Brief Communication

VARIATION OF HORSE PREALBUMINS IN ACIDIC STARCH GELS

Working with acidic starch gels (pH 5.9) Brænd & Efremov (1965) detected a large number of horse serum protein zones migrating faster than the albumins. In the present communication these proteins shall be called acidic prealbumins or just prealbumins.

The technique employed is based upon *Poulik*'s (1957) discontinuous system. The material comprised serum and plasma samples from 30 horses representing a variety of breeds which were tested monthly over half a year. In addition three horse families consisting of 30, 9 and 5 dam-offspring pairs respectively were investigated.

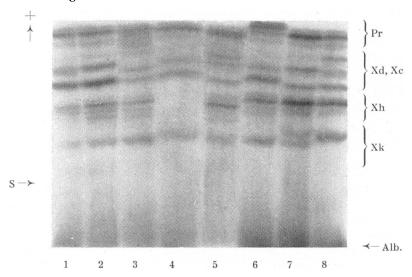


Figure 1. Prealbumins of 8 horses, gel buffer pH 5.0.

The appearance of acidic prealbumins is shown in Fig. 1. The fastest migrating are those studied by *Gahne* (1966) by him called the Pr system. Studies in our laboratory indicate that these proteins correspond to α_1 -antitrypsin.

The rest of the prealbumins are assumed to represent a variety of protein systems. They have been given tentative names Xk, Xh,

etc. where X stands for the unknown and the second letter for the order of migration. The slowest one is the Xk system where four different phenotypes were diagnozed. Most common is the KK phenotype which appears as one band only. Most of the samples in Fig. 1 are KK. A faster phenotype, FF, is seen in sample no. 8 and a two band, FK, type in no. 7. Another two band phenotype, KS, is seen in nos. 1 and 2. The position of the S band, which is rather faint on the photograph is marked with an arrow. These results suggested a genetic theory of three codominant alleles, Xk^F, Xk^K and Xk^S. Family studies agree with the theory.

In front of the Xk proteins another group of protein zones can be seen. These are assumed to represent another system, Xh. Most of the samples in Fig. 1 show the same Xh phenotype designated HH. A different phenotype, LL, is shown in no. 5. A combination between these two has been found as well as other phenotypes.

Between the Pr and Xh systems a rather great number of different protein bands have been found. The studies undertaken so far suggest two variable and partly overlapping systems tentatively called Xd and Xc.

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