

Effect of estrogens and gibberellic acid ( $GA_3$ ) on the growth of seedlings and cytokinin activity in extracts from 100 g of pea tissues

	Value in	Control	Estrone	Estradiol	Estriol	$GA_3$
Height <sup>a</sup> of pea seedlings	(mm)	70.8	99.6	100.4	98.1	98.7
	(%)	100.00	140.67	141.80	138.55	139.40
Yield of fresh tobacco tissue	(g/flask)	0.26	0.55	0.46	0.35	0.28
	(%)	100.00	211.53	176.92	134.61	107.69
Kinetic controls ( $\mu\text{g/l}$ )	0–0.12 <sup>b</sup>					
	1–0.35					
	5–0.76					
	25–3.16					

<sup>a</sup>96 h after application. <sup>b</sup>Yield of fresh tobacco tissue – (g/flask).

pea seedlings deprived of cotyledons 24, 48, 72 (AbA) and 96 h (cytokinins, AbA) after application.

AbA-like substances were extracted and fractionated according to the method described by RUDNICKI<sup>16</sup>. TLC with benzene-acetone-acetic acid (70:30:1) as solvent was used. The abscisic acid zone ( $R_f$  0.35–0.45) was then rechromatographed on Whatman No. 3 paper in redistilled water. The content of inhibitor was estimated by the wheat section straight growth test<sup>17</sup>. Growth inhibition was expressed in activity units. As activity unit 10% growth inhibition of the test plants in relation to control was taken. Cytokinins were determined according to HEIDE and SKOOG method<sup>18</sup>. A cation exchange column (250 ml, Dowex 50 W-X4 H<sup>+</sup> 50–100 mesh) was used for separation. Cytokinin activity was measured by the tobacco callus bio-assay<sup>19</sup>.

**Results and discussion.** The increase of 40% in the growth of the seedlings treated with estrogens and  $GA_3$  was observed 96 h after application (Table). Thus the two kinds of different hormones had an identical physiological effect. Our results (Table, Figure) show the positive effect of estrogens on the endogenous cytokinins content and the lack of the influence on the level of AbA-like substance. On the other hand, however, gibberellic acid, while showing the same final physiological effect, lowered the AbA amounts and did not change the cytokinins content. So it seems possible that estrogens and gibberellins effect the plant metabolism in various ways. The results obtained confirm also a relationship between estrogens and other plant hormones in regulating the growth and development processes in plants. The previous<sup>12, 13</sup> and the present papers show that estrogenic hormones influence the content of auxins, gibberellins and cytokinins in plant

tissues. This may be the cause of many important metabolic reactions. The investigations of the interrelations between steroid hormones and other groups of active substance were carried out in order to detect the strict control of growth and development processes which may exist in plants through the combined action of several regulatory substances.

**Zusammenfassung.** Die mit Oestrogenen (0.1  $\mu\text{g}$  pro Pflanze) und Gibberellinsäure ( $GA_3$ , 0,001  $\mu\text{g}$  pro Pflanze) behandelten Erbsenkeimlinge zeigten nach 96 h ein um 40% stärkeres Längenwachstum als die Kontrollpflanzen. Die Oestrogene erhöhten in den Keimlingen den Gehalt an Cytokininen, übten jedoch keinen Einfluss auf den Abscisinsäure-Gehalt aus. Die Gibberellinsäure andererseits setzte den Gehalt an Abscisinsäure herab, ohne die Konzentration der Cytokinine zu verändern.

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## STUDIORUM PROGRESSUS

### A Generalized Homology Correlation for Various Hormones and Proteins

An important archetypal connection between glucagon and secretin has been demonstrated and, in view of the disparities in both function and formation site for these two hormones, it was suggested that a search be instigated for other, less obvious genetic and biological relationships<sup>1</sup>. Along these lines, a computer alignment was then made between the above pair and pituitary and placental lactogen hormones<sup>2</sup>. The results were interesting, but inconclusive, since the complete sequences of the latter proteins were not known at the time. Recently, a different approach, based on glandular origin, was used to rank

various enterosecretory proteins<sup>3</sup>. Here the original glucagon-secretin correlation was extended to include portions of two growth hormones. The relationships seemed probable, yet suffered from the use of erroneous primary sequences and the restriction of homology to short, selected regions. In view of continual interest in this area, a new treatment differing in various details, positioning points, and area of coverage is now presented in the Table.

