

Pirkle, 1977). One of these, batroxobin (from *Rothras* venoms) has been cloned and sequenced (Itoh et al., 1987; 1988). Peptide paranitroaniline (pNA) Chemical: Biochemical Corp. and Sigma

For a list of references, see the end of the book. For a list of the authors, see the end of the book.

was made for the heparin-like sites. The reaction mixture was dialyzed into 50 mM Tris-HCl and for the pH 8.0 and applied to a 1.5 cm x 75 cm using ProPheArgNA and PheValArg. Sephadex G-75 gel filtration column. Flow: thrombin-like protease using ions pNA. An appropriate amount of substrate rate was 4.8 ml/hr with four fractions. Fractions 50-60 containing was added to give a total volume of 675 μ l collected/hr.

erized. A thrombin-like enzyme, *unclonin*, has been characterized. The active site, characterized with a caseinolytic protease, is inhibited by PRACK (Fig. 9). A *S. aureus* β -caseinase, *clonin*, is also characterized with another caseinolytic protease. 100 nM PRACK results in complete inhibition of *clonin* activity ($K_i = 0.5$ nM) within one minute (Fig. 9).

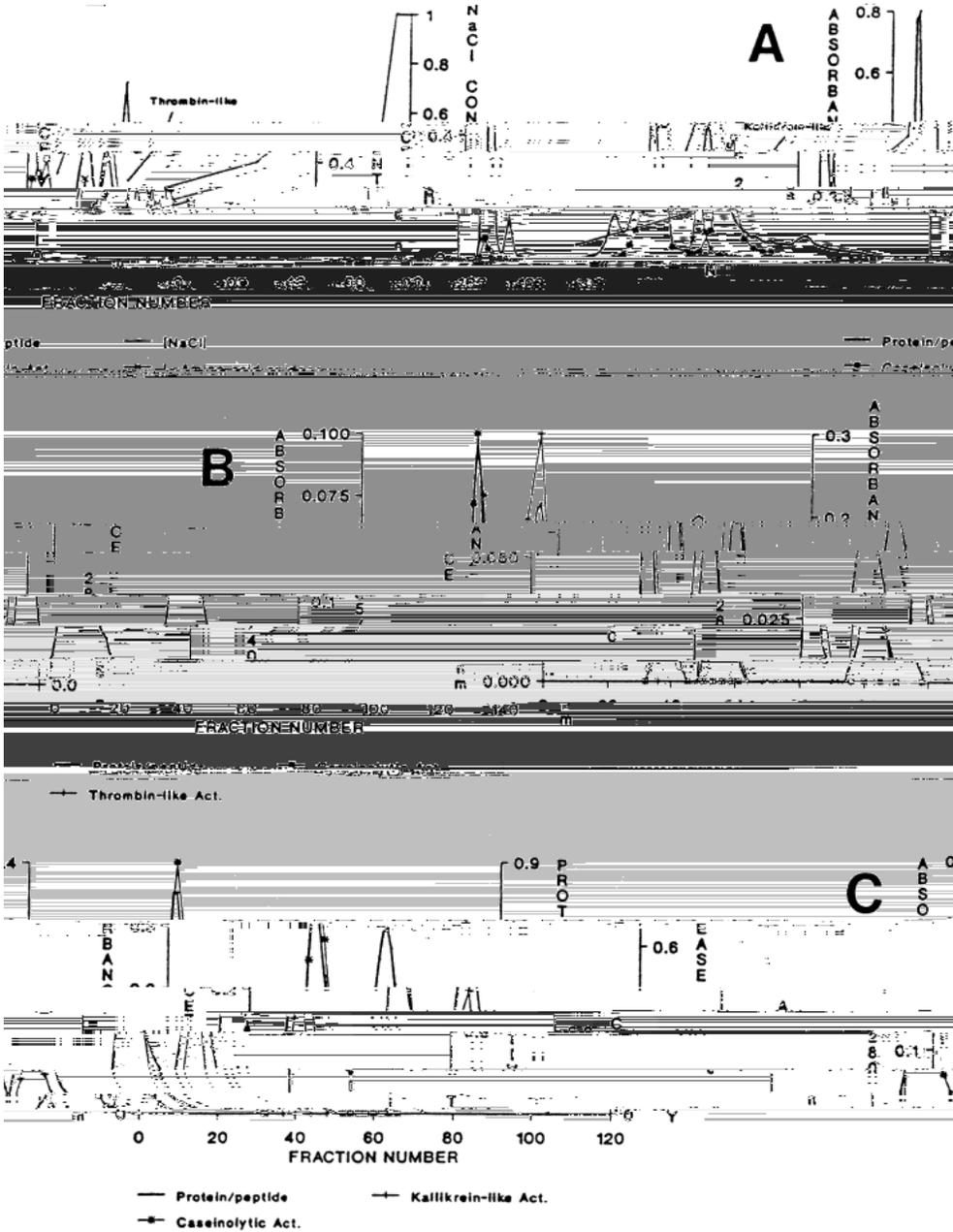


Fig. 1. Fractionation of combined fractions from *Urolophorus* venoms from *Urolophorus* *procerus* fractionated on a 1.5 cm x 25 cm DEAE-Sephacel ion exchange column. B. Elution profile of combined fractions from the thrombin-like protease peak chromatographed on a 1.5 cm x 75 cm Sephadex G-75 gel filtration column. Thrombin-like activity was completely separated from caseinolytic activity. C. Elution profile of combined fractions from a 1.5 cm x 75 cm DEAE-Sephacel ion exchange column. Kallikrein-like activity was completely separated from caseinolytic activity.

Hydrolyzed
Peptide

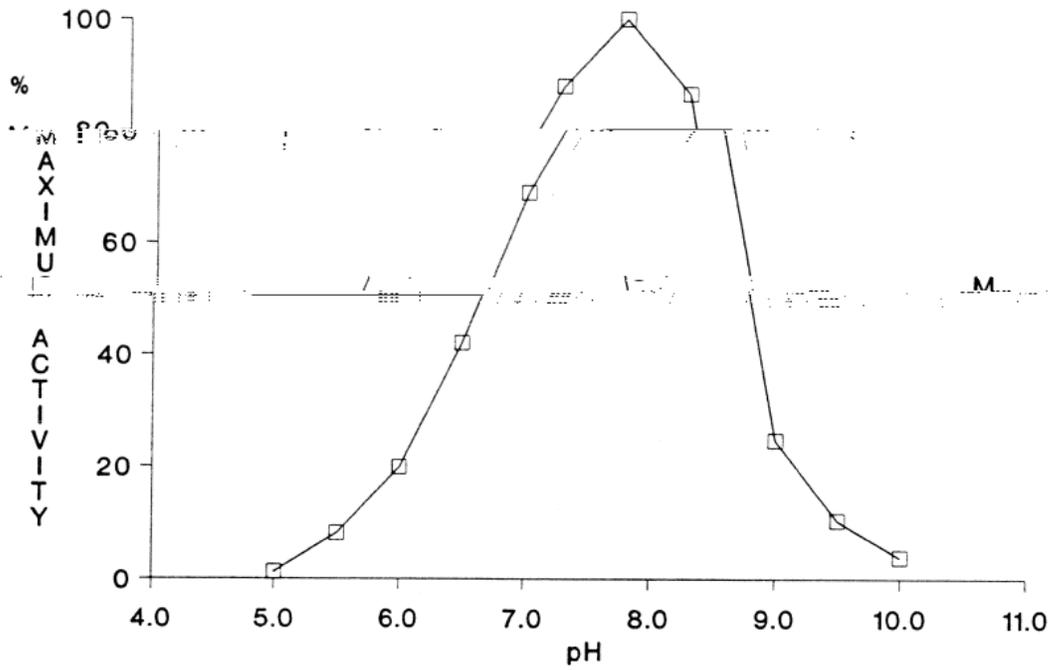


Figure 3. pH-profile for the hydrolysis of NBD-Casein by the adult mouse Chromobacterium (at pH 8.0). The activity is expressed as a percentage of the highest activity observed.

