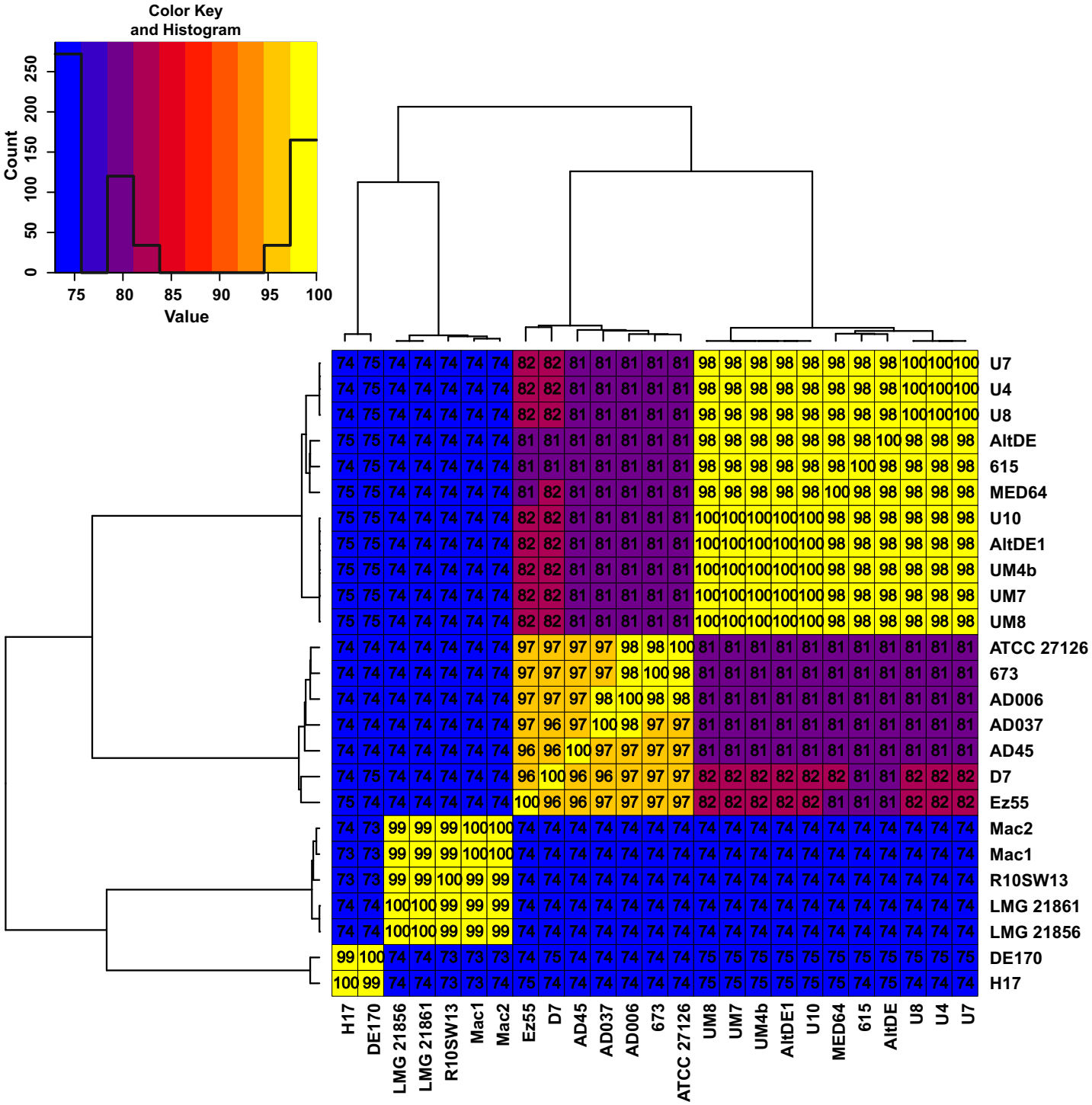
Pangenome evolution in the marine bacterium *Alteromonas*

Mario López-Pérez[1] and Francisco Rodriguez-Valera[1*]