Some comments are necessary on the specific primary structures selected for the various compounds. In rough

order, the data for equine growth hormone (EGRO)<sup>4</sup>, ovine lactogen (OLAC)<sup>5</sup>, porcine pepsinogen (PPSN)<sup>6, 7</sup>, bovine pepsinogen (BPSN)<sup>8</sup>, bovine rennin (BREN)<sup>9, 10</sup>, human, porcine, and salmon calcitonin (HCAL, PCAL, SCAL)<sup>11-15</sup>, porcine  $\beta$ -corticotropin (PCTP)<sup>16</sup>, mouse nerve growth factor (MNGF)<sup>17</sup>, human, porcine, bovine, and rat proinsulin (HPIN, PPIN, BPIN, RPIN-1, RPIN-2)<sup>18-23</sup>, angler fish insulin (AFIN)<sup>24</sup>, bovine parathyroid hormone (BPTH)<sup>25</sup>, ovine, bovine, porcine, and human growth hormone (OGRO, BGRO, PGRO, HGRO)<sup>26-38</sup>, human placental lactogen (HPLA)<sup>39</sup>, bovine  $\alpha$ - and  $\beta$ -thyrotropin (BTT $\alpha$ , BTT $\beta$ )<sup>40</sup>, ovine  $\alpha$ - and  $\beta$ -luteinizing hormone<sup>41-43</sup>, human  $\alpha$ - and  $\beta$ -chorionic gonadotropin (HCGT- $\alpha$  and - $\beta$ )<sup>44</sup>, human  $\alpha$ -luteinizing hormone (HLUT- $\alpha$ )<sup>45, 47</sup>, bovine  $\beta$ -luteinizing hormone (BLUT- $\beta$ )<sup>48</sup>, porcine cholecystokinin-pancreozymin (PCPZ)<sup>49</sup>, frog caerulein (FCRN)<sup>50</sup>, canine, porcine, ovine, and human gastrin (CGAS, PGAS, OGAS, HGAS)<sup>51-54</sup>, porcine motilin (PMOT)<sup>55, 56</sup>, porcine vasoactive intestinal polypeptide (PVIP)<sup>57</sup>, porcine secretin (PSEC)<sup>58</sup>, porcine glucagon (PGLU)<sup>59</sup>, and porcine gastric inhibitory peptide (PGIP)<sup>60</sup> are taken from the literature.

Bovine prolactin differs from ovine prolactin by having alanine at position 112 and a tyrosine at 151<sup>61, 62</sup>, while bovine growth hormone is known to possess a leucine-valine interchange at 141<sup>63, 64</sup>. Bovine  $\alpha$ -luteinizing hormone is apparently identical to bovine  $\alpha$ -thyrotropin<sup>65, 66</sup>, and a similarity has been discussed for the ovine com-

pound, too<sup>67</sup>. The ovine hormone differs only at residue 3; however, evidence has been presented for some allele substitutions in the bovine hormone present at amino acids 106, 109, and 115<sup>68, 69</sup>. Bovine  $\beta$ -luteinizing hormone seems the same as the ovine hormone, yet a few residues remain uncertain at the present time. Another interesting