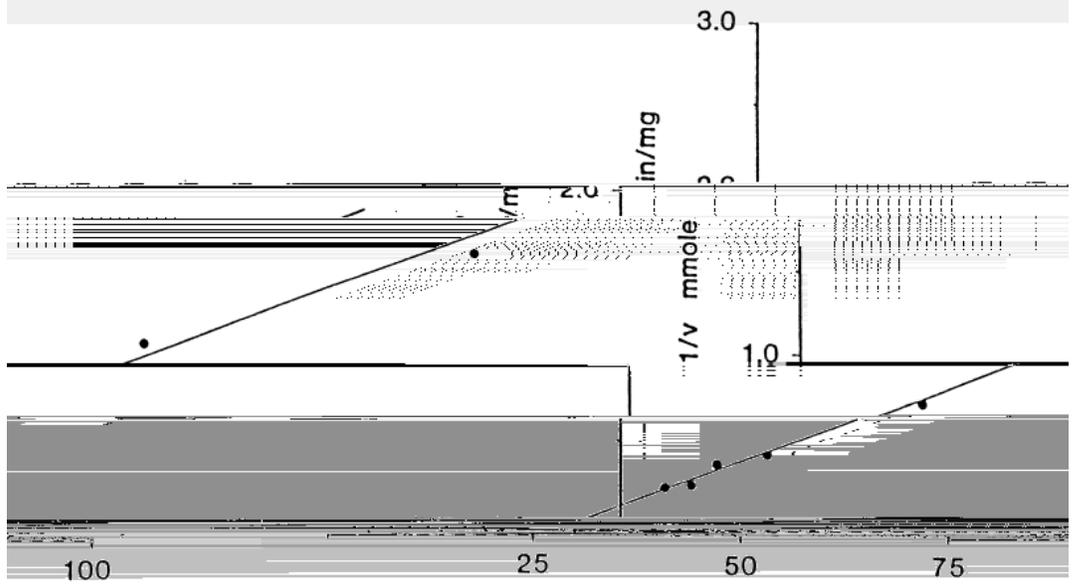


Figure 4. Double-reciprocal plot of $1/v$ vs. substrate concentration for the hydrolysis of NBD-Casein by the adult mouse Chromobacterium. Estimated kinetic constants: $K_m = 22 \mu M$, $V_m = 1.413 \text{ mmol/min/mg}$.

The kallikrein III
had a K_m of 0.00125 mM and a V_{max} of 0.000125 mM. It showed a high degree of specificity, as
hydrolyzed to a significant extent had \equiv kallikrein substrate ProPheArg-pNA with

