

NTDB	id	1250	GC085	RS13065	WP	027219989.1	MNNLQPLLALNRMKRIQPRT.....VLKLQKRWPDNLNMFQLSSVDLEEAGLPSWLAQTIKFNPDFI	63
NTDB	id	114	BSU	16110	NP	389493.1	MDQAAVCLTICRINQLSPSLILLKWWKADPSMSLTSPVLQTVT.....RDQIKA...AALK...NEIEQFYPKLPRVLA	68
NTDB	id	599	KW2	RS05940	WP	021037268.1	MITNFDLFRWKKAGMTNLGVNKLKFFRKYDRKISLRQMGQV...AQVKSIPNFIEQY.KNQDVKKL	63
NTDB	id	525	SMSK321	RS06295	WP	000705298.1	MKITNYEIIYKLKKSGLTNQQILKVLEYGENVDQELLGDIADI.SGCRNPAVFMERY.FQIDDAHL	64
NTDB	id	267	KZH43	RS05590	WP	000705306.1	MKITNYEIIYKLKKSGLTNQQILKVLEYGENVDQELLGDIADI.SGCRNPAVFMERY.FQIDDAHL	64
NTDB	id	226	SPD	RS05990	WP	000705306.1	MKITNYEIIYKLKKSGLTNQQILKVLEYGENVDQELLGDIADI.SGCRNPAVFMERY.FQIDDAHL	64
NTDB	id	192	SPR	RS05715	WP	000705306.1	MKITNYEIIYKLKKSGLTNQQILKVLEYGENVDQELLGDIADI.SGCRNPAVFMERY.FQIDDAHL	64
NTDB	id	157	SP	RS06205	WP	000705306.1	MKITNYEIIYKLKKSGLTNQQILKVLEYGENVDQELLGDIADI.SGCRNPAVFMERY.FQIDDAHL	64
NTDB	id	497	SM12261	RS05565	WP	000705318.1	MKITNYEIIYKLKKSGLTNQQVLAVLEYGENVDQELLGDIAEI.SGCRNPAVFMERY.FQIDDAHL	64
NTDB	id	40548	SSUST3	RS05080	WP	009909860.1	MNNFELFKWKKAGLTNLNVNKKVLAYQEKHGKKLTLRNVAVV.SECKNPVLFMENY.KNLDTKAC	62
NTDB	id	407	SMU	RS04605	WP	002262865.1	MDNFQLFKLKKAGLTNLNILNIIDYEERTQKSLSLRDMAVV.SKNKKPLIFMEHY.KNLDISKAL	62
consensus							*****	62

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