

Arylphorin from Larval Hemolymph of the Chinese Oak Silkworm, *Antheraea pernyi*

Nam Sook Park¹, Jae Yu Moon², Su Il Seong³, Sang Mong Lee¹,
Soohyun kim⁴, Cristina Colominas⁵, Pauline M. Rudd⁵,
Byung Rae Jin⁶ and Jae Sam Whang⁷

¹Miryang National University, ²College of Agriculture & Life Science
Seoul National University, ³The University of Suwon, ⁴Biomolecule
Research Team Korea Basic Science Institute, ⁵Glycobiology
Institute, University of Oxford, ⁶Dong-A University, ⁷Dept. of
Sericulture and Entomology, The National Institute of Agricultural
Science and Technology, RDA

Arylphorin (APA; *Antheraea pernyi* Arylphorin) has been purified from the 5th instar larval hemolymph of the Chinese oak silkworm, *Antheraea pernyi*. The hemolymph contained a lot of proteins including a major storage protein. APA was purified by a simple preparative PAGE and diffusion. The preparation was shown to be homogeneous more than 95% by SDS-PAGE. The native molecular weight of APA was 460 kDa with a 80 kDa single subunit, suggesting hexamer. The protein had high amounts (17.7%, w/w) of phenylalanine and tyrosine. Rabbit antibody prepared against the purified protein cross-reacted with the 5th instar larval hemolymph proteins of *A. pernyi* and *A. yamamai*, but not with those of *Bombyx mori* and *B. mandarina*. With these results, it was concluded that APA was fall into as an arylphorin family having close relationship with *A. yamamai*. Arylphorins were known to be glycosylated. APA was stained by Schiff's reagent, suggesting a glycoprotein. Arylphorin had 4.8% (w/w) sugar, and mannose and *N*-acetylglucosamine were major components. PNGase-F digestion or hydrazinolysis produced the same high-performance liquid chromatographic profile of 2-aminobenzamide labeled glycans, suggesting APA had only *N*-linked oligosaccharides. Matrix-assisted laser desorption ionization-mass spectrometry analysis confirmed that the major glycans consisted of high-mannose type structures. Sequencing of glycan pool for further characterization showed that arylphorin had a glycan family of high-mannose type structures with and without core fucose residue. Interestingly, the most abundant form was Man₁₀GlcNAc₂, which was not described for *N*-glycan in insects. This work presents the characteristics of a purified arylphorin from *A. pernyi* and provided the first and most complete view of glycan structures of any arylphorins studied to date.