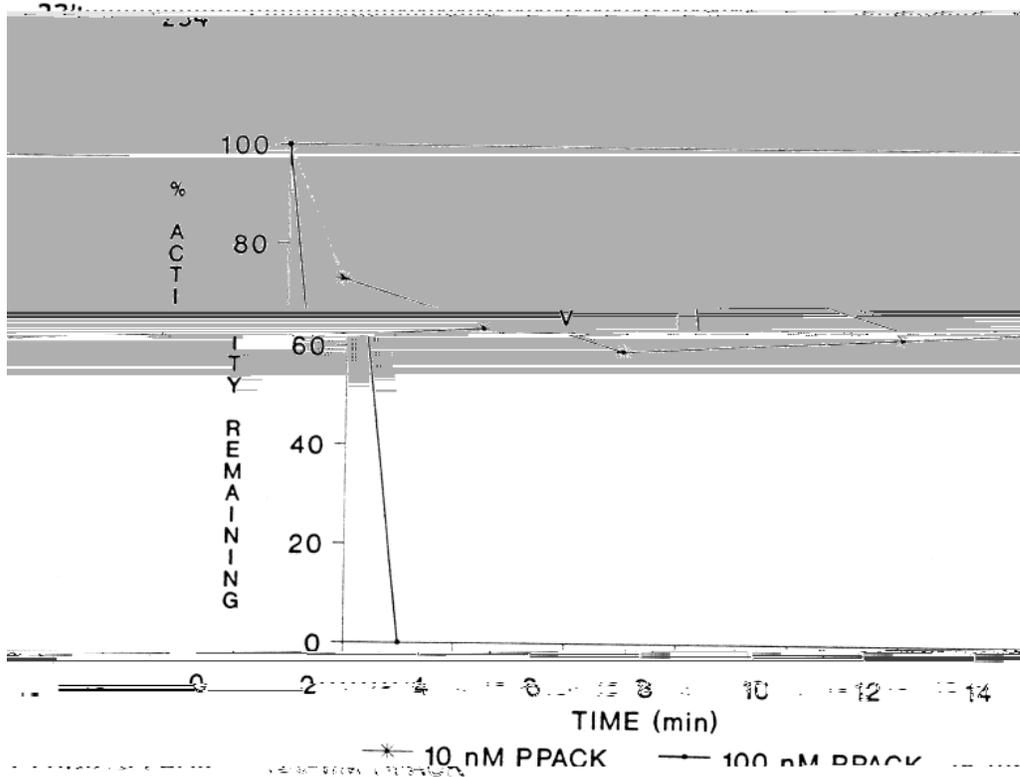


Figure 9
 Inhibition of vesicle-facilitated activity by PPACK. 100 nM PPACK resulted in the complete inhibition of ~0.5 nM enzyme within one minute.

It must be recalled that the kallikrein-like enzymes isolated from Crotalus veneniferus (Schwarz and Bröck, 1965) possess particular activities attributable to the venom. The approach of isolating venom constituents from *Bitis gabonica* venom (Vilicic, 1965) to study potential synergistic effects of venom constituents. In order to understand the biological role of venoms as biological

components, it is necessary to study the biological role of the venom components. The following sections discuss the biological role of the venom components.

resemble the summed effects of isolated components. Rattlesnake venoms have two major biological functions related to the venom components. Rather than being treated as individual independent components, they are treated as individual independent components.

They are analyzed for their potentially independent effects. Zeller (1977) has made the distinction between the primary importance and is accomplished by biochemistry with the view that venoms have evolved to serve specific biological functions. Zeller (1977) has made the distinction between the primary importance and is accomplished by biochemistry with the view that venoms have evolved to serve specific biological functions. Zeller (1977) has made the distinction between the primary importance and is accomplished by biochemistry with the view that venoms have evolved to serve specific biological functions.

in these venoms (Lee, 1979; Tu, 1983). In the past, the venom components have been studied in isolation. However, the biological role of the venom components is not understood. The biological role of the venom components is not understood. The biological role of the venom components is not understood.

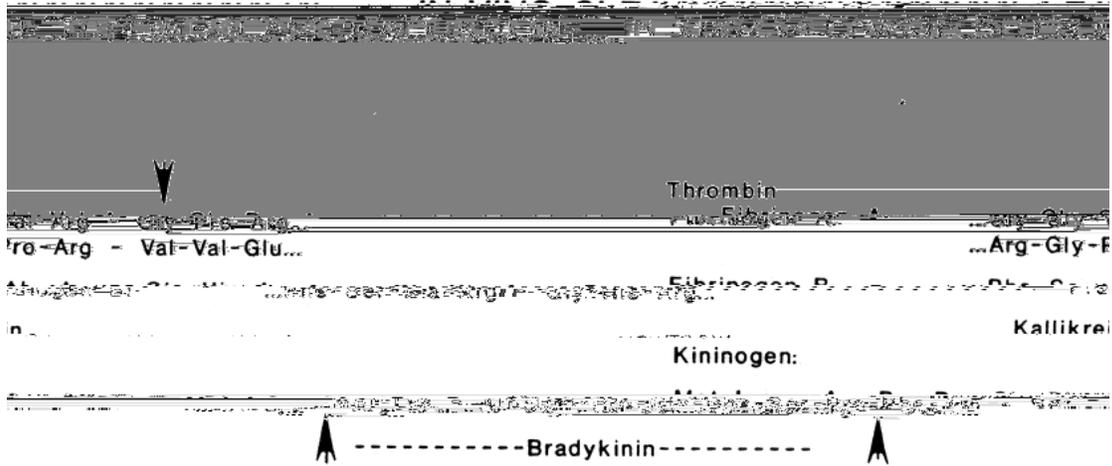


Figure 10. Several of the *in vivo* drug sites in cardiovascular pharmacology. Paranitrophenyl derivatives are based on these recognition sequences.

Proteolytic beginning toward the elucidation of these Bjarnason, J. B., and Fox, J. W. (1983).
 roles
 toxin e, a zinc protease isolated from the venom
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Roche e Silva, M., Beraldo, W. T., and Rosenfeld, Weisbach, H., Robertson, A. V., Witkop, B., and
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