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	31	35	40	45	50	55	60																												
OLAC	Asn	Leu	Ser	Ser	Glu	Met	Phe	Asn	Glu	Phe	Asp	Lys	Arg	Tyr	-	Ala	Gln	Gly	Lys	Gly	Phe	-	-	Ile	Thr	Met	Ala	Leu	Asn	Ser					
PPSN	Leu	Lys	Asp	Phe	Leu	Lys	Thr	His	Lys	-	His	Asn	Pro	Ala	Ser	Lys	Tyr	Phe	-	-	Pro	Ala	Glu	Ile	Gly	Asp	Glu	Pro	Leu	Glu					
BPSN	Leu	Lys	Glu	Phe	Leu	Lys	Thr	His	Lys	Val	Arg	Asn	Met	Gly	Ser	Lys	Tyr	Leu	Ile	Arg	Glu	Ala	Ala	Thr	Leu	Ser	Val								
BREN	Leu	Glu	Asp	Phe	Leu	Lys																													
HCAL	Cys	Met	Leu	Gly	Thr	Tyr	Thr	Gln	Asp	Phe	Asn	Lys	Phe	His	Thr	Phe	Pro	Gln	Thr	Ala	Ile	Gly	Val	Gly	Ala	Pro									
PCAL	Cys	Val	Leu	Ser	Ala	Tyr	Trp	Arg	Asn	Leu	Asn	Asn	Phe	His	Arg	Phe	Ser	Gly	Met	Gly	Phe	Gly	Pro	Glu	Thr	Pro									
SCAL	Cys	Val	Leu	Gly	Lys	Leu	Ser	Gln	Glu	Leu	His	Lys	Leu	Gln	Thr	Tyr	Pro	Arg	Thr	Asn	Thr	Gly	Ser	Gly	Thr	Pro									
PCTP	Trp	Gly	Lys	Pro	Val	Gly	Lys	Lys	Arg	Arg	Pro	Val	Lys	Val	Tyr	Pro	Asn	Gly	Ala	Glu	Asp	Glu	Leu	Ala	Glu	Ala	Phe	Pro	Leu	Glu					
MNGF	Glu	Phe	Ser	-	-	Val	Cys	Asp	Ser	Val	Ser	Val	Trp	Val	Gly	Asp	Lys	Thr	Ala	Thr	Asn	Ile	Lys	Gly	Lys	Glu	Val	Thr	Val						
HPIN	Glu	Ala	Leu	Tyr	Leu	Val	Cys	Gly	Glu	Arg	Gly	Phe	Phe	Tyr	Thr	Pro	Lys	Thr	Arg	Arg	Glu	Ala	Glu	Asp	Leu	Gln	Val	Gly							
PPIN	Glu	Ala	Leu	Tyr	Leu	Val	Cys	Gly	Glu	Arg	Gly	Phe	Phe	Tyr	Thr	Pro	Lys	Ala	Arg	Arg	Glu	Val	Glu	Gly	Pro	Gln	Val	Gly	Ala	Leu					
SPIN	Glu	Ala	Leu	Tyr	Leu	Val	Cys	Gly	Glu	Arg	Gly	Phe	Phe	Tyr	Thr	Pro	Lys	Ala	Arg	Arg	Glu	Ala	Glu	Asn	Pro	Gln	Ala	Gly	Ala	Leu					
RPIN-1	Glu	Ala	Leu	Tyr	Leu	Val	Cys	Gly	Glu	Arg	Gly	Phe	Phe	Tyr	Thr	Pro	Lys	Ser	Arg	Arg	Glu	Val	Glu	Asp	Pro	Gln	Val	Pro	Gln	Leu					
RPIN-2	Glu	Ala	Leu	Tyr	Leu	Val	Cys	Gly	Glu	Arg	Gly	Phe	Phe	Tyr	Thr	Pro	Met	Ser	Arg	Arg	Glu	Val	Glu	Asp	Pro	Gln	Val	Ala	Gin	Leu					
AFIN	Asp	Ala	Leu	Tyr	Leu	Val	Cys	Gly	Asp	Arg	Gly	Ph	Phe	Tyr	Asn	Pro	Lys																		
BPTH	Ser	Met	Glu	Arg	Val	Glu	Trp	Leu	Arg	Lys	Leu	Gln	Asp	Val	His	Asn	Phe	Val	Ala	Leu	Gly	Ala	Ser	Ile	Ala	Tyr	Arg	Asp	Gly						
OGRO	Gln	Leu	Ala	Ala	Asp	Thr	Phe	Lys	Glu	Phe	Glu	Arg	Thr	Tyr	Ile	Pro	Glu	Gly	Gln	Arg	Tyr	Ser	-	Ile	Gln	Asn	Thr	Gln	Val	Ala					
BGRO	Gln	Leu	Ala	Ala	Asp	Thr	Phe	Lys	Glu	Phe	Glu	Arg	Thr	Tyr	Ile	Pro	Glu	Gly	Gln	Arg	Tyr	Ser	-	Ile	Gin	Asp	Thr	Gln	Val	Ala					
HGRO	Gln	Leu	Ala	Phe	Asp	Thr	Tyr	Gln	Glu	Phe	Glu	Ala	Tyr	Ile	Pro	Lys	Glu	Gln	Lys	Tyr	Ser	Ph	Leu	Gln	Asp	Pro	Glu	Thr	Ser						
HPLA	Gln	Leu	Ala	Ile	Asp	Thr	Tyr	Gln	Glu	Phe	Glu	Thr	Tyr	Ile	Pro	Lys	Asp	Gln	Lys	Tyr	Ser	Ph	Leu	His	Asp	Ser	Glx	Thr	Ser						
BTTa	Glu	Phe	Thr	Met	Gln	Gly	Cys	Pro	Gln	Cys	Lys	Leu	Lys	Glu	Asn	Lys	Tyr	Phe	Ser	Lys	Pro	Asp	Ala	Pro	Ile	Tyr	Gln	Cys	Met	Gly					
HCGT-a	Ala	Pro	Asx	Val	Glx	Asx	Cys	Pro	Glx	Cys	Thr	Leu	Glx	Glx	Asx	Pro	Phe	Ph	Ser	Glx	Pro	Gly	Ala	Pro	Ile	Leu	Glx	Cys	Met	Gly					
HLUT-a				Val	Gln	Asp	Cys	Pro	Glu	Cys	Thr	Leu	Gln	Glu	Asn	Pro	Phe	Ph	Ser	Gln	Pro	Gly	Ala	Pro	Ile	Leu	Gln	Cys	Met	Gly					
HCGT-B	Ala	Val	Glx	Lys	Glx	Gly	Cys	Pro	Val	Cys	Ile	Asn	-	-	-	-	-	-	Val	-	Thr	Thr	Ile	Cys	Aia	Gly	Tyr	Cys	Pro	Thr	Met	Thr			
BLUT-B	Ala	Ala	Glu	Lys	Glu	Ala	Cys	Pro	Val	Cys	Ile	Thr	-	-	-	-	-	-	Phe	Thr	Thr	Ser	Ile	Cys	Ala	Gly	Tyr	Cys	Pro	Ser	Met	Lys			
PLUT-B	Ala	Ala	Glx	Asx	Glx	Ala	Cys	Pro	Val	Cys	Ile	Thr	-	-	-	-	-	-	Phe	Thr	Thr	Ser	Ile	Cys	Ala	Gly	Tyr	Cys	Pro	Ser	Met	Arg			
BBTB	His	Val	Glu	Arg	Lys	Glu	Cys	Ala	Tyr	Cys	Leu	Thr	-	-	-	-	-	-	Ile	Asn	Thr	Thr	Val	Cys	Ala	Gly	Tyr	Cys	Met	Thr	Arg	Asx			
PCPZ	Gln	Ser	Leu	Asp	Pro	Ser	His	Arg	Ile	Ser	Asp	Arg	Asp	Tyr	Met	Gly	Trp	Met	Asp	Ph															
FCRN							<Glu	Gln	Asp	Tyr	Thr	Gly	Trp	Met	Asp	Ph																			
CGAS							Glu	Gly	Pro	Trp	Met	Glu	Glu	Ala	Glu	Ala	Tyr	-	Gly	Trp	Met	Asp	Ph												
PGAS							Glu	Gly	Pro	Trp	Met	Glu	Glu	Glu	Glu	Ala	Tyr	-	Gly	Trp	Met	Asp	Ph												
OGAS							Glu	Gly	Pro	Trp	Val	Glu	Glu	Glu	Glu	Ala	Ala	Tyr	-	Gly	Trp	Met	Asp	Ph											
HGAS							Glu	Gly	Pro	Trp	Leu	Glu	Glu	Glu	Glu	Ala	Tyr	-	Gly	Trp	Met	Asp	Ph												
PMOT	Tyr	Gly	Glu	Leu	Gln	Arg	Met	Glu	Glu	Lys	Glu	Arg	Asn	Lys	Gly	Gin																			
PVIP	Leu																																		
PSEC	Leu	Val																																	
PGLU	Leu	Met	Asn	Thr																															
PGIP	Leu	Leu	Ala	Gln	Gln	Lys	Gly	Lys	Lys	Ser	Asp	Trp	Lys	His	Asn	Ile	Thr	Gln																	

