

TABLE 2.  $V_{\max}$  and  $K_m$  of glucose transport and specific growth rates<sup>a</sup> of the wild-type, *HXT1*, *HXT7*, *TM1*, *TM2*, *TM3*, *TM4*, *TM5*, *TM6\**, *TM11*, and *TM12* strains

Relevant genotype	$V_{\max}$ of (nmol/mg of protein/min)	$V_{\max}$ (% of wild-type value)	$K_m$ (mM)	$\mu$ ( $\text{h}^{-1}$ )	$\mu$ (% of wild-type value)
Wild type	663 ± 31	100	76 ± 9.2	0.35	100
<i>HXT1</i>	806 ± 140	122	250 ± 74	0.34	97
<i>TM12</i>	302 ± 13	46	44 ± 4.3	0.35	100
<i>TM11</i>	522 ± 47	79	72 ± 4.0	0.36	103
<i>HXT7</i>	125 ± 7.6	19	2.5 ± 0.6	0.34	97
<i>TM1</i>	73 ± 5.2	11	4.6 ± 1.3	0.30	86
<i>TM2</i>	113 ± 3.1	17	7.0 ± 0.7	0.30	86
<i>TM3</i>	129 ± 5.6	19	6.2 ± 1.4	0.28	80
<i>TM4</i>	103 ± 7.0	16	7.6 ± 2.1	0.26	74
<i>TM5</i>	35 ± 1.9	5	4.9 ± 0.9	0.27	77
<i>TM6*</i>	61 ± 2.1	9	3.5 ± 0.5	0.23	66

<sup>a</sup> Specific-growth-rate values were calculated during the respirofermentative phase. The specific growth rate of the *TM6\** strain, which shows only one phase, was determined during respiratory metabolism.