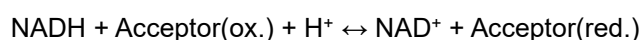


# DIAPHORASE 22 (Di-22)

[EC 1. 8. 1. 4]

from recombinant *E.coli*

## SPECIFICATION

State	: Lyophilized	
Specific activity	: more than 150 U/mg protein	
Contaminants	: (as Diaphorase activity = 100 %)	
	Adenylate kinase	< 0.01 %
	NADH oxidase	< 0.20 %

## PROPERTIES

Molecular weight	: ca. 110,000	
Subunit molecular weight	: ca. 50,000	
Optimum pH	: 8.0	(Fig,1)
pH stability	: 6.0 – 9.0	(Fig.2)
Thermal stability	: No detectable decrease in activity up to 70 °C.	(Fig. 3, 4)
Michaelis constants	: (50 mM HEPES buffer, pH 7.0, at 30 °C)	
	3-(4,5-Dimethyl-2-thiazolyl)-2,5-diphenyl-2 <i>H</i> -tetrazolium bromide (MTT)	0.345 mM
	NADH	0.033 mM
		(Table 1)
Substrate specificity	: NADH	100 %
	NADPH	1 %
	MTT	100 %
	Lipoate	103 %
		(Table 1)

## STORAGE

Store at -20°C

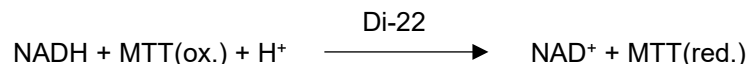
## APPLICATION

The enzyme is useful for measurement of various dehydrogenase reactions in the visible spectral range.

## ASSAY

### Principle

The change in absorbance is measured at 565 nm according to the following reaction.



### Unit Definition

One unit of activity is defined as the amount of Diaphorase that forms 1  $\mu\text{mol}$  of  $\text{NAD}^+$  per minute at 30 °C

### Solutions

- I Buffer solution ; 100 mM HEPES, pH 7.0
- II 3-(4,5-Dimethyl-2-thiazolyl)-2,5-diphenyl-2*H*-tetrazolium bromide (MTT) solution ; 10 mM (20 mg MTT disodium salt·2H<sub>2</sub>O/5 mL distilled water)
- III NADH solution ; 13.1 mM (0.100 g NADH disodium salt·3H<sub>2</sub>O /10 mL distilled water)
- IV Triton solution ; 10 % (1 mL TritonX-100 dilute with distilled water up to 10 mL)

### Preparation of Enzyme Solution

Dissolve the lyophilized enzyme with distilled water and dilute to 1 to 5 U/mL with 50 mM potassium phosphate buffer, pH 7.5, 1 mg/mL BSA.

### Procedure

1. Prepare the following reaction mixture and pipette 3.00 mL of reaction mixture into a cuvette.
 

Solution I	15.00 mL	Solution IV	1.50 mL
Solution II	1.50 mL	H <sub>2</sub> O	10.80 mL
Solution III	1.20 mL		
2. Incubate at 30 °C for about 3 minutes.
3. Add 0.01 mL of enzyme solution into the cuvette and mix.
4. Read absorbance change at 565 nm per minute ( $\Delta\text{Abs}_{565}$ ) in the linear portion of curve.

### Calculation

$$\text{Volume activity (U/mL)} = \frac{(\Delta\text{Abs}_{565}) \times (3.00 + 0.01)}{20.0 \times 0.01} \quad \text{X d.f.}$$

$$\text{Specific activity (U/mg protein)} = \frac{\text{Volume activity (U/mL)}}{\text{Protein concentration (mg/mL)}^*}$$

d.f. ; dilution factor

20.0 ; millimolar extinction coefficient of MTT ( $\text{cm}^2/\mu\text{mol}$ )

\*Protein concentration ; determined by Bradford's method

## REFERENCE

1. Packman, L.C., and Perham. R.N.; *FEBS Lett.*, **139**, 155 (1982)

**Table 1. SUBSTRATE SPECIFICITY OF DIAPHORASE 22**

Acceptor	MTT	Lipoate
Km <sup>Acceptor</sup> (mM)	0.345	2.0
Km <sup>NADH</sup> (mM)	0.033	0.01
Relative Activity	100	103
Assay Mixture	HEPESI (pH 7.0) 50 mM NADH 0.5 mM MTT 0.5 mM Triton X-100 0.5 %	Potassium Phosphate (pH 6.5) 70.5 mM NADH 0.2 mM NAD 0.3 mM Lipoate 10.2 mM EDTA 0.81 mM BSA 0.7 mg/mL
Wavelength for Measurement (nm)	565	340
Extinction coefficient (cm <sup>2</sup> /μmol)	20	6.22