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	61	65	70	75	80	85	90
OLAC	-	Cys His Thr Ser Ser Leu Pro Thr Pro Glu Asp Lys Glu Gln Ala Gln Gln Thr His His Glu Val Leu Met Ser Leu Ile Leu Gly					
PPSN	Asn	Tyr Leu Asx Thr Glu Tyr Phe					
BREN	Asn	Tyr Leu Asx Ser Glx Tyr Phe Gly Lys Ile Thr					
PCTP		Phe					
MNGF	Leu Ala Glu Val Asn Ile Asn Asn Ser Val Phe Arg Gln Tyr Phe Phe Glu Thr Lys Cys Arg Ala Ser Asn Pro Val Glu Ser Gly Cys						
HPIN	Glu Leu Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu Ala Leu Glu Gly Ser Leu Gln Lys Arg Gly Ile Val Gly Gln Cys Cys						
PPIN	Glu Leu Gly Gly Gly Leu Gly - - - Gly Leu Gln Ala Leu Ala Leu Glu Gly Pro Pro Gln Lys Arg Gly Ile Val Glu Gln Cys Cys						
BPIN	Glu Leu Ala Gly Gly Pro Gly Ala Gly Gly - - - - Leu Glu Gly Pro Pro Gln Lys Arg Gly Ile Val Glu Gln Cys Cys						
RPIN-1	Glu Leu Gly Gly Gly Pro Glu Ala Asp Gly Leu Gln Thr Leu Ala Leu Glu Val Ala Arg Gln Lys Arg Gly Ile Val Asp Gln Cys Cys						
RPIN-2	Glu Leu Gly Gly Gly Pro Gly Ala Asp Gly Leu Gln Thr Leu Ala Leu Glu Val Ala Arg Gln Lys Arg Gly Ile Val Asp Gln Cys Cys						
AFIN							
BPTH	Ser Ser Gln Arg Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Gln Lys Ser Leu Gly Glu Ala Asp Lys Ala Asp Val Asp						
OGRO	Phe Cys Phe Ser Glu Thr Ile Pro Ala Pro Thr Gly Lys Asn Glu Ala Gln Gln Lys Ser Asp Leu Glu Leu Arg Ile Ser Leu Leu						
BGRO	Phe Cys Phe Ser Glu Thr Ile Pro Ala Pro Thr Gly Lys Asn Glu Ala Gln Glu Lys Ser Asp Leu Glu Leu Arg Ile Ser Leu Leu						
HGRO	Leu Cys Phe Ser Glu Ser Ile Pro Thr Pro Ser Asn Arg Glu Glu Thr Gln - - Lys Ser Asn Leu Gln Leu Leu Arg Ile Ser Leu Leu						
HPLA	Phe Cys Phe Ser Asx Ser Thr Pro Thr Pro Ser Asx Met Gix Glx Thr Glx - - Lys Ser Asx Leu Gln Leu Leu Arg Ile Ser Leu Leu						
BBT <sub>a</sub>	Cys Cys Phe Ser Arg Ala Tyr Pro Thr Pro Ala Arg Ser Lys Lys Thr Met Leu Val Pro Lys Asn Ile Thr Ser Glu Ala Thr Cys Cys						
HCGT- $\alpha$	Cys Cys Phe Ser Arg Ala Tyr Pro Thr Pro Leu Arg Ser Lys Lys Thr Met Leu Val Gln Lys Asn Val Thr Ser Gix Ser Thr Cys Cys						
HLT- $\alpha$	Cys Cys Phe Ser Arg Ala Tyr Pro Thr Pro Leu Arg Ser Lys Lys Thr Met Leu Val Gln Lys Asn Val Thr Ser Gix Ser Thr Cys Cys						
HCGT- $\beta$	Arg Val Leu Gln Gly Val Leu Pro Ala Leu Pro Gln Leu - - Val Cys Asx Tyr Arg Asx Val Arg Phe Gix Ser Ile Arg Leu Pro						
BLUT- $\beta$	Arg Val Leu Pro Val Ile Leu Pro Pro Met Pro Gln Arg - - Val Cys Thr Tyr His Glu Leu Arg Phe Ala Ser Val Arg Leu Pro						
PLUT- $\beta$	Arg Val Leu Pro Ala Ala Leu Pro Pro Val Pro Gln - - Pro Val Cys Thr Tyr Arg Glu Leu Ile Phe Ala Ser Ser Arg Leu Pro						
BBT <sub>b</sub>	Val Asx Gly Lys Leu Phe Leu Pro Lys Tyr Ala Leu Ser Gln Asp Val Cys Thr Tyr Arg Asp Phe Met Tyr Lys Thr Ala Glu Ile Pro						

