SVEEY	YTK
	313	475	559	650	777

Figure S9 Alignment of the Abz1 amino acid modified in the strain 59A with the corresponding region seven other *Saccharomyces cerevisiae* strains and three other *Saccharomyces* species.