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**Comparative Immunohistochemical Analysis of *Antheraea pernyi* Arylphorin with Storage Proteins of the Wild Silkworms, *Antheraea yamamai*, *Samia cynthia pryeri* and *Actias gnoma***

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The occurrences of proteins related to *A. pernyi* arylphorin in haemolymph, fatbody, integument, mid gut and silk gland of the wild silkworms, *A. yamamai*, *A. pernyi*, *S. cynthia pryeri* and *A. gnoma* were analysed by immunohistochemical analysis using mouse polyclonal antibody for *A. pernyi* arylphorin as probe. In *A. yamamai*, *A. pernyi* and *A. gnoma*, the major immunoreactive proteins with a molecular weight of 80kDa were observed in the haemolymph, integument, fat body and mid gut except silk gland. In *S. cynthia pryeri*, the immunoreactive proteins with a molecular weight of 80kDa were also observed in the haemolymph, integument and fat body except mid gut and silk gland. Also the *A. pernyi* arylphorin were detected immunohistochemically in cuticle, fat body, midgut, testis, posterior silk gland, middle silk gland, Malpighian tubule and ovary of the larva. These results reveals that the *A. pernyi* arylphorin has almost same identity with those of the wild silkworms, *A. yamamai*, *S. cynthia pryeri* and *A. gnoma* with respect to immunological reactivity and is distributed in several tissues and reproductive organs of the worm eventhough it is synthesized only in fat-body cells. In addition, it can be considered that the *A. pernyi* arylphorin has been evolved from a common ancestor gene of the wild silkworms described as the above.