retention of sequence occurs with glucagon, where the bovine and human forms are the same as the porcine material<sup>70,71</sup>. The different heterogenetic modifications are noted in the alignment framework, but the various duplicate sequences have been omitted at this time. The use of gaps was held to a minimum between completely different proteins; however, the device was employed judiciously within similar groups.

The homology pattern for the placental and growth hormones, such as EGRO, OLAC, OGRO, EGRO, PGRO, HGRO, and HPLA, is very obvious and has been discussed in part by other workers<sup>72-74</sup>. Attention is drawn to the point that the present Table is the first to include data from all of these compounds. Subsequently, several improvements are apparent at this time, which includes the detection of possible sequencing errors located at positions 93-94, 139-140, and 194-195 in the various growth compounds. One must note that the relationship of OLAC to this group of hormones in the 110-160 region is poor and may reflect major placement errors or even an incorrect primary sequence. However, the sequence of HCGT- $\alpha$  in this area is very helpful in partially locating a relationship with OLAC. In any event, it seems OLAC and EGRO represent older, more intact, proteins based on both their longer N-terminal sections and the high degree of individual residue changes. Indeed, the present resemblance between the two proteins at the N-terminal region suggests EGRO may be the equivalent ELAC, rather than the presumed horse growth hormone. Of interest is the degree of commonality displayed in the 12-25 region between HGRO, PVIP, PSEC, PGLU, and PGIP, which implies a strong familial relationship. The completion of the sequence work on PGRO would no doubt serve to confirm these assumptions.

The gastrins are a well-known group of hormones and present no problems in terms of homology. The very recent disclosure of a structure for cholecystokinin-pancreozymin permits one to locate caerulein and the gastrins at the C-terminal sequence of PCPZ. The similarity at the 38-44 section for HGRO-HPLA and the various gastrins serves as a second reference point and ties them into the various growth hormones. Based on these arguments, a rather interesting fact emerges; namely, secretin terminates at position 32, vasoactive intestinal

peptide stops at position 33, and glucagon ends at position 34, while the gastrins appear to commence in the same region. Such a break suggests the possible existence of a common zymogen, either past or present, which upon activation yields this family of biologically active hor-

