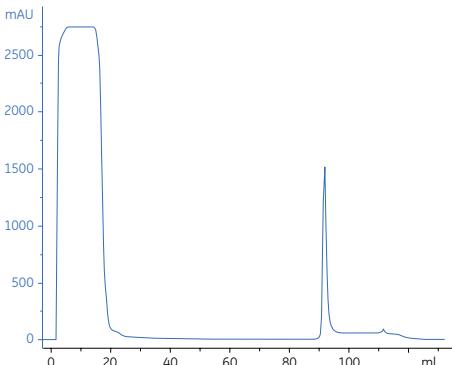


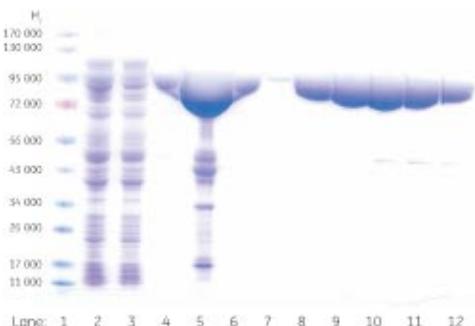
# 代謝疾患に関するMBP 融合タンパク質 (MCAD : medium-chain acyl-CoA dehydrogenase) の精製

Column: MBPTrap HP 5 ml  
Sample: N-terminal MBP-MCAD in E. coli lysate  
Sample volume: 15 ml  
Flow rate: 5.0 ml/min  
(0.5 ml/min during sample loading)  
Binding buffer: 20 mM Tris-HCl, 200 mM NaCl, 1 mM EDTA, 1 mM DTT, pH 7.4  
Elution buffer: 10 mM maltose in binding buffer  
System: ÄKTAprime plus

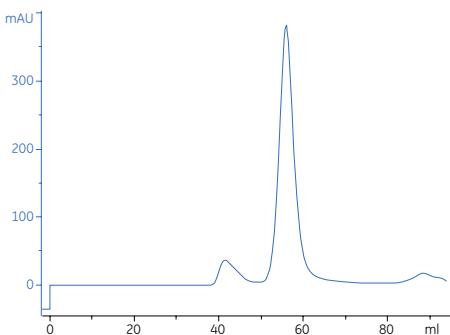


## Lane

- 1 Molecular weight marker
- 2 Start material, N-terminal MBP-MCAD in E. coli lysate, dil. 6 ×
- 3 Flow through MBPTrap HP, dil. 6 ×
- 4–6 Eluted fractions from MBPTrap HP
- 7–12 Eluted fractions from gel filtration



Column: HiLoad 16/60 Superdex 200pg  
Sample: Eluted fraction from MBPTrap HP 5 ml  
Sample volume: 2 ml  
Flow rate: 0.4 ml/min  
Buffer: 20 mM HEPES, 200 mM NaCl, pH 7.0  
System: ÄKTAprime plus



MBPTrap HP で精製した後、続いて HiLoad 16/60 Superdex 200 pg で最終精製を行いました。還元条件下の SDS-PAGE により、極めて純度高く目標のタンパク質を精製できていることがわかります。