

**\*\*REPRESENTATIVE DATASHEET\*\***

**Matched-Pair Antibody Set  
for ELISA of  
Murine Fibrinogen antigen (Fg)**

Sufficient reagent for 4 x 96 well plates

**Product #:** MFG-EIA  
**Lot #:** SAMPLE  
**Expiry Date:** SAMPLE

**Store at -10 to -20°C**

For Research Use Only  
Not for use in diagnostic procedures.

**Description of Fibrinogen (Fg)**

Fibrinogen is an abundant plasma protein (5-10  $\mu$ M) produced in the liver. The intact protein has a molecular weight of 340 kDa and is composed of 3 pairs of disulphide-bound polypeptide chains named  $\alpha$ ,  $\beta$  and  $\gamma$ . Fibrinogen is a triglobular protein consisting of a central E domain and terminal D domains. Proteolysis by thrombin results in release of Fibrinopeptide A (FPA,  $\alpha$ 1-16) followed by Fibrinopeptide B (FPB,  $\beta$ 1-14) and the fibrin monomers that result polymerize in a half-overlap fashion to form insoluble fibrin fibrils. The chains of fibrin are referred to as  $\alpha$ ,  $\beta$  and  $\gamma$ , due to the removal of FPA and FPB. The polymerised fibrin is subsequently stabilized by the transglutaminase activated Factor XIII that forms amide linkages between  $\gamma$  chains and, to a lesser extent,  $\alpha$  chains of the fibrin molecules. Proteolysis of fibrinogen by plasmin initially liberates C-terminal residues from the  $\alpha$  chain to produce fragment X (intact D-E-D, which is still clottable). Fragment X is further degraded to non-clottable fragments Y (D-E) and D. Fragment Y can be digested into its constituent D and E fragments. Digestion of non-crosslinked fibrin with plasmin is very similar to the digestion of fibrinogen, which results in production of fragments D and E. Degradation of crosslinked fibrin by plasmin results in fragment DD (D-Dimer consisting of the D domains of 2 fibrin molecules crosslinked via the  $\gamma$  chains), fragment E (central E domain) as well as DDE in which fragment E is non-covalently associated with DD. For human crosslinked fibrin, the relative weights of the cleavage fragments produced are: 184 kDa for fragment DD, 92 kDa for D, 50 kDa for E, 1.54 kDa for FPA and 1.57 kDa for FPB<sup>1-3</sup>.

**Principle of Sandwich-style ELISA**

Affinity-purified polyclonal antibody to Fg is coated onto the wells of a microtitre plate. The plates are washed and plasma or other fluids containing Fg are applied. The coated antibody will capture the Fg in the sample. After washing the plate to remove unbound material, a peroxidase conjugated second antibody to Fg is added to the plate to bind to the captured Fg. After washing the plate to remove unbound conjugated antibody, the peroxidase activity is expressed by incubation with o-phenylenediamine (OPD). After a fixed development time the reaction is quenched with the addition of  $H_2SO_4$  and the colour produced is quantified using a microplate reader. The colour generated is proportional to the concentration of Fg in the sample.

**Supplied Materials:**

- 1. Capture Antibody (MFG-EIA-C):** One yellow-capped vial containing 0.4 ml of affinity-purified polyclonal anti-fibrinogen antibody for coating plates.
- 2. Detecting Antibody (MFG-EIA-D):** One red-capped vial containing 0.4 ml of peroxidase conjugated polyclonal anti-fibrinogen antibody for detection of captured fibrinogen.

**Note:** Antibodies are supplied in a 50% (v/v) glycerol solution for storage at -10 to -20°C. Keep vials tightly capped. Do not store in frost-free freezers.

**Materials Required but not Provided:**

**1. Coating Buffer:** 50 mM Carbonate  
1.59g of  $Na_2CO_3$  and 2.93g of  $NaHCO_3$  up to 1 litre. Adjust pH to 9.6. Store at 2-8°C up to 1 month.

