

SUPPORTING INFORMATION

Capnophilic lactic fermentation from *Thermotoga neapolitana*: a resourceful pathway to obtain almost enantiopure L-lactic acid.

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Figure S1. GC-MS analysis of O-trifluoracetyl-(-)-menthyl ester derivatives of DL-lactic acid, L-lactic acid and natural lactate of *T. neapolitana*, co-elution of DL-Lactic acid and natural lactic acid. (Rt 10.21 min = L-lactic acid derivative; Rt 10.25 min = D-lactic acid derivative).

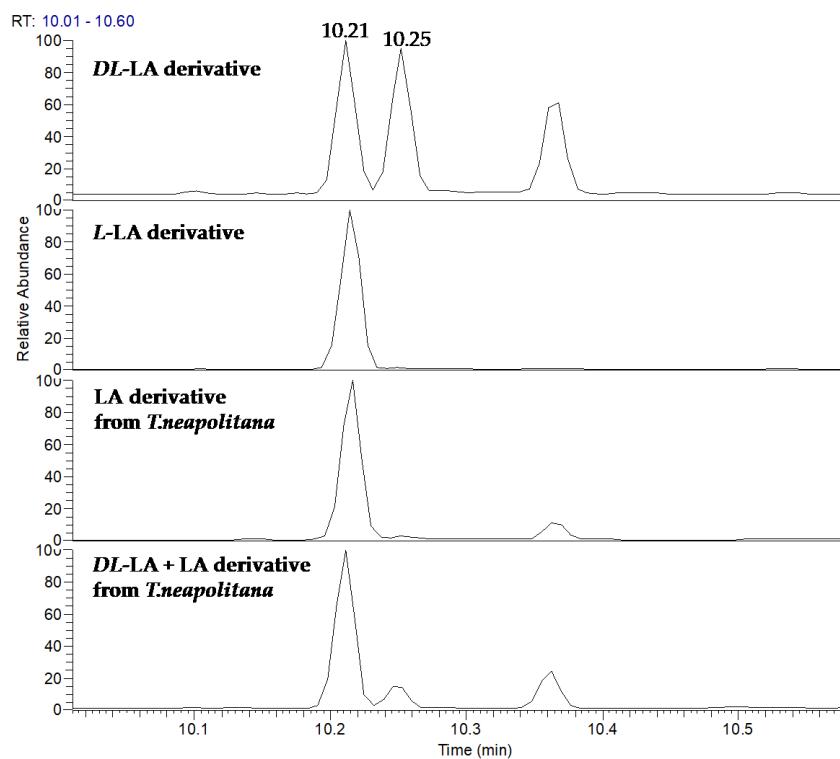


Figure S2. GC-MS chromatogram of O-trifluoracetyl-(-)-menthyl ester of lactic acid from *T. neapolitana* and integration of peaks area of *L*- and *D*-LA.

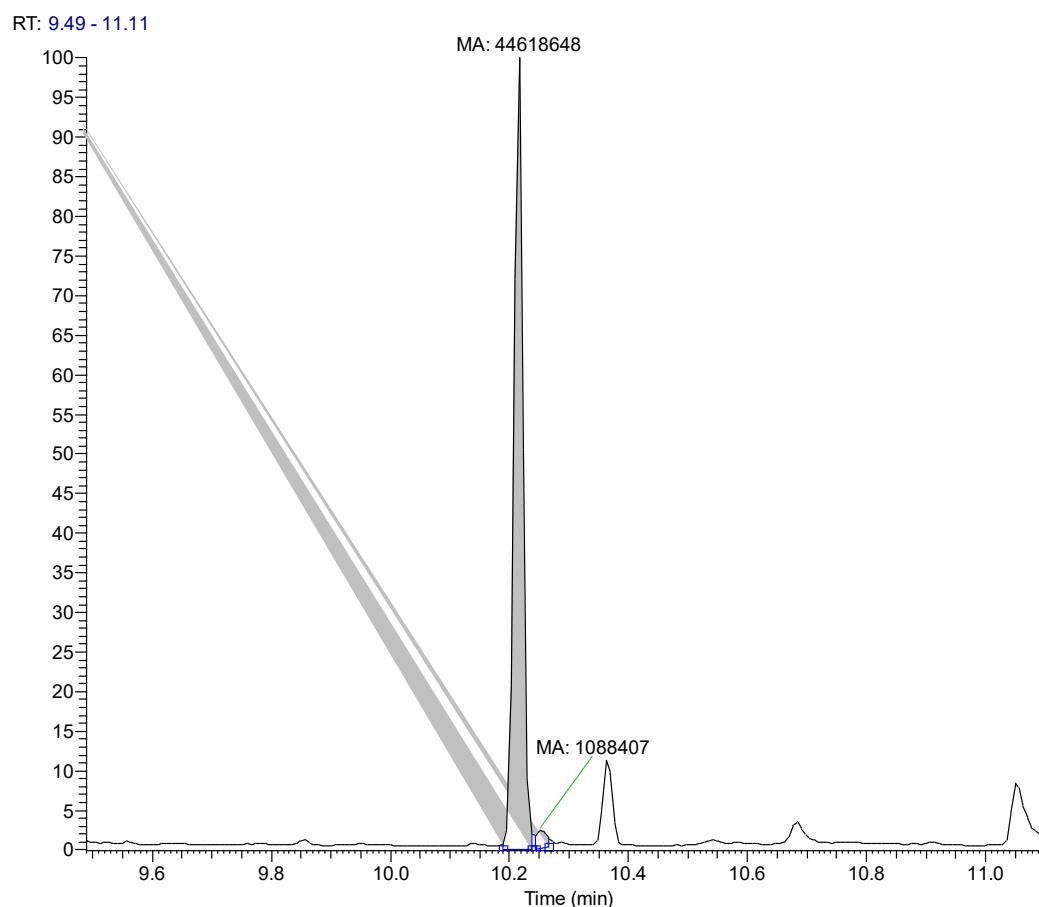


Figure S3. Sequence alignment of LDH from members of the bacterial order Thermotogales

