



# Promix™ Columns

## Two-Dimensional Separation of Proteins & Peptides

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The ever-increasing utility of proteomics in biotechnology has spurred demand for effective HPLC separation of mixtures involving proteins and peptides. However, analysis is complicated by the complexity and homology of these biomolecules.

**Promix™ Technology** enables two-dimensional separation of peptides and proteins for purification and proteomics applications. Columns combine both ionic and reverse-phase interactions to **maximize resolution**. Promix can separate peptides that differ in a single amino acid.

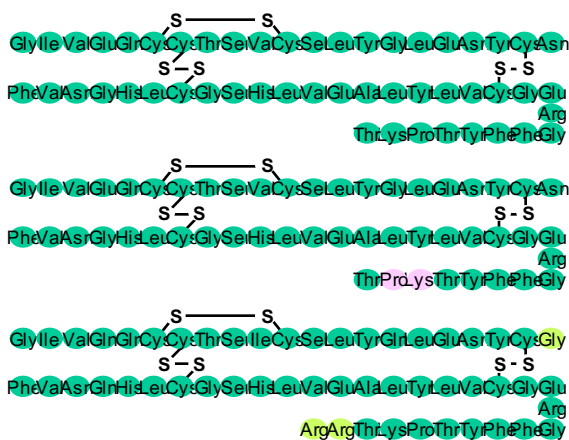
Promix offers **enhanced selectivity** for closely related peptides and proteins, and **increased peak capacity** compared to ion-exchange and reverse-phase columns alone. Promix is **completely scalable** from capillary to preparatory applications. Columns are compatible with formic acid and ammonium acetate, which makes them **perfectly suited for LC-MS**.

### Separation of Insulin Analogs

Synthetic insulin analogs differ from normal insulin by as little as a single amino acid pair (right).

Combining ionic and hydrophobic interactions enables Promix MP to separate these analogs by exploiting slight differences in isoelectric charge (pI) (bottom).

Independent adjustment of ionic and organic concentrations provides a diverse array of separation conditions.