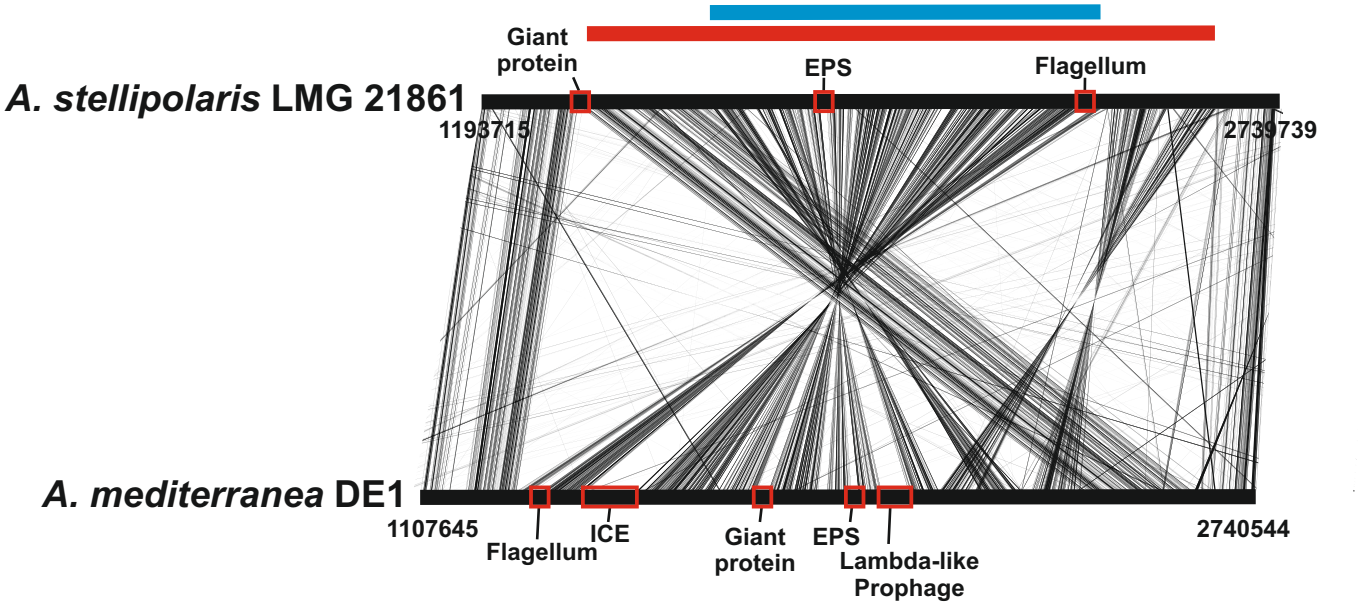
[1]Evolutionary Genomics Group, Departamento de Producción Vegetal y Microbiología, Universidad Miguel Hernández, San Juan de Alicante, 03550, Alicante, Spain

*Corresponding author: frvalera@umh.es
Evolutionary Genomics Group, División de Microbiología, Universidad Miguel Hernández, Apartado 18, San Juan 03550, Alicante, Spain
Phone +34-965919313, Fax +34-965 919457