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	91	95	100	105	110	115	120
OLAC	Leu Arg - Ser Trp Asn Asp Pro Leu Tyr His Leu Val Thr Glu Val Arg Gly Met Lys Gly Val Pro Asp Ala Ile Leu Ser Arg Ala						
MNGF	Arg Gly Ile Asp Ser Lys His - Trp Asn Ser Tyr Cys Thr Thr His Thr Phe Val Lys Ala Leu Thr Thr Asp Glu Lys Gin Ala						
HPIN	Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn						
PPIN	Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn						
BPIN	Ala Ser Val Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn						
RPIN-1	Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn						
RPIN-2	Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn						
AFIN	His Arg Pro Cys Asn Ile Phe Asp Leu Gln Asn Tyr Cys Asn						
BPTH	Val Leu Ile Lys Ala Lys Pro Gln						
OGRO	Leu Ile Gln Ser Trp Leu Gly Pro Leu Gln Phe Leu Ser Arg Val Phe Thr Asp Ser Leu Val Phe Gly Thr Ser Asp Arg - Val Tyr						
BGRO	Leu Ile Ser Glu Trp Leu Gln Pro - Gly Phe Leu Arg - Val Phe Thr Asn Ser Leu Val Phe Gly Thr Ser Asp Arg - Val Tyr						
HGRO	Leu Ile Gln Ser Trp Leu Glu Pro Val Gln Phe Leu Arg Ser Val Phe Ala Asn Ser Leu Val Tyr Gly Ala Ser Asn Ser Asp Val Tyr						
HPLA	Leu Ile Gln Ser Trp Leu Gln Pro Val Arg Phe Leu Arg Ser Met Phe Ala Asx Asx Leu Val Tyr Asx Thr Ser Asx Asx Ser Tyr						
BTT $\alpha$	Val Ala Lys Ala Phe Thr Lys Ala Thr Val Met Gly Asn Val Arg Val Gln Asn His Thr Glu Cys His Cys Ser Thr Cys Tyr Tyr His						
HGCT- $\alpha$	Val Ala Lys Ser Tyr Asx Arg Val Thr Val Met Gly Gly Phe Lys Val Gln Asn His Thr Ala Cys His Cys Ser Thr Cys Tyr Tyr His						
HLUT- $\alpha$	Val Ala Lys Ser Tyr Asn Arg Val Thr Val Met Gly Gly Phe Lys Val Gln Asn His Thr Ala Cys His Cys Ser Thr Cys Tyr Tyr His						
HGCT- $\beta$	Gly Cys Pro Arg Gly Val Asx Pro Val Val Ser Tyr Ala Val Ala - - - Leu - - - Cys Arg - Ser						
BLUT- $\beta$	Gly Cys Pro Pro Gly Val Asx Pro Thr Val Ser Phe Pro Val Ala - - - Leu Ser Cys His Cys Gly Pro Cys Arg Leu Ser						
PLUT- $\beta$	Gly Cys Pro Pro Gly Val Asx Pro Thr Val Ser Phe Pro Val Ala - - - Ile Ser Cys His Cys Gly Pro Cys Arg Leu Ser						
BBTB	Gly Cys Pro Arg His Val Thr Pro Tyr Phe Ser Tyr Pro Val Ala - - - Ile Ser Cys Lys Cys Gly Lys Cys Asx Thr Asx						
	121	125	130	135	140	145	150
OLAC	Ile Glu Ile Glu Glu Glu Asn Lys Arg Leu Leu Glu Gly Met Glu Met Ile Phe Gly Gln Val Ile Pro Gly Ala Lys Glu Thr Glu Pro						
MNGF	Ala Trp Arg Phe Ile Arg Ile Asn Thr Ala Cys Val Cys Val Leu Ser Arg Lys Ala Thr Arg						
OGRO	- - - - Ile Lys Leu Lys Asp Leu Ile Ile Gly Ile Leu Ala Leu Met Arg Glu Leu Glu Asp Val Thr Pro Arg Ala Gly Gln						
BGRO	- - - - Glu Lys Leu Lys Asp Leu Glu Glu Gly Ile - Ala Leu Met Arg Glu Val Glu Asp Gly Thr Pro Arg Ala Gly Gln						
PGRO	- - - - Asp Leu Leu Lys Asp Leu Glu Glu Gly Ile Glu Thr Leu Met Gly Arg Leu Glu Asp Gly Ser Pro Arg Ala Gly Gln						
HGRO	- - - - His Leu Leu Lys Asx Leu Gln Gln Ile Gln Thr Leu Met Gly Arg Leu Glu Asp Gly Ser Pro Arg Thr Gly Gln						
HPLA	- - - - His Leu Leu Lys Asx Leu Gln Gln Ile Gln Thr Leu Met Gly Arg Leu Glu Asp Gly Ser Arg Arg Thr Gly Gln						
BTT $\alpha$	Lys Ser						
HGCT- $\alpha$	Lys Ser						
HLUT- $\alpha$	Lys Ser						
HGCT- $\beta$	Thr Thr Asx Cys Gly Gly Pro Lys Asx His Pro Leu Thr Cys Asx Pro Arg Phe Gln Asx Ser Ser Ser Lys Ala Pro Pro Pro Ser						
BLUT- $\beta$	Ser Thr Asp Cys Gly Pro Gly Arg Thr Glu Pro Leu Ala Cys Asp His Pro Pro Leu Pro Asp Ile Leu						
PLUT- $\beta$	Ser Ser Asx Cys Gly Pro Gly Arg Ala Gln Pro Leu Ala Cys Asx Arg Pro Pro Leu Pro Gly Leu Leu						
BBTB	Tyr Ser Asx Cys Ile His Glu Ala Ile Lys Thr Asn Tyr Cys Thr Lys Pro Gln Lys Ser Tyr Met						
	151	155	160	165	170	175	180
OLAC	Tyr Pro Val Trp Ser Gly Leu Pro Ser Leu - Gln Thr Lys Asp Glu - Asp Ala Arg His Ser Ala Phe Tyr Asn Leu Leu His Cys						
OGRO	Ile Leu Lys Gln Thr Tyr Asp Lys Phe Asp Thr Asn Met Arg Ser Asp - Asp Ala Leu Lys Asn - Tyr Gly Leu Leu Ser Cys						
BGRO	Ile Leu Lys Gln Thr Tyr Asp Lys Phe Asp Thr Asn Met Arg Ser Asp - Asp Ala Leu Lys Asn - Tyr Gly Leu Leu Ser Cys						
PGRO	Ile Leu Lys Gln Thr Tyr Asp Lys Phe Asp Thr Asn Leu Arg Ser Asp - Asp Ala Leu Lys Asn - Tyr Gly Leu Leu Ser Cys						
HGRO	Ile Phe Lys Gln Thr Tyr Ser Lys Phe Asp Thr Asn Ser His Asn Asp - Asp Ala Leu Lys Asn - Tyr Gly Leu Leu Tyr Cys						
HPLA	Ile Leu Lys Gln Thr Tyr Ser Lys Phe Asx Thr Asx Ser His Asx Asx His Asx Ala Leu Lys Asx - Tyr Gly Leu Leu Tyr Cys						
HGCT- $\beta$	Leu Pro Ser Pro Arg Leu Pro Gly Pro Asx Thr Pro Ile Leu Pro Gln Ser Leu Pro						
	181	185	190	195	200	205	
EGRO							
OLAC	Leu Arg Arg Asp Ser Ser Lys Ile Asp Thr Tyr Leu Lys Leu Leu Asn Cys Arg Ile Ile Tyr Asn Asn Asn Cys						
OGRO	Phe Arg Lys Asp Leu His Lys Thr Glu Thr Tyr Leu Arg Val Met Lys Cys Arg Arg Phe Gly Glu Ala Ser Cys Ala Phe						
BGRO	Phe Arg Lys Asp Leu His Lys Thr Glu Thr Tyr Leu Arg Val Met Lys Cys Arg Arg Phe Gly Glu Ala Ser Cys Ala Phe						
HGRO	Phe Arg Lys Asp Met Asp Lys Val Glu Thr Phe Leu Arg Ile Val Gln Cys Arg - Ser Val Glu Gly Ser Cys Gly Phe						
HPLA	Phe Arg Lys Asx Met Asx Lys Val Gln Thr Phe Leu Arg Met Val Gln Cys Arg - Ser Val Glu Gly Ser Cys Gly Phe						