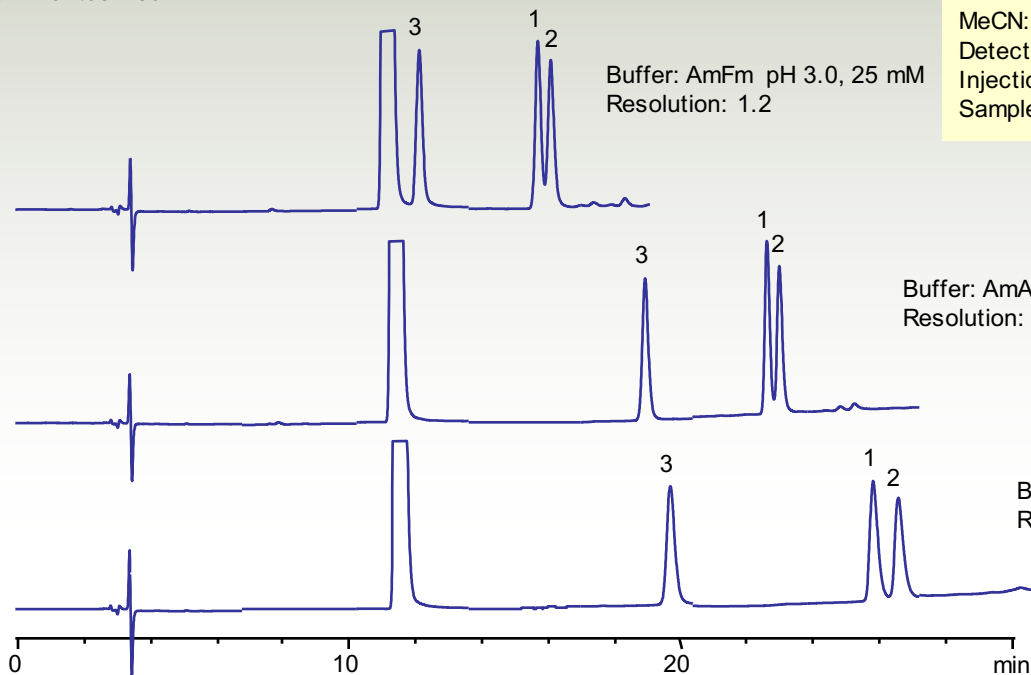
Normal human insulin

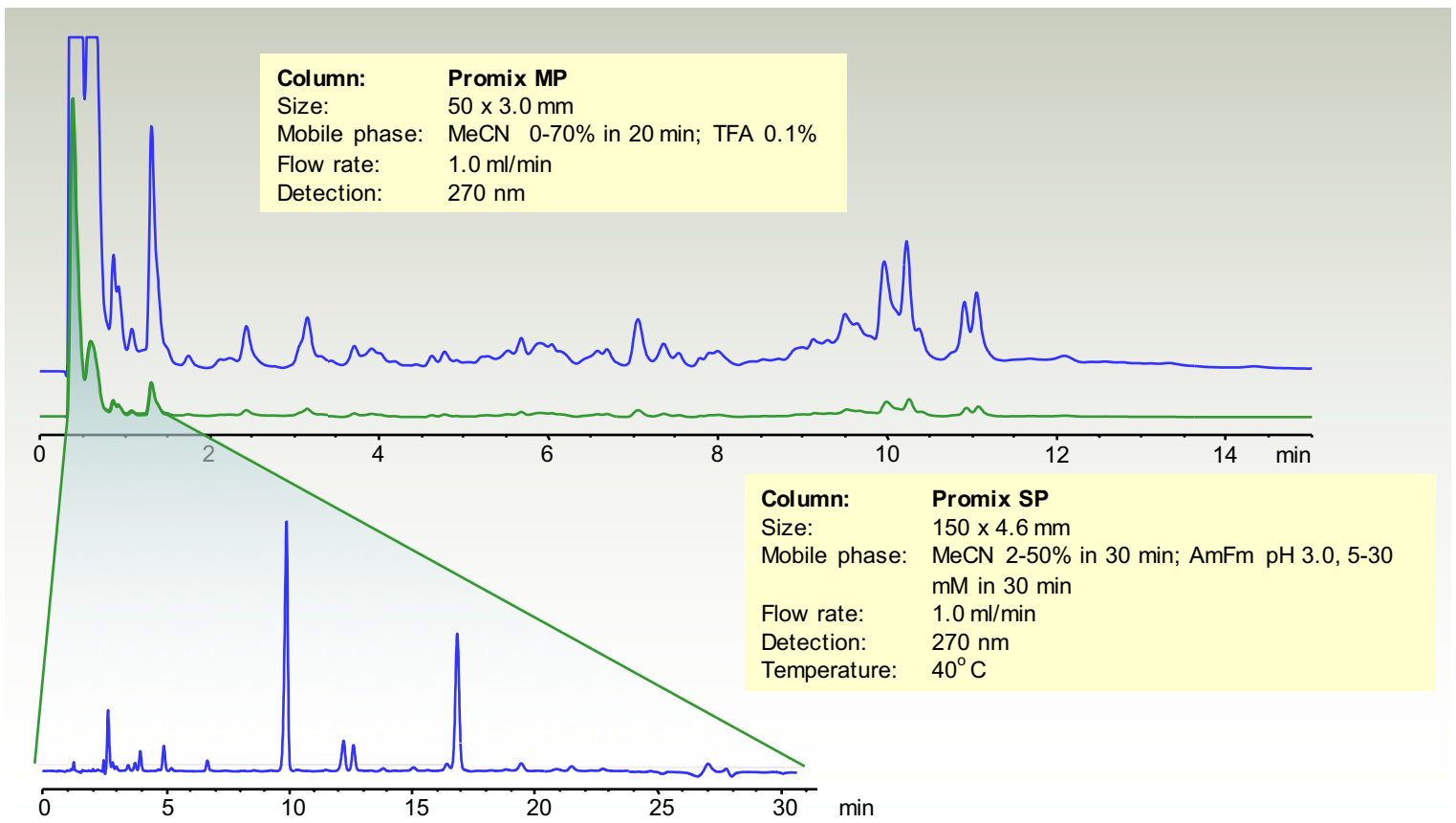
Synthetic insulin Humalog

Synthetic insulin Lantus

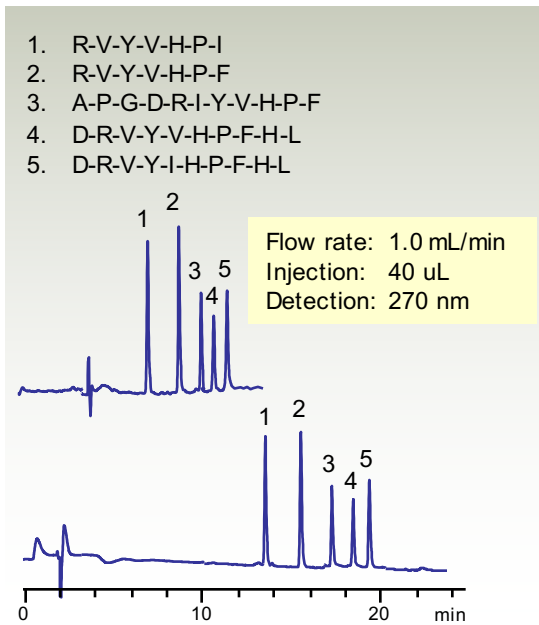
1. Normal human insulin
2. Humalog insulin
3. Lantus insulin

**Column:** Promix™ MP  
**Size:** 250 x 4.6 mm  
**Flow rate:** 1.0 ml/min  
**MeCN:** 20-40% in 30 min  
**Detector:** 270 nm  
**Injection:** 5 µL  
**Sample:** 30 units/mL in water





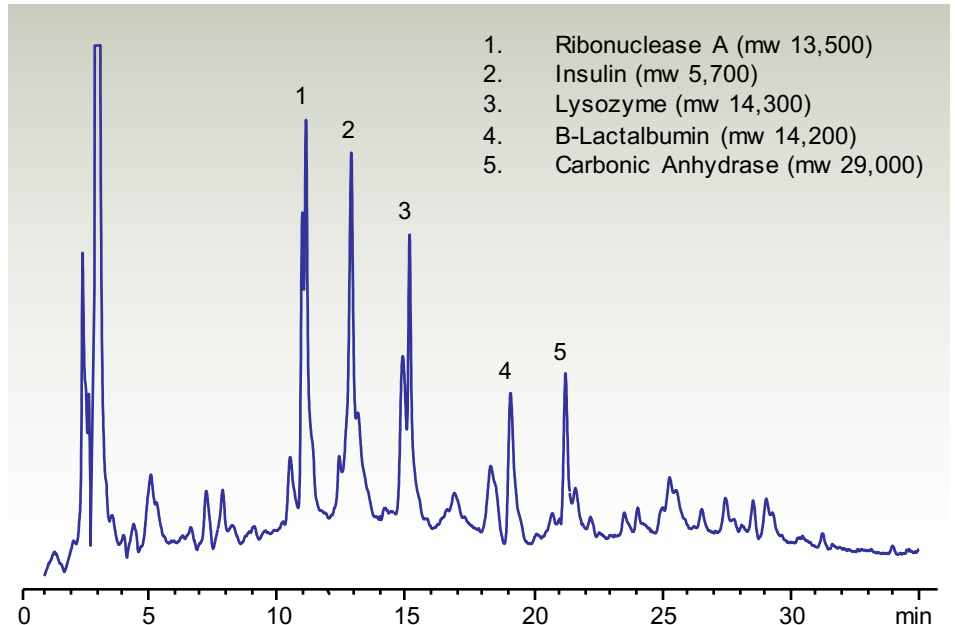
**Separation of a Complex Mixture** containing both large and small peptides can be achieved using two different Promix columns. Small peptides eluted in the first 1.5 min using Promix MP were collected and resolved using Promix SP.



**Separation of small oligopeptides.**

**Top:**  
**Column: Promix MP**, 4.6 x 250 mm  
 Mobile phase: MeCN 15-35% in 15 min;  
 TFA 0.4-0.05% in 15 min

**Bottom:**  
**Column: Promix MP**, 4.6 x 150 mm  
 Mobile phase: MeCN 0-40% in 30 min;  
 TFA 0.4-0.05% in 30 min



**Separation of large (>5 kDa) peptides.**

**Column: Promix MP**  
 Size: 4.6 x 150 mm  
 Mobile phase: MeCN 20-60 % in 30 min; TFA 0.25-0.05% in 30 min  
 Flow rate: 1 ml/min  
 Injection: 100 uL  
 Detection: 270 nm