**2. PBS:** (base for wash buffer and blocking buffer)  
8.0g NaCl, 1.15g  $Na_2HPO_4$ , 0.2g  $KH_2PO_4$  and 0.2g KCl, up to 1 litre. Adjust pH to 7.4, if necessary. Store up to 1 month at 2-8°C, discard if there is evidence of microbial growth.

**3. Wash Buffer:** PBS-Tween (0.1%,v/v)  
To 1 litre of PBS add 1.0 ml of Tween-20.  
Check that the pH is 7.4. Store at 2-8°C up to 1 week.

**4. Blocking Buffer:** PBS-BSA (1%, w/v)  
Dissolve 2.5 g of Bovine Serum Albumin (Sigma-RIA grade) in 200 ml of PBS. Adjust pH to 7.4, if required, then make up to 250 ml with PBS. Aliquot and store frozen at -20°C.

**5. Sample and Conjugate Diluent:** HBS-BSA-T20  
5.95g HEPES (free acid), 1.46 g NaCl, 2.5 g Bovine Serum Albumin (Sigma, RIA grade) dissolved in 200 ml  $H_2O$ . Add 0.25 ml of Tween-20, check and adjust pH to 7.2 with NaOH, then make up to a final volume of 250 ml with  $H_2O$ . Aliquot and store frozen at -20°C.

**6. Substrate Buffer:** Citrate-Phosphate buffer pH 5.0  
2.6g Citric acid and 6.9g  $Na_2HPO_4$  up to a final volume of 500 ml with purified  $H_2O$ . Store at 2-8°C up to 1 month.

**7. OPD Substrate:** (o-Phenylenediamine.2HCl) TOXIC!  
(5mg tablets: Sigma # P-6912). Make up immediately before use. Dissolve 5mg OPD in 12 ml substrate buffer then add 12  $\mu$ l 30%  $H_2O_2$ . Do not store.

**8. Stopping Solution:** 2.5 M  $H_2SO_4$   
Caution: VERY CORROSIVE! GENERATES HEAT ON DILUTION! Where stock sulphuric acid is 18 Molar, add 13.9 ml to 86 ml  $H_2O$ . Store at room temperature.

**9. Other:**  
Microplates, 96-well Immulon 4-HBX (<http://www.labsystems.fi>)  
Microplate washer (optional)  
Microplate reader.

## Assay Procedure:

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### 1. Coating of plates:

Dilute the capture antibody 1/100 in coating buffer (preferably in a polypropylene tube) and immediately add 100 µl to every well in the plate. Incubate 2 hours at 22°C or overnight at 2-8°C.

### 2. Blocking:

Empty contents of plate and add 150 µl of blocking buffer to every well and incubate for **60 minutes** @ 22°C.

Wash plate X 3 with wash buffer.

### 3. Samples:

Normal mouse plasma is diluted 1/4,000 (100%) then serial 1/2's down to 1/128,000 (3.13%). Sample plasmas are diluted 1/8,000, 1/16,000 & 1/32,000. All dilutions are made in HBS-BSA-T20 sample diluent. Apply 100 µl/well and incubate plate @ 22°C for 90 minutes. Wash plate X 3 with wash buffer.

### 4. Detecting Antibody:

Dilute the detecting antibody 1/100 in HBS-BSA-T20 diluent and apply 100 µl to each well. Incubate plate @ 22°C for 60 minutes. Wash plate X 3 with wash buffer.

### 5. OPD Substrate:

Apply 100 µl of freshly prepared OPD substrate to every well. Allow colour to develop for 10-15 minutes then stop colour reaction with the addition of 50 µl/well of 2.5 M H<sub>2</sub>SO<sub>4</sub>. The plate can be read at a wavelength of 490 nm.