The amino acid sequences of equine growth hormone (EGRO), ovine lactogen (OLAC), porcine pepsinogen (PPSN), bovine pepsinogen (BPSN), bovine rennin (BREN), human calcitonin (HCAL), porcine calcitonin (PCAL), salmon calcitonin (SCAL), porcine  $\beta$ -corticotropin (PCTP), mouse nerve growth factor (MNGF), human proinsulin (HPIN), porcine proinsulin (PPIN), bovine proinsulin (BPIN), rat proinsulin-1 and -2 (RPIN-1 and -2), angler fish insulin (AFIN), bovine parathyroid hormone (BPTH), ovine growth hormone (OGRO), bovine growth hormone (BGRO), porcine growth hormone (PGRO), human growth hormone (HGRO), human placental lactogen (HPLA), bovine  $\alpha$ -thyrotropin (BTT $\alpha$ ), human  $\alpha$ -luteinizing hormone (HLUT- $\alpha$ ), human chorionic gonadotropin- $\alpha$  and - $\beta$  (HGCT- $\alpha$  and - $\beta$ ), bovine  $\beta$ -luteinizing hormone (BLUT- $\beta$ ), porcine  $\beta$ -luteinizing hormone (PLUT- $\beta$ ), bovine  $\beta$ -thyrotropin (BTT $\beta$ ), porcine cholecystokinin-pancreozymin (PCPZ), frog caerulein (FCRN), canine gastrin (CGAS), porcine gastrin (PGAS), ovine gastrin (OGAS), human gastrin (HGAS), porcine motilin (PMOT), porcine vasoactive intestinal polypeptide (PVIP), porcine secretin (PSEC), porcine glucagon (PGLU), and porcine gastric inhibitory polypeptide (PGIP) aligned for maximum external correspondence. Deletions are indicated with a dash.

mones. The name 'prosecogastrin' is assigned at this time to the hypothetical parent. Support for this latter proposal has already come from a different corner; namely, the detection of 'Big Gastrin', a polypeptide with a molecular weight of about 7,000 and possessing the same physiological properties as gastrin<sup>75</sup>. Here, on digestion with trypsin, a smaller gastrin-like peptide is released in the gut; thus, the

larger molecule may be the actual precursor protein. The recent report of a 'Big, Big Gastrin' lends further weight to this proposal<sup>76</sup>.

