

Logo



NTDB id 626 LCA RS04960 WP 011374694.1	ITANLALQANRNVYAIPGRIDQSLSQGCNQLIAAGATPILLDKNILIEELRYFD.....	288
NTDB id 129639 LG36 RS06185 WP 038600928.1	ITCERALEEGRDIFAVPGNIADGFSQGCNHLIQGAKLIYQAQDILEEYLYD.....	282
NTDB id 599 KW2 RS05940 WP 021037268.1	ITCERALEEGRDIFAVPGNIADGTSQGCNHLIQGAKLVYQAQDILEEYLYN.....	282
NTDB id 407 SMU RS04605 WP 002262865.1	ITCERAMEEGRDVFVPGNILDGQSEGGCHHLIQEGAKCITSGFDILNEFNF.....	280
NTDB id 525 SMSK321 RS06295 WP 000705298.1	ITCERAMEEGRDVFVPGNILDGQSEGGCHHLIQEGAKCITSGFDILNEFNF.....	282
NTDB id 267 KZH43 RS05590 WP 000705306.1	ITCERAMEEGRDVFVPGNILDGQSEGGCHHLIQEGAKCITSGFDILNEFNF.....	282
NTDB id 226 SPD RS05990 WP 000705306.1	ITCERAMEEGRDVFVPGNILDGQSEGGCHHLIQEGAKCITSGFDILNEFNF.....	282
NTDB id 192 SPR RS05715 WP 000705306.1	ITCERAMEEGRDVFVPGNILDGQSEGGCHHLIQEGAKCITSGFDILNEFNF.....	282
NTDB id 157 SP RS06205 WP 000705306.1	ITCERAMEEGRDVFVPGNILDGQSEGGCHHLIQEGAKCITSGFDILNEFNF.....	282
NTDB id 497 SM12261 RS05565 WP 000705318.1	ITCERAMEEGRDVFVPGNILDGQSEGGCHHLIQEGAKCITSGFDILNEFNF.....	282
NTDB id 1361 HI 0985 AAC22646.1	ITARYALEQNRVFAVPGNIQNKSSQGCNHLIKQGAMLVENAKDILETLYQHSI.HSQTEIDFDQIAVPNYTPPPDPRRLVEAPS	321
NTDB id 1021 ACIAD RS00975 WP 004930462.1	ITANKAAEQGKVVFAIPGHIYSEFHQGCNHLIREGAILVDHPEQITIEDLALPTHWHSQSTAS.GEANADD.SGKTDDA..DLPDD	327
consensus	!!***!*****!! !*****!***!***!***** ** *	

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NTDB id 1361 HI 0985 AAC22646.1	HPKLYSRIGYTPVSIDDLAEFNLSVDVLLVQLLDLELQDLIISENGLYKRV....	373
NTDB id 1021 ACIAD RS00975 WP 004930462.1	LLPLYQLLDWQQDLDQLAQHYQGSTAELTAQLMELELLGVCIQSGRYLRCRSPY	383
consensus		

- X non conserved
- X similar
- X ≥ 50% conserved