

## Anti-NADH dehydrogenase subunit 9, C-terminal antibody

Catalog: PHY3697S

## **Product Information**

**Description:** Rabbit polyclonal antibody

**Background:** 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

NAD9 (ATMG00070) is one of the subunit.

Synonyms: NAD9, NADH DEHYDROGENASE SUBUNIT 9

**Immunogen:** KLH-conjugated synthetic peptide (15 aa from C terminal section) derived from

Arabidopsis thaliana NAD9 (ATMG00070).

Form: Lyophilized

**Quantity**: 150 μg **Purification**: Serum

Peptide affinity form antibody available upon request at info@phytoab.com.

**Reconstitution:** Reconstitution with 150µl 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".

Stability &Storage: 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.

**Shipping:** The product is shipped at 4 °C. Upon receipt, store it immediately at the

temperature recommended above.

## **Application Information**

**Recommended Dilution:** Western Blot(1:1000-1:2000)

Note: Optimal dilutions/concentrations should be determined by the

end user.

Expected / apparent MW: 23 kDa



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 Triticum

aestivum, Oryza sativa, Solanum tuberosum, Brassica napus,

Hordeum vulgare, Glycine max, Setaria viridis, Nicotiana tabacum,

Zea mays, Solanum lycopersicum, Cucumis sativus, Gossypium

raimondii, and 80-99% homologues with the sequence in Vitis

vinifera.

For more species homologues information, please contact tech

support at tech@phytoab.com.