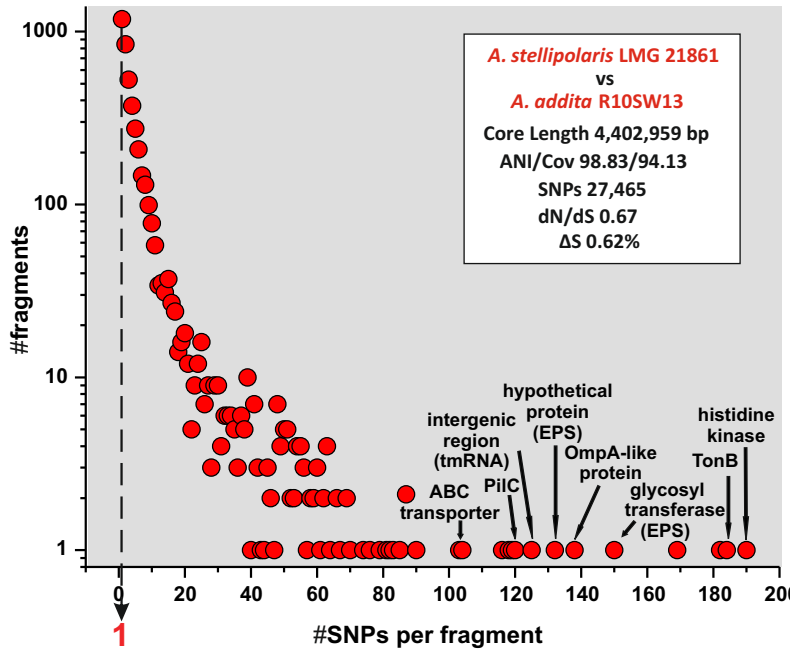
Supplementary Figure S1

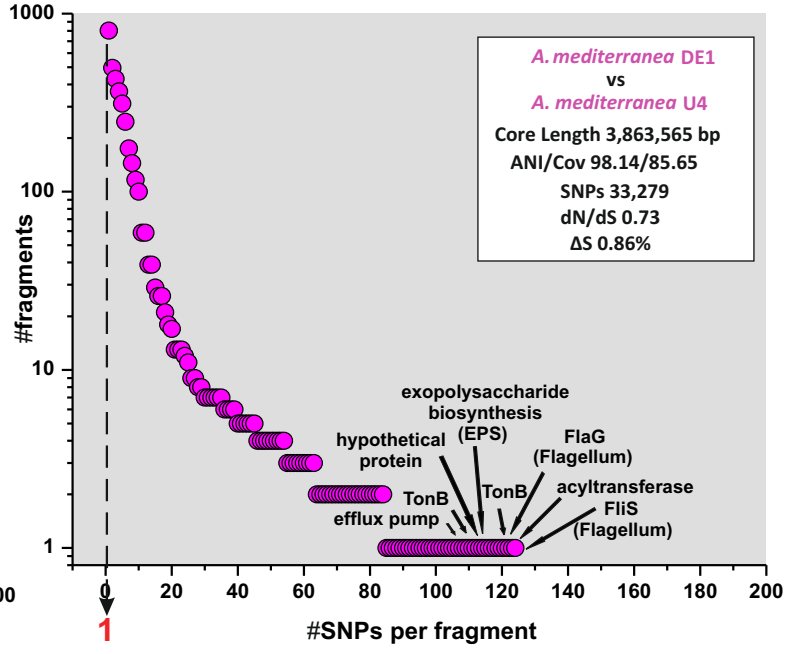


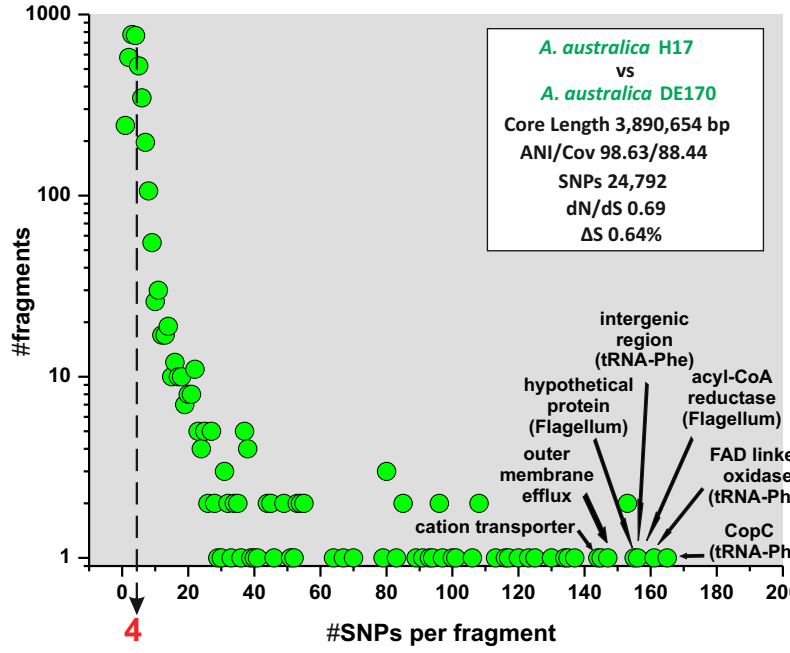
