

Product no **AS08 294****ALD | Fructose-1,6 bisphosphate aldolase****Product information**

<b>Immunogen</b>	overexpressed cytosolic fructose 1,6 bisphosphate aldolase (ALD) based on the sequence from <i>Arabidopsis thaliana</i> Q9LF98, At3g52930
<b>Host</b>	Rabbit
<b>Clonality</b>	Polyclonal
<b>Purity</b>	Serum
<b>Format</b>	Lyophilized
<b>Quantity</b>	100 µl
<b>Reconstitution</b>	For reconstitution add 100 µl of sterile water
<b>Storage</b>	Store lyophilized/reconstituted at -20°C; once reconstituted make aliquots to avoid repeated freeze-thaw cycles. Please remember to spin the tubes briefly prior to opening them to avoid any losses that might occur from material adhering to the cap or sides of the tube.

**Application information**

<b>Recommended dilution</b>	1 : 5000 (WB)
<b>Expected   apparent MW</b>	38   38 kDa
<b>Confirmed reactivity</b>	<i>Arabidopsis thaliana</i> , <i>Brassica napus</i> , <i>Cicer arietinum</i> , <i>Eragrostis tef</i> , <i>Gracilaria gracilis</i> (red algae), <i>Oryza sativa</i> , <i>Plasmodium chabaudi</i> , <i>Plasmodium falciparum</i> , <i>Thellungiella salsuginea</i>
<b>Predicted reactivity</b>	<i>Glycine max</i> , <i>Nicotiana attenuata</i> , <i>Nicotiana tabacum</i> , <i>Oryza sativa</i> , <i>Picea sitchensis</i> , <i>Physcomitrium patens</i> , <i>Pisum sativum</i> , <i>Populus jackii</i> , <i>Spinacia oleracea</i> , <i>Vitis vinifera</i> , <i>Zea mays</i> Species of your interest not listed? <a href="#">Contact us</a>
<b>Not reactive in</b>	<i>Synechocystis</i> sp.
<b>Additional information</b>	This product can be sold containing ProClin if requested
<b>Selected references</b>	<p><a href="#">Wang</a> et al. (2018). iTRAQ-based quantitative proteomics analysis of an immature high-oleic acid near-isogenic line of rapeseed. <i>Molecular Breeding</i> January 2018, 38:2.</p> <p><a href="#">Kamies</a> et al. (2017). A Proteomic Approach to Investigate the Drought Response in the Orphan Crop <i>Eragrostis tef</i>. <i>Proteomes</i>. 2017 Nov 15;5(4). pii: E32. doi: 10.3390/proteomes5040032.</p> <p><a href="#">Foley</a> et al. (2017). A Global View of RNA-Protein Interactions Identifies Post-transcriptional Regulators of Root Hair Cell Fate. <i>Dev Cell</i>. 2017 Apr 24;41(2):204-220.e5. doi: 10.1016/j.devcel.2017.03.018.</p> <p><a href="#">Parveen</a> et al. (2016). Chickpea Ferritin CaFer1 Participates in Oxidative Stress Response, and Promotes Growth and Development. <i>Sci Rep</i>. 2016 Aug 9;6:31218. doi: 10.1038/srep31218.</p> <p><a href="#">Yam</a> et al. (2016). Characterization of the Plasmodium Interspersed Repeats (PIR) proteins of <i>Plasmodium chabaudi</i> indicates functional diversity. <i>Sci Rep</i>. 2016 Mar 21;6:23449. doi: 10.1038/srep23449.</p> <p><a href="#">Dixit</a> (2015). Sulfur alleviates arsenic toxicity by reducing its accumulation and modulating proteome, amino acids and thiol metabolism in rice leaves. <i>Sci Rep</i>. 2015 Nov 10;5:16205. doi: 10.1038/srep16205.</p> <p><a href="#">Vera-Estrella</a> et al. (2014). Comparative 2D-DIGE analysis of salinity responsive microsomal proteins from leaves of salt-sensitive <i>Arabidopsis thaliana</i> and salt-tolerant <i>Thellungiella salsuginea</i>. <i>J Proteomics</i>. 2014 Jun 2. pii: S1874-3919(14)00288-7. doi: 10.1016/j.jprot.2014.05.018.</p>