

contig00022 MEGEVETFAFQAEIAQLMSLINTFSNKEIFIRELTISNSSDALDKIRYESLTDPTKLEAKELFIKLIPDKNEGTLT 80  
 contig10288 -  
 contig00026 -  
 contig02405 -  
 contig02406 -  
 contig00025 -

contig00022 DSGVGMTKADELINNLGTIAKSGTKAFMEAASAGADSMIGQFGVGFY SAYLVADKVVVTSKCNNDDEOYVWESSAGGSFTI 160  
 contig10288 -  
 contig00026 -  
 contig02405 -  
 contig02406 -  
 contig00025 -

contig00022 RTDNSEPIGRGTTKIVLHLKEDMTDYCEEKTVKEIVKKHQSFIGYPIKLLVEKERDLEISDEEEEDKEKMQIRKRKMRTN 240  
 contig10288 -  
 contig00026 -  
 contig02405 -  
 contig02406 -  
 contig00025 -

contig00022 QKELKMLVRMKMQIRMLIKTKRKKTVKKEKYTEDEEINKTKPLWTRNPDDVSTEELYGEFYKSITNDWEDHLAVKHFSVEGQL 320  
 contig10288 -  
 contig00026 -  
 contig02405 -  
 contig02406 -  
 contig00025 -

contig00022 EFRALLEFLPRRAAFDLFENKKQKNKIKLYVRRVEMENCEDLIPDYLNFLTGVVDSEDLPNTISREMLQQNKILKVKIRKN 400  
 contig10288 -  
 contig00026 -  
 contig02405 -  
 contig02406 -  
 contig00025 -

contig00022 LVKKAMELIEEICEDKENYTFKYYENFSKNNLKGTHEDSNRKKLADFLRYHTSSGDDEMTSIKDYVSRMKETQKQIYYIT 480  
 contig10288 DKEQYKKFYDNGKNVKLAHDSTNRKKFAEFLRYHTSHSGEEMCSKEYVSRMKENQSQIYIFT 66  
 contig00026 -  
 contig02405 -  
 contig02406 -  
 contig00025 -

contig00022 GESKDQVANSAFVERVKKRGFEVIYMTEPIDEYCYQQKEYDGKTLVSITKEGLELPEDEEEKKKFEEQKTKMENLCKVM 560  
 contig10288 GEGREQITNSSFVERVKKGGFEVIYMTPIDEYSITQKEYEGKQLI SVTKEGLELPEDDDD -  
 contig00026 -  
 contig02405 -  
 contig02406 -  
 contig00025 -

contig00022 KDIIDKRVEKVLVSNRLETSPCCIVTSQHWTANMERIMKAQALRDTSTMGYMAAKKHLEINPDHSII - 629  
 contig10288 -  
 contig00026 -  
 contig02405 -  
 contig02406 -  
 contig00025 -

contig00022 KDIIDKRVEKVVVSNRLVTSPCCIVTSQYWTANMERIMKAQALRDTSTMGYMAAKKHLEINPDHSIVE - 628  
 contig10288 -  
 contig00026 -  
 contig02405 -  
 contig02406 -  
 contig00025 -

contig00022 HSIVETLRQKADADKN 16  
 contig10288 -  
 contig00026 -  
 contig02405 -  
 contig02406 -  
 contig00025 RKKADADKN 9

contig00022 - 629  
 contig10288 - 128  
 contig00026 - 77  
 contig02405 DKSVKDILVMILLFETSSLSSGFSLLEDPGIHASRIVRMIRLGGLIDEEDLPVEDSNTAAEEMPPLECDDEDASRMEEVD\* 93  
 contig02406 DKSVKDILVMILLFETSSLSSGFSLLEDPGIHASRIVRMIRLGGLIDEEDLPVEDSNTAAEEMPPLECDDEDASRMEEVD\* 93  
 contig00025 DKSVKDILVMILLFETSSLTSGFSLEDPGVHSSRIVRMIKLGGLIDEEEEAPVKDSNVEEEMPSLECDDEDASR - 80