Supplementary Figure S2

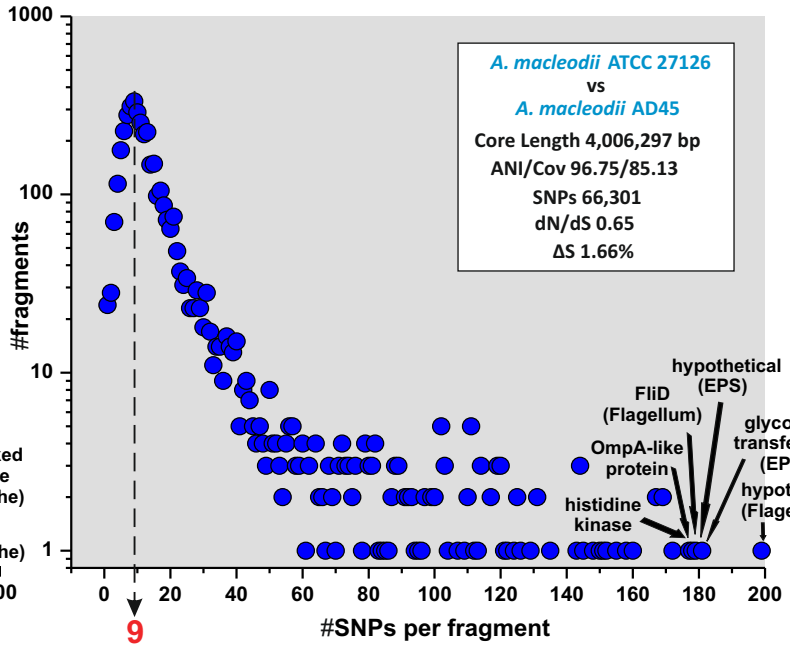


Supplementary Figure S3

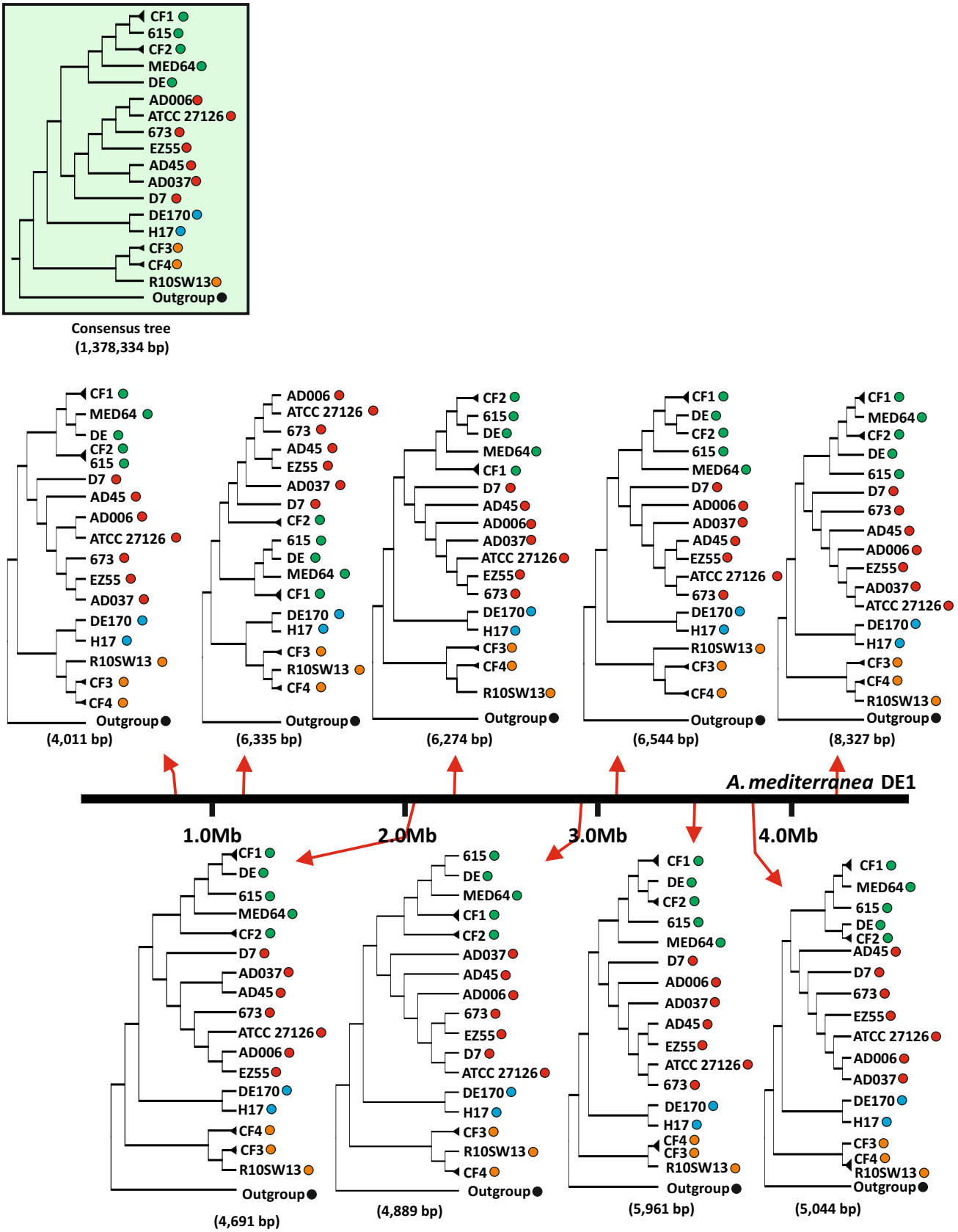








