

**S1 Table. *ACE1* cluster expression in *Δbip1* during barley infection 24 hai.**

Gene ID	Gene name in <i>ACE1</i> cluster	Expression in wild type P1-2 *	Expression in <i>ΔBIP1:hph</i> *	Fold change **
<i>MGG_12447</i>	<i>ACE1</i>	$6.3 \times 10^{-4} \pm 3.7 \times 10^{-4}$	$8.8 \times 10^{-6} \pm 4.4 \times 10^{-6}$	0.01
<i>MGG_08386</i>	<i>BC2</i>	$1.3 \times 10^{-5} \pm 5.2 \times 10^{-6}$	$2.4 \times 10^{-4} \pm 1.0 \times 10^{-4}$	19
<i>MGG_08387</i>	<i>CYP1</i>	$1.2 \times 10^{-3} \pm 3.7 \times 10^{-4}$	$1.1 \times 10^{-4} \pm 5.6 \times 10^{-5}$	0.09
<i>MGG_15928</i>	<i>CYP2</i>	$1.3 \times 10^{-4} \pm 1.8 \times 10^{-5}$	$1.4 \times 10^{-3} \pm 5.0 \times 10^{-4}$	11
<i>MGG_08379</i>	<i>CYP3</i>	$1.8 \times 10^{-4} \pm 5.3 \times 10^{-4}$	$3.1 \times 10^{-6} \pm 3.4 \times 10^{-6}$	0.02
<i>MGG_08378</i>	<i>CYP4</i>	$1.4 \times 10^{-3} \pm 2.1 \times 10^{-4}$	$2.4 \times 10^{-5} \pm 2.7 \times 10^{-5}$	0.02
<i>MGG_08377</i>	<i>OME1</i>	$1.7 \times 10^{-4} \pm 1.1 \times 10^{-4}$	$6.7 \times 10^{-5} \pm 4.1 \times 10^{-5}$	0.40
<i>MGG_08381</i>	<i>ORF3</i>	$3.5 \times 10^{-3} \pm 1.1 \times 10^{-3}$	$1.6 \times 10^{-4} \pm 1.6 \times 10^{-4}$	0.05
<i>MGG_08391</i>	<i>RAP1</i>	$4.5 \times 10^{-4} \pm 1.9 \times 10^{-4}$	$1.4 \times 10^{-4} \pm 1.6 \times 10^{-4}$	0.30
<i>MGG_08380</i>	<i>RAP2</i>	$1.2 \times 10^{-3} \pm 3.7 \times 10^{-4}$	$6.7 \times 10^{-5} \pm 4.1 \times 10^{-5}$	0.05

\* Expression was determined by qRT-PCR. \*\* The ratio of expression between the *Δbip1* mutant and the wild-type control was calculated using the formula  $2^{-\Delta Ct \Delta bip1} / 2^{-\Delta Ct \text{ wild type}}$ .