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Attention is next paid to the two bovine thyrotropins, as well as the various gonadotropins and luteinizing hormones. These proteins are related through various common sequences<sup>77,78</sup> and are fitted to placental lactogen at positions 61–63 and 68–70. It is recognized that these compounds have a larger number of differences with the growth hormones and are probably only distantly related at this time. The placement of BLUT- $\beta$  and BTT $\beta$  with respect to BTT $\alpha$  is based on common residues found at positions 112–117, yet can stand improvement in the initial N-terminal section. On turning to the various proinsulins, which includes nerve growth factor and probably parathyroid hormone and  $\beta$ -corticotropin, alignments are easily found, especially with the cysteines located at positions 37, 62, and 89–90. Angler fish insulin is included now since it contains both an additional residue and various changes in the N-terminus of the  $\beta$ -chain. The homology between these groups disappears rather rapidly after the 44–47 area, possibly as a consequence of the initiation of the connecting peptide sequence found in the insulins, or of the need to introduce gap areas in order to maintain the alignments. The recent detection of a proparathyroid hormone is of much interest<sup>79,80</sup>, since it possesses 15–20 additional residues, which presumably must lie to the left and right of the present parathyroid sequence and should support the present assignments. The inclusion of porcine pepsinogen, bovine pepsinogen, and bovine rennin at this time is based on their resemblance to the initial sections of glucagon and secretin, as well as similar regions in the various growth hormones. Finally, the three calcitonins appear to be generally related to both OLAC and the proinsulins.

The compounds discussed here must have arisen from a series of ancestral homotypes that evolved by the general mechanism of gene duplication. The original parent was probably present in the digestive tract and in time was modified to fit various roles through changes occurring in the gut and accessory regions. It is likely that the commonality in the first 30 residues for many of these compounds may be a result of the conservation of binding to a specific receptor site, followed by the activation of the adenylate cyclase system, or by the release of inorganic ions<sup>81</sup>, transmitter substances, and by serving at the level of transcription<sup>82</sup>. The empirical relationships established here clearly show that the various gastric hormones constitute a homologous block of compounds, which served in time to develop the insulins, and, ultimately

the growth and luteinizing hormones. On these grounds, one can find reduplicated sections beginning at positions 51, 91, and 131, suggestive of internal gene doubling patterns. Computer techniques that have been proposed for the testing of homology<sup>83–86</sup> will be used in the near future in an attempt to confirm these assignments. Even now indications exist that thrombin is a member of this large family. In summary, attention is called to the old folk proverb that maintains, 'the way to a man's heart is through his stomach'. This statement is undoubtedly true and probably should be modified to include other organs as well<sup>87</sup>.

**Résumé.** La calcitonine, la cholécystokinine, la motilin, la pancreozymine, le peptide inhibiteur gastrique, le peptide vasoactif, la gastrine, le glucagon, et la secretine sont comparables par leur composition en amino acides à l'hormone de croissance, à l'hormone luteinigène, la gonadotropine, au facteur de croissance nerveux, à l'hormone parathyroïdienne, au lactogène du placenta, à la proinsuline, à la rénine et à la thyrotropine. Ces peptides proviennent probablement d'un génotype dont l'origine se trouve dans le système digestif.

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## PRO EXPERIMENTIS

### The Purification of Peptic Antibody Fragments from Rabbit Immunoglobulin G

Univalent antibody fragments of the type Fab' are obtained by sequential treatment of immunoglobulins with pepsin and thiol compounds<sup>1</sup>. However, experience in our laboratory has shown that published methods do not produce physically homogeneous proteins. Therefore, we have devised a scheme that effectively uses gel permeation chromatography for purification<sup>2</sup>.

To 1 volume of rabbit serum was added saturated  $(\text{NH}_4)_2\text{SO}_4$  to 40% saturation<sup>3</sup>. The precipitated globulins were recovered by centrifugation (2,500 g for 10 min), dissolved in 1 volume water, and twice reprecipitated as before. The final precipitate was taken up in 1/6 volume water and extensively dialyzed against 15 mM potassium phosphate, pH 8.0. Insoluble material was removed by

centrifugation (8,000 g for 10 min). The supernatant was chromatographed on diethylaminoethyl (DEAE)-cellulose<sup>4</sup> using the same buffer. The immunoglobulin G (IgG) peak emerging in the void volume was dialyzed against 0.1 M sodium acetate, pH 4.5, and the contents of the dialysis bag were concentrated with Ficoll<sup>5</sup> to 15 mg/ml ( $A_{280 \text{ nm}}^{mg/ml}$  1.29).

To 500–750 mg IgG was added pepsin<sup>6</sup> at a pepsin-to-globulin ratio of 1:50. The mixture was gently agitated at 37 °C for 8 h. The reaction was stopped by raising the pH of the solution to 7.6 and by chilling. Insoluble material was removed by centrifugation (24,000 g for 10 min). After concentration the supernatant was applied to a column (2.8 × 45 cm) of Sephadex G-150 superfine<sup>5</sup>.