Supplementary Figure S4



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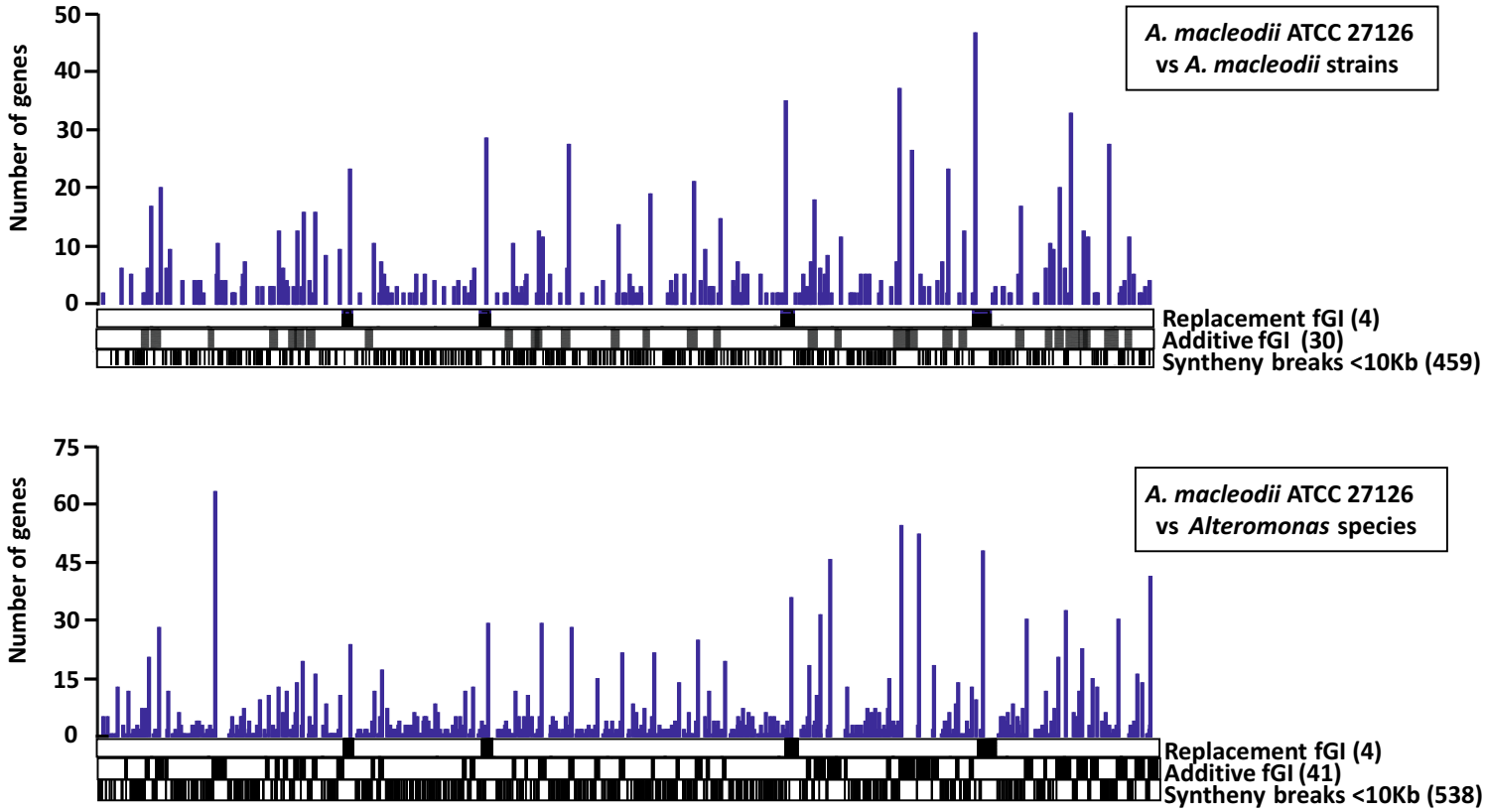