Column size: 250 x 4.6 mm  
 Flow rate: 1.0 mL/min  
 Detection: 210 nm

**Column:** Promix AP

**Mobile phase:** Gradient of MeCN from 15% to 40% in 20 min with H<sub>2</sub>SO<sub>4</sub> 0.2%

**Column:** Promix MP

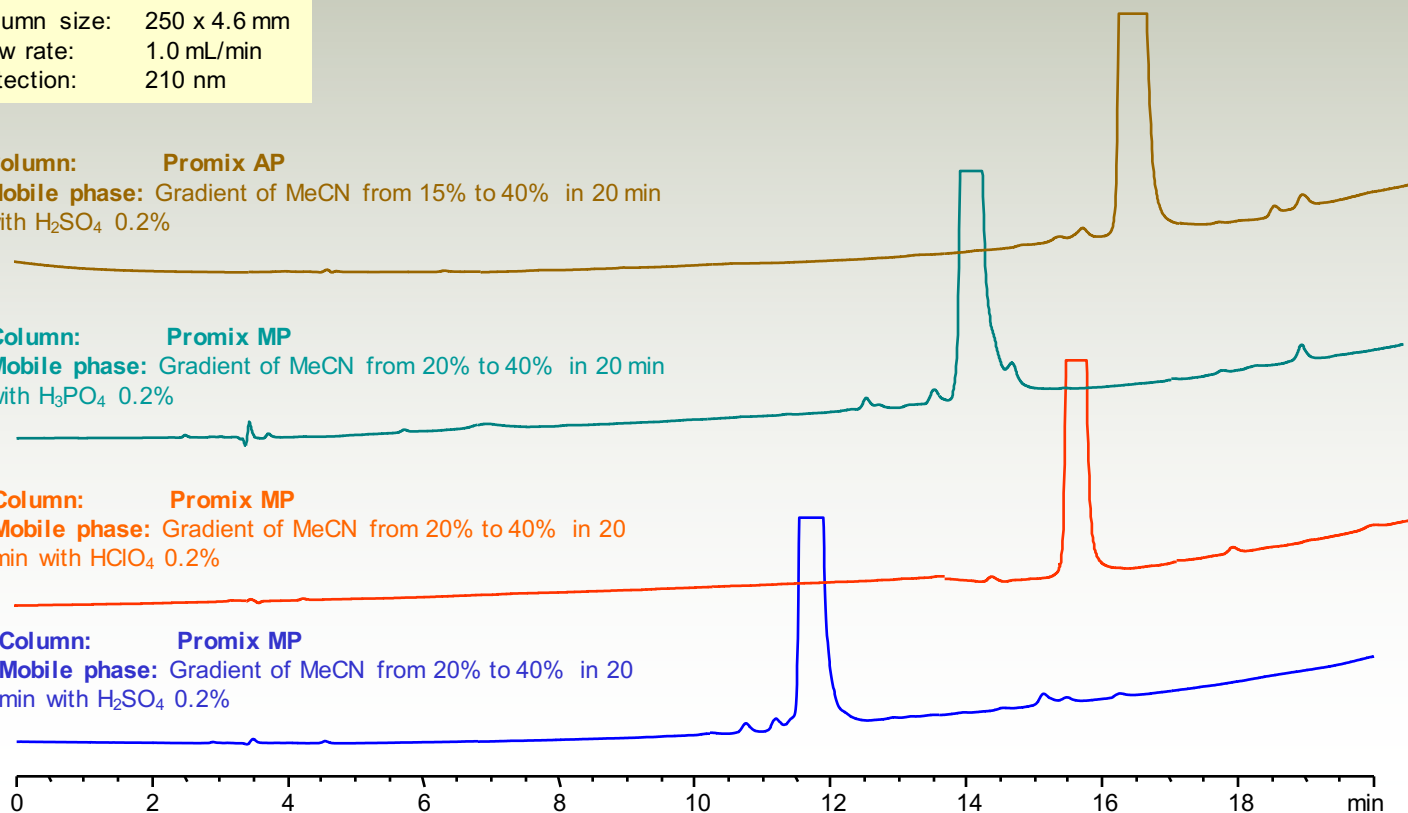
**Mobile phase:** Gradient of MeCN from 20% to 40% in 20 min with H<sub>3</sub>PO<sub>4</sub> 0.2%

