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gi|17647529|ref|NP_523899.1| FC SLADYVSRMKD NOKHVYFITGESKDVSN S AFVERVKRGFEVVMTEP IDEYVI QHLKEYKGLVSVTK EGLELPD -EAEKKKREBEDKAKFESLCKLMKSLDNKVEKVVVSNRLVDS PCCIVTSQFGWSANMERIMKAQALRDT 607
gi|118778689|ref|XP_308800.3| YCSLNDYVGRMKENQTIYFIITGESIDVKN S AFVERVKRGRFVIMTEP IDEYVIQOLKEKYGKQLVSVTK EGLELPD -EAEKKKREBEDKAKFENLCKVMKSVLESKVEKVMVSNRLVDS PCCIVTSQFGWSANMERIMKAQALRDS 602
gi|73964025|ref|XP_537557.2| MVSLKDYCTR MKENOKHIYIITGETKDQVAN S AFVERLRKHGLEVIYMI EPID EYCVQOLKEFEFGKTLVSVTK EGLELPD -EBEKKKQEBEKKTKFENLCKIMKDILEKKVEKVVVSNRLVTS PCCIVTSTYGWITANMERIMKAQALRDN 505
gi|60592792|ref|NP_001012688.1| MVSLKDYCTR MKENOKHIYIITGETKDQVAN S AFVERLRKHGLEVIYMI EPID EYCVQOLKEFEFGKTLVSVTK EGLELPD -EBEKKKQEBEKKTKFENLCKIMKDILEKKVEKVVVSNRLVTS PCCIVTSTYGWITANMERIMKAQALRDN 623
gi|6754254|ref|NP_034610.1| MVSLKDYCTR MKENOKHIYIITGETKDQVAN S AFVERLRKHGLEVIYMI EPID EYCVQOLKEFEFGKTLVSVTK EGLELPD -EBEKKKQEBEKKTKFENLCKIMKDILEKKVEKVVVSNRLVTS PCCIVTSTYGWITANMERIMKAQALRDN 623
gi|28467005|ref|NP_786937.1| MVSLKDYCTR MKENOKHIYIITGETKDQVAN S AFVERLRKHGLEVIYMI EPID EYCVQOLKEFEFGKTLVSVTK EGLELPD -EBEKKKQEBEKKTKFENLCKIMKDILEKKVEKVVVSNRLVTS PCCIVTSTYGWITANMERIMKAQALRDN 623
gi|114654793|ref|XP_001160832.1| MVSLKDYCTR MKENOKHIYIITGETKDQVAN S AFVERLRKHGLEVIYMI EPID EYCVQOLKEFEFGKTLVSVTK EGLELPD -EBEKKKQEBEKKTKFENLCKIMKDILEKKVEKVVVSNRLVTS PCCIVTSTYGWITANMERIMKAQALRDN 745
gi|153792590|ref|NP_001017963.2| MVSLKDYCTR MKENOKHIYIITGETKDQVAN S AFVERLRKHGLEVIYMI EPID EYCVQOLKEFEFGKTLVSVTK EGLELPD -EBEKKKQEBEKKTKFENLCKIMKDILEKKVEKVVVSNRLVTS PCCIVTSTYGWITANMERIMKAQALRDN 744
gi|113681112|ref|NP_001038538.1| MVSLKDYVTR MKDTOKHIYIITGETKDQVAN S AFVERLRKAGLEVIYMI EPID EYCVQOLKEFEFGKTLVSVTK EGLELPD -EBEKKKQEBEKKSKFENLCKIMKDILEKKVEKVT VSNRLVSS PCCIVTSTYGWITANMERIMKAQALRDN 624
gi|15237214|ref|NP_200076.1| MTSFKDYVTR MKEGQKDFIYIITGESKAVENS PFLERLRKRGYEVLYMVDALDEYAVGQLKEYD GKGLVSA K EGLELPD EEBEKKKREBEKKKSFENLCKITKEILGDKVEKVVVSDRIVDSPCCIVTGEYGWITANMERIMKAQALRDS 600
          .....610.....620.....630.....640.....650.....660.....670.....680.....690.....700.....710.....720.....730.....740.....750

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gi|17647529|ref|NP_523899.1| AITMGYMAKKHLEINPDHPIVE TLRQKAEADKNDKSVKDLVILLYETALLSSGFSLEDPTOTANRIYRMIKLGLGIDEEDPTADDSSAA-----VTEEMPPLEGD-DDTSRMEVD 717
gi|118778689|ref|XP_308800.3| SAMGYMAKKHLEINPDHPIVE TLRQKAEADKNDKSVKDLVILLYETALLSSGFSLEDPTOTANRIYRMIKLGLGIDEEDPTADDSSAA-----VTEEMPPLEGD-DDTSRMEVD 720
gi|73964025|ref|XP_537557.2| STMGYMAAKHLEINPDHSIIE TLRQKAEADKNDKSVKDLVILLYETALLSSGFSLEDPTOTANRIYRMIKLGLGIDEEDPTADDSSAA-----VTEEMPPLEGD-DDTSRMEVD 615
gi|60592792|ref|NP_001012688.1| STMGYMAAKHLEINPDHSIIE TLRQKAEADKNDKSVKDLVILLYETALLSSGFSLEDPTOTANRIYRMIKLGLGIDEEDPTADDSSAA-----VTEEMPPLEGD-DDTSRMEVD 733
gi|6754254|ref|NP_034610.1| STMGYMAAKHLEINPDHSIIE TLRQKAEADKNDKSVKDLVILLYETALLSSGFSLEDPTOTANRIYRMIKLGLGIDEEDPTADDSSAA-----VTEEMPPLEGD-DDTSRMEVD 733
gi|28467005|ref|NP_786937.1| STMGYMAAKHLEINPDHSIIE TLRQKAEADKNDKSVKDLVILLYETALLSSGFSLEDPTOTANRIYRMIKLGLGIDEEDPTADDSSAA-----VTEEMPPLEGD-DDTSRMEVD 733
gi|114654793|ref|XP_001160832.1| STMGYMAAKHLEINPDHSIIE TLRQKAEADKNDKSVKDLVILLYETALLSSGFSLEDPTOTANRIYRMIKLGLGIDEEDPTADDSSAA-----VTEEMPPLEGD-DDTSRMEVD 855
gi|153792590|ref|NP_001017963.2| STMGYMAAKHLEINPDHSIIE TLRQKAEADKNDKSVKDLVILLYETALLSSGFSLEDPTOTANRIYRMIKLGLGIDEEDPTADDSSAA-----VTEEMPPLEGD-DDTSRMEVD 854
gi|113681112|ref|NP_001038538.1| STMGYMAAKHLEINPDHPIVE TLRQKAEADKNDKSVKDLVILLYETALLSSGFSLEDPTOTANRIYRMIKLGLGIDEEDPTADDSSAA-----VTEEMPPLEGD-DDTSRMEVD 734
gi|15237214|ref|NP_200076.1| MIMGYMAAKHLEINPDHPIVE TLRQKAEADKNDKSVKDLVILLYETALLSSGFSLEDPTOTANRIYRMIKLGLGIDEEDPTADDSSAA-----VTEEMPPLEGD-DDTSRMEVD 705
          .....760.....770.....780.....790.....800.....810.....820.....830.....840.....850.....860.....

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