

EXPB1 20 LTARATWYgqpngagapdngGACgiknvnlpypsg
 GH45 1 ADGRSTRY wdck PSCgwakkapvnpvfscnanfgritdfdaksgcepvgvayscadqtpwaynddfal

I → II →

EXPB1 55 MTACGNVPIFKDGKGCGSCYEVRCKEKPECSGNPVTTVYITDMNYEPIAPYHFDLSgkafgslakpglndklr 126
 GH45 71 GFAATSIAGSNEAGWCCCACYELTFTSG-PVAGKKMVVQSTSTGGDLG-SNHFDLNipgggvgifdgctpqfg 140

II → III → IV → V → VI →

III → IV → V → VI →

α_a α_b α_c

EXPB1 127 HCGIMDVEFRRVRCKY 142
 GH45 141 glpggryggissrnecdrfpdalkpgcywrfdwfkNADNPSFSFRQVQCPAelvartgcrnrnddgnfpav 210

VI → VII →

α_a α_b α_c