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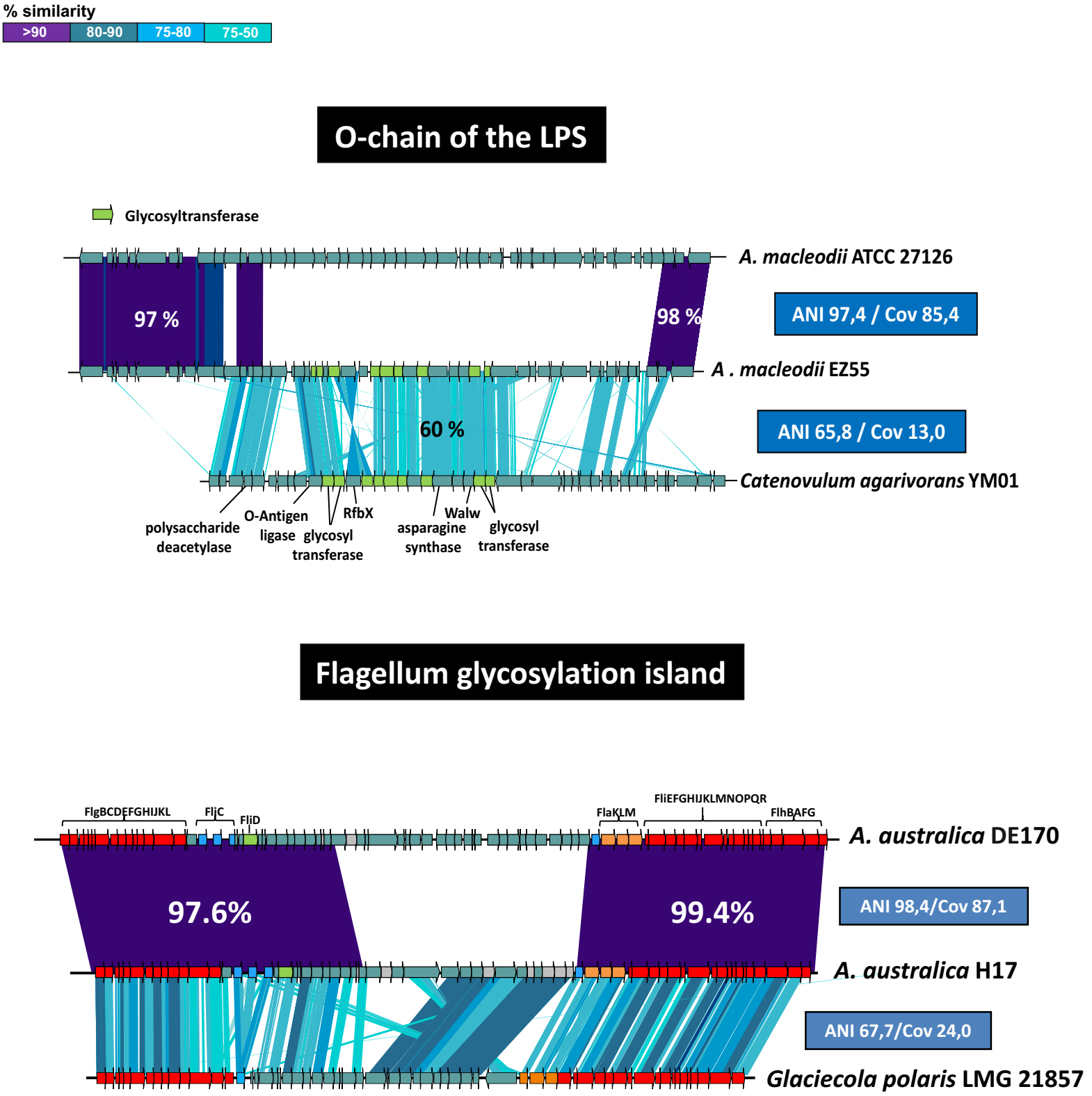