**Column:** Promix MP

**Mobile phase:** Gradient of MeCN from 20% to 40% in 20 min with HClO<sub>4</sub> 0.2%

**Column:** Promix MP

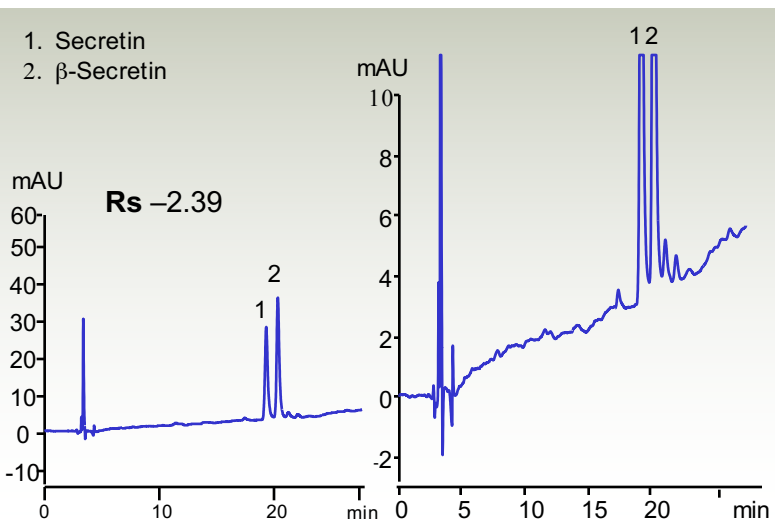
**Mobile phase:** Gradient of MeCN from 20% to 40% in 20 min with H<sub>2</sub>SO<sub>4</sub> 0.2%



**Selectivity adjustment by buffer type.** The choice of organic and ionic buffers, and their concentrations, significantly influences elution profile.

#### Separation of Secretin and $\beta$ -Secretin.

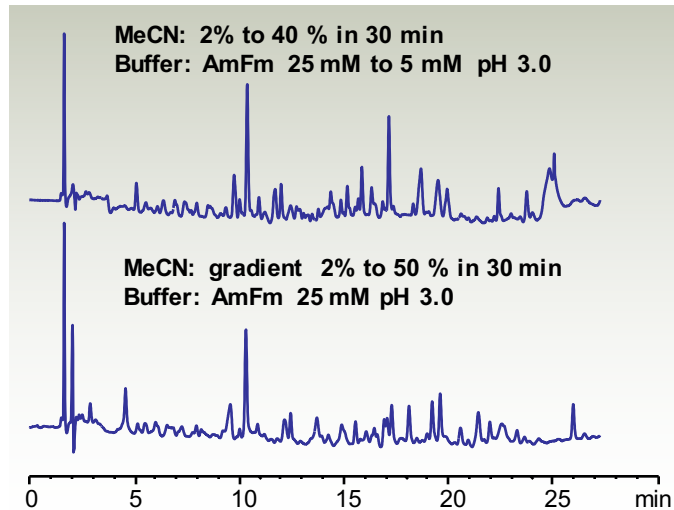
Secretin is a protein comprised of the following 11 amino acids:  
 H-S-D-G-T-S-E-L-S-R-L-R-D-S-A-R-L-Q-R-L-L-Q-G-L-V.



**Column:** Promix MP  
**Column size:** 250 x 4.6 mm  
**Mobile phase:** MeCN 10-25% in 30 min, H<sub>3</sub>PO<sub>4</sub> pH 2.0, 10 mM  
**Detection:** 210 nm  
**Flow:** 1.0 mL/min  
**Temperature:** 45°C

#### Effect of Buffer Concentration on Albumin Selectivity.

Complex mixtures such as protein digests exhibit different profiles at different ionic strengths. Multiple gradients can be applied to complex samples: organic gradients, ionic gradients, or both. Each provides a different selectivity and elution profile.



**Column:** Promix MP  
**Size:** 150 x 4.6 mm  
**Flow rate:** 1.0 mL/min  
**Detector:** 270 nm  
**Injection:** 50  $\mu$ L  
**Sample:** 0.3 mg/ml

Contact SIELC for information on column performance, method development, and availability.  
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