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NTDB id 1245 Cj1673c YP 002345041.1	R	L	D	V	R	K	V	A	T	L	.	K	Q	N	E	P	I	G	N	R	V	K	V	V	V	K	N	K	V	A	P	P	F	R	Q	A	E	F	D	V	M	F	G	E	G	I	S	R	E	G	E	L	I	D	Y	C	V	K	L	D	I	V	D	K	S	G	A	W	F	S	Y	K	D	K	296					
NTDB id 1214 NCTC11637 00359 SQJ03206.1	R	I	D	I	R	R	I	A	A	L	.	K	Q	N	E	Q	H	I	G	N	R	A	K	A	V	V	V	K	N	K	V	A	P	P	F	R	E	A	E	F	D	I	M	F	G	E	G	I	S	K	E	G	E	I	D	Y	C	V	K	L	D	I	V	D	K	S	G	A	W	L	S	Y	Q	D	K	298				
NTDB id 1211 C694 RS00795 WP 000963128.1	R	I	D	I	R	R	I	A	S	L	.	K	Q	N	E	Q	H	I	G	N	R	A	K	A	V	V	V	K	N	K	V	A	P	P	F	R	E	A	E	F	D	I	M	F	G	E	G	I	S	K	E	G	E	I	D	Y	C	V	K	L	D	I	V	D	K	S	G	A	W	L	S	Y	Q	D	K	298				
NTDB id 1334 RA0C RS04870 WP 004918236.1	R	I	D	I	R	R	S	L	S	D	T	G	K	I	K	D	K	S	G	E	V	I	G	N	R	T	K	V	V	V	K	N	K	V	A	P	P	F	K	S	A	E	F	D	I	M	Y	G	E	C	V	S	K	V	G	E	I	D	V	A	T	D	L	D	I	I	Q	K	S	G	W	F	S	Y	Q	D	T	309						
NTDB id 1389 A4U84 RS07560 WP 021112712.1	R	L	D	I	R	R	C	G	V	V	.	K	D	G	E	I	I	G	S	E	T	K	V	V	V	K	N	K	V	A	P	P	F	R	E	V	Q	F	D	I	M	Y	G	E	G	I	S	R	M	N	E	L	L	I	L	A	E	A	N	G	F	I	K	S	G	A	W	F	S	Y	D	G	T	312						
NTDB id 1130 NGFG RS03960 WP 003688695.1	R	L	D	I	R	R	T	G	S	I	.	K	K	G	E	E	V	L	G	N	E	T	R	V	K	V	I	K	N	K	V	A	P	P	F	R	Q	A	E	F	D	I	L	Y	G	E	G	I	S	W	E	G	E	L	I	D	I	G	V	K	N	D	I	I	N	K	S	G	A	W	Y	S	I	N	G	A	296			
NTDB id 1124 NGFG RS03960 WP 003688695.1	R	L	D	I	R	R	T	G	S	I	.	K	K	G	E	E	V	L	G	N	E	T	R	V	K	V	I	K	N	K	V	A	P	P	F	R	Q	A	E	F	D	I	L	Y	G	E	G	I	S	W	E	G	E	L	I	D	I	G	V	K	N	D	I	I	N	K	S	G	A	W	Y	S	I	N	G	A	296			
NTDB id 1123 OK783 RS03845 WP 003688695.1	R	L	D	I	R	R	T	G	S	I	.	K	K	G	E	E	V	L	G	N	E	T	R	V	K	V	I	K	N	K	V	A	P	P	F	R	Q	A	E	F	D	I	L	Y	G	E	G	I	S	W	E	G	E	L	I	D	I	G	V	K	N	D	I	I	N	K	S	G	A	W	Y	S	I	N	G	A	296			
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	R	L	D	I	R	R	T	G	A	I	.	K	E	G	E	E	V	L	G	N	E	T	R	I	K	V	V	K	N	K	I	A	A	P	F	K	E	A	N	T	Q	I	M	Y	G	Q	G	F	N	R	E	G	E	L	I	D	L	G	V	K	H	K	M	V	E	K	S	G	A	W	Y	S	I	N	G	D	295			
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NTDB id 1205 PSJM300 05545 AFN77183.1	R	L	D	I	R	R	T	G	A	V	.	K	E	G	E	E	V	L	G	S	E	T	R	V	K	V	V	K	N	K	V	A	P	P	F	R	Q	A	E	F	D	I	L	Y	G	A	C	V	S	R	E	G	E	I	D	L	G	V	E	A	K	V	V	E	K	S	G	A	W	Y	S	I	N	G	E	295				
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NTDB id 600 KW2 RS01765 WP 011675413.1	R	L	D	V	R	G	S	T	K	I	E	E	G	S	G	D	N	K	T	Q	I	G	K	I	T	K	V	V	K	N	K	V	A	P	P	F	K	V	A	L	V	D	I	M	F	G	E	G	I	S	S	T	G	E	L	I	N	I	A	V	E	E	G	I	I	K	K	S	G	A	W	F	A	Y	N	D	E	314		
NTDB id 1142081 DQL01 RS10030 WP 061459621.1	R	L	D	V	R	G	N	T	Q	I	K	G	T	G	D	Q	K	E	T	N	V	G	K	E	T	K	I	K	V	V	K	N	K	V	A	P	P	F	K	E	A	V	V	E	I	M	Y	G	E	G	I	S	K	T	G	E	L	L	K	I	A	S	D	L	D	I	I	K	K	A	G	A	W	Y	S	I	K	D	E	313
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consensus
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