	1	10	20	30	40	50	60																								
PthLDH	-----	MKVSIFGAGRVGISIAYSSLH	T	I	A	E	M																								
TmLDH	-----	M	KIGIVGLGRVGSSTA	F	A	L	M																								
TnLDH	M	PSPCLYSITTEVIS	M	KIGIVGLGRVGSSTA	F	A	L																								
TnpLDH	-----	M	KIGIVGLGRVGSSTA	F	A	L	M																								
TRQ7LDH	-----	M	KIGIVGLGRVGSSTA	F	A	L	M																								
PthLDH	YHSTPFLKR	CNITAGNPHDILNSDFVI	I	TAGASQSAGESRLS	L	T	KRNVKII																								
TmLDH	IHGT	HGTPTRRANIYAGDYADL	KGS	DVIVIAAGVPQ	KPGETR	L	QLLGRNARVMKEIARNVSK																								
TnLDH	IHGT	HGTPTRRNIYAGDYADL	KGS	DVIVIAAGVPQ	KPGETR	L	QLLGRNARVMREIARNVSK																								
TnpLDH	IHGT	HGTPTRRNIYAGDYADL	KGS	DVIVIAAGVPQ	KPGETR	L	QLLGRNARVMREIARNVSK																								
TRQ7LDH	IHGT	HGTPTRRNIYAGDYADL	KGS	DVIVIAAGVPQ	KPGETR	L	QLLGRNARVMREIARNVSK																								
PthLDH	YSPDAIVINVSNPVDVL	SYVLUW	KETKF	NWRKV	I	G	T	G	T	I	D	T	A	R	F	R	A	L	V	A	K	Q	C	G	V	S	P	M	S	V	
TmLDH	YAPDSIVIVV	TNPVDVL	TYFFL	KESGM	DPR	KVF	FGSGT	VLD	TAR	L	R	L	I	A	Q	H	C	G	F	S	P	R	S	V							
TnLDH	YAPDSIVIVV	TNPVDVL	TYFFL	KESGM	DPR	KVF	FGSGT	VLD	TAR	L	R	L	I	A	Q	H	C	G	F	S	P	R	S	V							
TnpLDH	YAPDSIVIVV	TNPVDVL	TYFFL	KESGM	DPR	KVF	FGSGT	VLD	TAR	L	R	L	I	A	Q	H	C	G	F	S	P	R	S	V							
TRQ7LDH	YAPDSIVIVV	TNPVDVL	TYFFL	KESGM	DPR	KVF	FGSGT	VLD	TAR	L	R	L	I	A	Q	H	C	G	F	S	P	R	S	V							
PthLDH	AYIIGEHGDSELLVWSNATIGGVSI	KR	F	QC	F	C	T	N	R	C	T	F	L	F	E	Q	T	K	N	A	Y	E	I	I	E	R	K				
TmLDH	VVIGE	HGDSE	V	PVWSGAMIGGI	PL	QNM	C	Q	I	C	Q	K	C	D	S	K	I	L	E	F	A	E	K	T	K						
TnLDH	VVIGE	HGDSE	V	PVWSGAMIGGI	PL	QNM	C	Q	I	C	Q	K	C	D	S	H	I	L	E	F	A	E	K	T	K						
TnpLDH	VVIGE	HGDSE	V	PVWSGAMIGGI	PL	QNM	C	Q	I	C	Q	K	C	D	S	H	I	L	E	F	A	E	K	T	K						
TRQ7LDH	VVIGE	HGDSE	V	PVWSGAMIGGI	PL	QNM	C	Q	I	C	Q	K	C	D	S	H	I	L	E	F	A	E	K	T	K						
PthLDH	TNL	AI	G	T	A	A	L	V	E	I	Y	R	D	E	K	R	V	W	T	V	S	V	F	Q	D						
TmLDH	THY	A	I	A	L	A	V	D	I	V	E	S	I	F	D	E	K	R	V	L	T	S	V	L	E	D					
TnLDH	THY	A	I	A	L	A	V	D	I	V	E	S	I	F	D	E	K	R	V	L	T	S	V	L	E	D					
TnpLDH	THY	A	I	A	L	A	V	D	I	V	E	S	I	F	D	E	K	R	V	L	T	S	V	L	E	D					
TRQ7LDH	THY	A	I	A	L	A	V	D	I	V	E	S	I	F	D	E	K	R	V	L	T	S	V	L	E	D					
PthLDH	NSVEKEAFERSKEV	IKYIKE	GEK	SER	ESS	NN-																									
TmLDH	NEEELEAFRK	SASIL	KN	NA	I	E	E	N	K	H	Q	N	T	S	G																
TnLDH	SEEEMKA	FRESA	KIL	K	S	A	I	E	E	I	L	A	E	E	N	K	H	Q	N	T	S	G									
TnpLDH	NEEELEAFRK	SASIL	KN	NA	I	E	E	N	K	H	Q	N	T	S	G																
TRQ7LDH	SEEEMKA	FRESA	KIL	K	S	A	I	E	E	I	L	A	E	E	N	K	H	Q	N	T	S	G									