

# Anti-NADH-ubiquinone oxidoreductase chain 1 antibody

Catalog: PHY1076S

## Product Information

<b>Description:</b>	Rabbit polyclonal antibody
<b>Background:</b>	Complex I is the largest protein complex of the oxidative phosphorylation system in mitochondrial and it catalyzes NADH-quinone oxidoreduction. Complex I represents the main entrance site for electrons into the respiratory electron transfer chain. In Arabidopsis, Complex I have at least 49 subunits and NAD1 (ATMG00516/ATMG01120/ATMG01275) is one of the subunit.
<b>Synonyms:</b>	NAD1, NADH DEHYDROGENASE 1
<b>Immunogen:</b>	KLH-conjugated synthetic peptide (15 aa from C terminal section) derived from <i>Arabidopsis thaliana</i> NAD1A (ATMG01275), NAD1B (ATMG01120) and NAD1C (ATMG00516).
<b>Form:</b>	Lyophilized
<b>Quantity:</b>	150 µg
<b>Purification:</b>	Serum Peptide affinity form antibody available upon request at <a href="mailto:info@phytoab.com">info@phytoab.com</a> .
<b>Reconstitution:</b>	Reconstitution with 150 µl of sterile water. "Note: please spin tube briefly prior to opening it to avoid any losses that might occur from lyophilized material adhering to the cap or sides of the tube".
<b>Stability &amp;Storage:</b>	Use a manual defrost freezer and avoid repeated freeze-thaw cycles. 12 months from date of receipt, -20 to -70°C as supplied. 6 months, -20 to -70°C under sterile conditions after reconstitution. 1 month, 2 to 8°C under sterile conditions after reconstitution.
<b>Shipping:</b>	The product is shipped at 4°C. Upon receipt, store it immediately at the temperature recommended above.

## Application Information

<b>Recommended Dilution:</b>	Western Blot(1:1000-1:2000) Note: Optimal dilutions/concentrations should be determined by the end user.
<b>Expected / apparent MW:</b>	36 kDa

Research Use Only

**Confirmed Reactivity:**

Coming soon

**Predicted Reactivity:**

Among species analyzed, the sequence of the synthetic peptide used for immunization is 100% homologues with the sequence in *Vitis vinifera*, *Glycine max*, *Cucumis sativus*, *Solanum tuberosum*, *Medicago truncatula*, *Nicotiana tabacum*, *Oryza sativa Indica Group*, *Brassica rapa subsp. oleifera*, *Zea mays subsp. mays*, *Brassica napus*, and 80-99% homologues with the sequence in *Oryza sativa Japonica Group*, *Hordeum vulgare*, *Gossypium raimondii*, *Triticum aestivum*, *Physcomitrium patens*.

For more species homologues information, please contact tech support at [tech@phytoab.com](mailto:tech@phytoab.com).