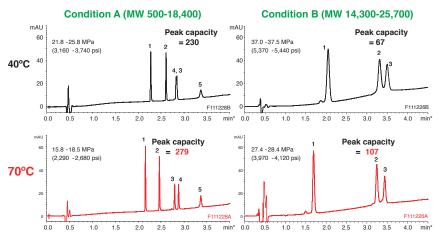
Effect of column temperature on separation of peptides and proteins



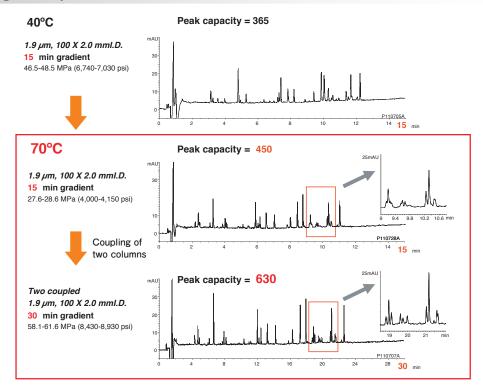
Analytes	MW	Peak width ½ (min)	
		40°C	70°C
Condition A			
1. Oxytocin	1,007	0.017	0.014
2. Leu-Enkephalin	556	0.015	0.015
β-Endorphin	3,465	-	0.016
4. Insulin	5,733	-	0.015
β-Lactoglobulin A	18,400	0.043	0.030
Condition B			
1. Lysozyme	14,300	0.069	0.044
2. α-Chymotrypsinogen	25,700	0.080	0.049
β-Lactoglobulin A	18,400	0.080	0.048

PC (peak capacity) = 1 + (gradient time/peak width*)
*peak width = 2Wo.sh average

The effect of temperature on separation of peptides and proteins with a variety of molecular weights (MW) is estimated. The separations at 40° C and 70° C are compared.

By increasing column temperature to 70°C, selectivity change is observed, and peaks become sharper, and improved resolution especially for larger molecules is obtained. Generally, larger molecules diffuse very slowly compared to small molecules. An elevated temperature can improve efficiency and peak shape by lowering mobile phase viscosity and improving mass transfer. Temperature is a simple and effective tool to increase resolution in separation of proteins and peptides.

Improvement of resolution by increasing column temperature and coupling of 1.9 μ m columns



Column : YMC-Triart C18 (1.9 µm, 12 nm)
Eluent : A) water/TFA (100/0.1)
B) acetonitrile/TFA (100/0.08)
5-40%8 (0-15 min) for a single column
5-40%8 (0-30 min) for two coupled columns
Flow rate : 0.4 mL/min
Detection : UV at 220 nm
Injection : 10 µL for a single column
20 µL for two coupled columns
Sample : Tryptic digest of Bovine Hemoglobin
System : Agilient 1290

23% more peaks can be resolved by increasing the column temperature to 70°C in the separation of tryptic digest of Hemoglobin.

The outstanding efficiency obtained by a coupling of two 100 mm length of Triart 1.9 μ m columns reduces co-elution of peaks and allows the precise separation in an analysis of complicated samples, such as peptide mapping.