## Calculation of Results:

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The construction of a proper reference curve is of no less importance than any other aspect of the assay. A reference curve should be constructed by plotting the known concentration of standards versus absorbance. This can be done manually using graph paper, or by using curve-fitting computer software. In our experience, the dose response curves of most immunoassays tend to be sigmoid in shape. Although linear regions can be identified within the curve, the best overall fit is often obtained using an algorithm that provides a weighted theoretical model of fit throughout the entire curve, such as a 4-parameter or 5-parameter logistic curve fit<sup>4,5</sup>. In general, the simplest model that defines the concentration-response relationship should be used<sup>6</sup>.

The "back-fit" test is a simple and reliable method to determine if a curve-fitting method is appropriate. In this test, the apparent concentrations for the absorbance values of each standard point are read from the reference curve. The derived values are compared to the assigned values. An appropriate curve fitting method will produce derived values that closely match assigned values throughout the range of the curve, within user-defined limits<sup>6</sup>. The coefficient of determination ( $R^2$ ) is a valuable indicator of the overall fit, but should not be used by itself in the selection of a curve fitting method, as a poor fit in a particular region of the curve may not be evident from this value alone<sup>5,6</sup>.

**In the quality control of this product we have determined that under the conditions described above, a reference curve that is constructed using serial dilutions of normal pooled plasma, will produce a correlation coefficient ( $R^2$ ) of at least 0.980 using a semi-log fit, and an  $R^2$  of at least 0.990 using a 4-parameter logistic curve fit algorithm.** However, the performance characteristics of in-house assays developed using this product in other laboratories may vary slightly from ours. Different curve fitting methods may be employed but we recommend that the back-fit test be applied as evidence that the fitting method is appropriate.

## Technical Notes:

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- This paired antibody product is intended to facilitate the end user in establishing an in-house immunoassay for research purposes only. Assay validation is the responsibility of the end user and should be done according to user-defined protocols<sup>6</sup>.
- Reference calibrators should be of the same matrix and anticoagulant as the samples to be tested (example serum or plasma, citrate or EDTA).
- Do not use samples diluted less than 1/100, as falsely high readings may result.
- The optimal colour development time should be determined empirically as the time required to obtain an absorbance of at least 1.000 at 490 nm for the 100% reference point, not to exceed 20 minutes.
- The presence of heterophilic antibodies in samples may interfere in ELISA by binding to the capture and/or detecting antibodies.
- The wells should not be allowed to become dry. Keep plate covered or in a humid chamber during incubations.
- Antibodies are supplied in a 50% glycerol solution and can be centrifuged briefly in a micro-centrifuge to gather residual reagent from the cap and walls of the tube.

## References:

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1. Hantgan RR, Francis CW, Marder VJ; Fibrinogen Structure and Physiology; in Hemostasis and Thrombosis, 3<sup>rd</sup> Edition, eds. RW Colman, J Hirsh, VJ Marder and EW Salzman, pp 277-300, J.B. Lippincott Co., Philadelphia PA, USA, 1994.
2. Shafer JA, Higgins DL; Human Fibrinogen; CRC Critical Reviews in Clinical Laboratory Sciences 26, pp 1-41, 1988.
3. Binnie CG, Lord ST; The Fibrinogen Sequences that Interact with Thrombin; Blood 81, pp 3186-3192, 1993.
4. Nix, B, Wild D, in Immunoassays, A Practical Approach, editor J.P. Gosling, pp. 239-261, Oxford University Press, 2000.
5. NCCLS. Evaluation of the Linearity of Quantitative Analytical Methods; Proposed Guideline – Second Edition. NCCLS Document EP6-P2 (ISBN 1-56238-446-5, NCCLS, Wayne, Pennsylvania USA, 2001.
6. FDA Guidance for Industry. Bioanalytical Method Validation; May 2001, available on the internet: [www.fda.gov/cder/guidance/index.htm](http://www.fda.gov/cder/guidance/index.htm)