TABLE S1. *Alteromonas* isolates used in this study and their origin. Genomes that reported here for the first time are highlighted in red

***Alteromonas* strain**	**Origen isolated**	**Location**	**Date**	**Depth of Sample (m)**
A. mediterranea				
DE^T	Adriatic Sea	41°37′N, 17°22′E	2003	1000
DE1	Adriatic Sea	41°37′N, 17°22′E	2003	1000
615	English Channel	50º15'N, 4º16'W	2005	Surface
U4	Ionian Sea	35º13.83'N, 21º28.29'E	1998	3475
U7	Ionian Sea	35º13.83'N, 21º28.29'E	1998	3500
U8	Ionian Sea	35º13.83'N, 21º28.29'E	1998	3500
U10	Ionian Sea	35º13.83'N, 21º28.29'E	1998	3500
UM7	Ionian Sea	35º13.83'N, 21º28.29'E	1998	3475
UM8	Ionian Sea	35º13.83'N, 21º28.29'E	1998	3475
UM4b	Ionian Sea	35º13.83'N, 21º28.29'E	1998	3455
MED64	Aegean Sea	32º81'N, 34º69'E	2000	Surface
A. macleodii				
ATCC 27126^T	Pacific Ocean	21º17´23´´N 157º58´34´´W	1972	Surface
673	English Channel	50º15'N, 4º16'W	2005	Surface
AD45	Balearic Sea	--	2003	Surface
D7	Andaman Sea	08º00'N, 98º00'E	2000	Surface
EZ55	Sargasso Sea	--	2008	Surface
AD006	Malaysia: Port Dickson	2º31´21´´N 101º48´14´´E	2014	Surface
AD037	Malaysia: Port Dickson	2º31´21´´N 101º48´14´´E	2014	Surface
A. australica				
H 17^T	Tasman Sea	37º52´10´´S 144º58´27´´E	2007	Surface
DE170	Adriatic Sea	41°37′N, 17°22′E	2003	1000
A. addita				
R10SW13^T	Sea of Japan	42º40´23´´N 131º38´42´´E	2000	Surface
A. stellipolaris				
LMG 21861^T	Antarctic Sea	66° 20·0′ S 08° 53·4' E	1999	25
LMG 21856	Antarctic Sea	66° 21·9′ S 33° 46·7′ E	1999	25
***Alteromonas* sp.**				
Mac1	Mediterranean Sea	38°25′48´´ N 0°22′25´´ W	2014	Surface
Mac2	Mediterranean Sea	38°25′48´´ N 0°22′25´´ W	2014	Surface

TABLE S2. The twenty genes with the highest dN /dS values obtained within the *A. mediterranea* strains.

A. mediterranea **DE1 coordinates**	**d*N*/d*S***	**Gene product**
3,596,902 - 3,598,002	3.97	permease YjgP/YjgQ CDS
168,106 - 168,915	3.78	hypotethical protein
2,984,023 - 2,983,268	3.43	hypothetical protein
1,513,585 - 1,512,839	3.29	Enoyl-CoA hydratase/isomerase
1,216,789 - 1,219,386	3.16	alanyl-tRNA synthetase
2,252,675 - 2,254,264	2.97	hypothetical protein TPR (tetratricopeptide repeat)
2,403,655 - 2,405,973	2.89	Predicted exporters of the RND superfamily
4,214,839 - 4,212,998	2.55	amino acid ABC transporter/signal transduction system
487,898 - 489,037	2.27	RND family efflux transporter MFP subunit
2,291,861 - 2,294,728	2.24	TonB-dependent receptor
4,295,677 - 4,298,625	2.18	TonB-dependent receptor
1,508,773 - 1,506,896	2.11	ABC transporter ATP-binding protein/permease
4,256,192 - 4,254,591	2.10	p-aminobenzoyl-glutamate transport
3,531,718 - 3,529,364	1.98	Histidine kinase (aerobic respiration control sensor protein ArcB)
2,434,188 - 2,432,557	1.96	Glutamate synthase
2,355,773 - 2,354,532	1.95	Radical SAM
2,422,725 - 2,421,418	1.92	deoxyribodipyrimidine photolyase
1,494,510 - 1,494,094	1.88	hypothetical protein
1,768,924 - 1,767,671	1.76	proton/glutamate symporter
480,575 - 482,266	1.71	gamma-glutamyltransferase