

-- VIPDTPSPTVSVITISSDTDEEEEQKHAPTSTVSKQRKNVISCVTVHD 916
 SPYSDSSSNTSPYSVQQRAGHNNANAFDTKGSLENHCTGNPRTIIVPP
 LKTQASEVLVECSLV--
 ↑
 977

Putative caspase cleavage sites within the HIPK2 sequence mediating its degradation. The positions of aspartic acids 977 and 916 are indicated.