

Putative signal peptide

Pru p 3.01 : MAYSAMT-KLA-LVVAL-CMVVSVPIA--QAITCGQVSSSLAPCI PYVRGGGAVPPACCNGIRNVNNLAR : 65
 Pru du 3.01 : MAYSAMT-KLA-LVVAL-CMVVSVPIA--QAITCGQVSSNLAPCI PYVRGGGAVPPACCNGIRNVNNLAR : 65
 Pru p 3.02 : MASSGQLLKLVLCLVAVMCCMAVGGPKA-MAAVSCGQVVNNLT PCIN YVANGGALNPSCCTGVRSLYSLAQ : 69
 Pru du 3.02 : MASSGQLLKLVLCLVAVMCCMAVGGPKA-MAAVSCGQVVNNLT PCIN YVANGGALNPSCCTGVRSLYSLAQ : 69
 Pru p 3.03 : MAS----LKVVCALF-M-CMVVAAPLITEAALTCPQIQAGLAPCLGYLQRGGVPAGGCCPGIKRLVGSAT : 64
 Pru du 3.03 : MAS----LKVVCALF-M-CMVVAAPLITEAALTCPQIQAGLAPCLGYLQRGGVPAGGCCPGIKRLVGSAT : 64

Pru p 3.01 : TTPDRQAACNCLKQLSASVPGVNPNN--AAALPGKCGVSI PYKISASTNCATVK : 117
 Pru du 3.01 : TTPDRQAACNCLKQLSASVPGVNPNN--AAALPGKCGVNI PYQISPSTNCANVK : 117
 Pru p 3.02 : TTADRQSI CNCLKQAVNGI PYTNANAGLAAGLPGKCGVNI PYKISPSTDCKSIK : 123
 Pru du 3.02 : TTADRQSI CNCLKQAVNGI PYTNANAGLAAGLPGKCGVNI PYKISPSTDCKSIK : 123
 Pru p 3.03 : TTADRQNACKCLKTVAGAVKGIN TGY--AAALPSLCGVKI PYKISASTNCNSVK : 116
 Pru du 3.03 : TTADRQNACKCLKTVAGAVKGINPGY--AAALPSLCGVKI PYKISASTNCNSVK : 116