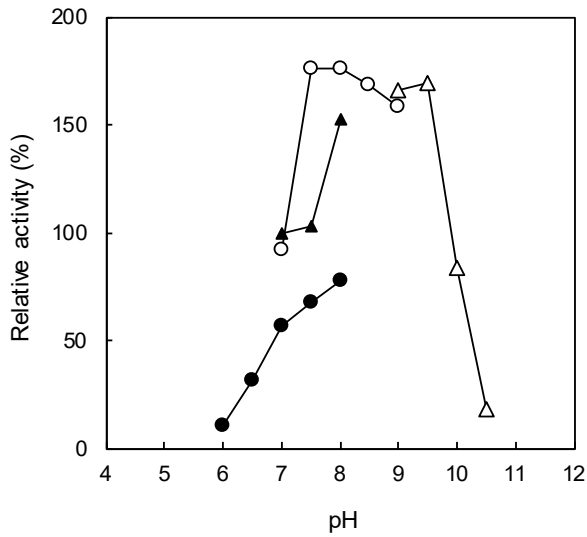


Fig. 1 pH profile

( △ Gly-KOH, ○ Bicine,  
● phosphate, ▲ HEPES )

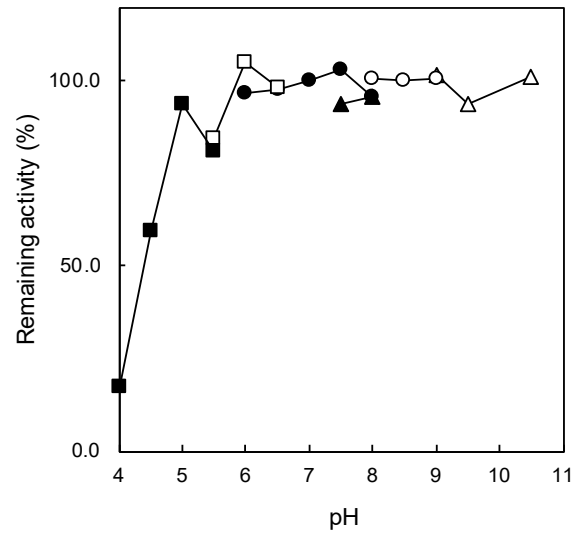


Fig. 2 pH stability

( treated for 24 hr at 4 °C in the following  
buffer solution (0.1 M) :  
△ Gly-KOH, ○ Bicine,  
● phosphate, ▲ HEPES,  
□ MES, ■ Citrate )

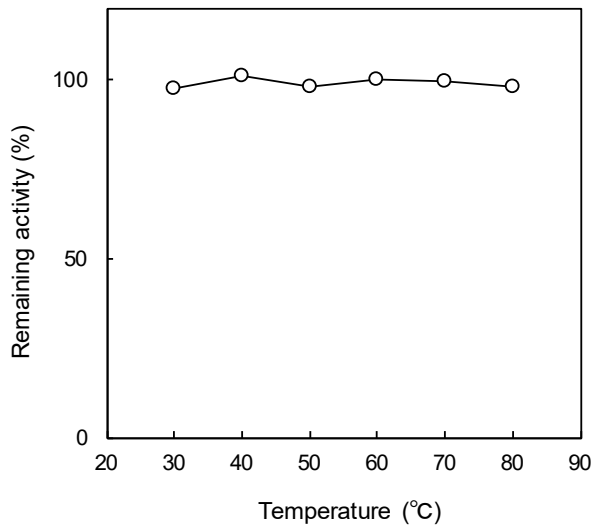


Fig. 3 Thermal stability

( treated for 15 min in 0.1M  
potassium phosphate buffer,  
pH 7.5 )

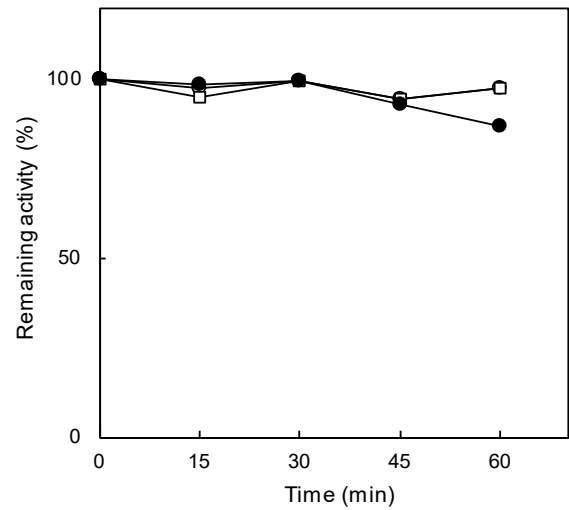


Fig. 4 Thermal stability

( treated for in 0.1M potassium  
phosphate buffer, pH 7.5  
○ 60 °C, □ 70 °C, ● 80 °C )