A STUDY OF POLYMERIZATION SITES IN BOVINE FIBRINOGEN. L. Tranqui, A. Chapel, N. Suscillon. Lab. HématoLogie, IRF. Centre d'Etudes Nucléaires, Brennoble, France.

Abnormalities in bovine fibrinogen polymerization have been brought about by slight proteolysis on the native molecule by two purified aminopeptidases that we have called A and B and which come from bacteria. These degraded molecules have a molecular weight between 120,000 and 250,000. The aminopeptidase A induces the polymerization of fibrinogen into ordered structures, the period being 8 nm, when the ionic strength (I) is between 0.1 and 0.2. Fibrinogen modified by the aminopeptidase B precipitates at 4°C (0.075 < I < 0.2) into fibrils with a major period of 66 nm. The action of thrombin on these modified fibrinogen molecules at I = 0.3 gives fibres each with two minor striations, whose distribution was however different. These structures were totally different from those of fibrin fibres with its typical major period of 23 nm and its three minor striations. The analysis of the polypeptide chains, the N-terminal acids and the amino-acids composition of these degraded fibrinogen molecules enables us to differentiate the respective roles of the polymerization sites in the N and C terminal parts of the Aα and Bβ chains.

Therefore, according to our hypothesis the fibre structures obtained from fibrinogen depend on the type of the sites which are discovered on fibrinogen by enzymatic splits and by slight conformational changes.

QUANTITATIVE ABNORMALITIES OF Aα CHAIN MOLECULAR WEIGHT IN THE FIBRINOGEN OF CIRRHOTIC PATIENTS. Mark Weinstain and Daniel Dayton. Boston VA Hospital, Boston, Massachusetts.

A novel two-stage SDS gel electrophoretic procedure was devised to examine the molecular weight heterogeneity of fibrinogen in small samples of whole plasma from 12 normal and 7 cirrhotic individuals. Fibrinogen was first separated from other plasma proteins on a large pore gel, cut out of the gel, reduced, and separated into its component Aα, Bβ, and γ chains on a second gel. Two major mol wt species—fibrinogen I and fibrinogen II—were seen on the second gel. The 25000 mol wt difference between these two forms reflected a decrease primarily in the size of one of the fibrinogen II Aα chains. In both normals and cirrhotic patients fibrinogen II comprised 30% of the total (range 29-35%). Fibrinogen I and II each contained two major high mol wt Aα chains—Aα1 and a smaller Aα2—that differ by 2000 mol wt. In normal fibrinogen I, Aα2 comprised 32% of the total Aα chains (range 37-41%). In contrast the fibrinogen I of 6 out of the 7 patients had a lower per cent of Aα2 (range 10-25%). Similar quantitative differences were seen in the decreased fraction of Aα1 in the fibrinogen II of cirrhotic patients compared to normals. No correlation was found between per cent fibrinogen II and per cent Aα2 in either normal subjects or cirrhotics. These results suggest that at least two independent processes are responsible for the observed levels of Aα chain heterogeneity in normals and cirrhotics and that one of these processes yields a lower than normal fraction of Aα2 chains in the fibrinogen of cirrhotic individuals.


A prolonged thrombin and reptilase time with normal fibrinogen concentration in a 9-years old girl, with no history of bleeding, prompted us to study the behaviour of her fibrinogen. Thrombin and reptilase time in different conditions, immunoelectrophoresis, immunodiffusion and fibrin polymerisation were performed. The thrombin time was partially corrected by calcium, ionic strength and increasing concentration of thrombin. Fibrin polymerisation rate and monomer aggregation time were moderately abnormal. Immunodiffusion and immunoelectrophoresis showed lines of identity with the normal. The electrophoretic mobility of the γ and β fibrin chains was normal, but there was a larger amount of a chain left in the cross-linked fibrin, in the presence of factor XIII. The above results suggest the presence of another fibrinogen variant.