

			GGDC eDDH	Genome ID	AAIr	%BBH	OS
Flavobacterium sp. KJJ		JARP00000000					
Flavobacterium sp. URHB0058		AIJEU00000000		<a href="#">1298868.6</a>	86.8	82.4	62.1
Flavobacterium sp. Root420		LMEF00000000	27.3	<a href="#">1736533.4</a>	86.6	85.0	63.8
Flavobacterium aquidurens	RC62 - not type	JRLF00000000	26.5	<a href="#">362413.4</a>	86.1	82.2	60.9
Flavobacterium sp. Leaf82		LMMA00000000	26.8	<a href="#">1736238.4</a>	86.1	81.9	60.7
Flavobacterium sp. Fl		JQJY00000000	25.9	<a href="#">1506583.4</a>	85.9	82.6	60.9
Flavobacterium chilense			26.1	<a href="#">946677.6</a>	85.8	80.5	59.3
Flavobacterium denitrificans	DSM 15936	AUDL00000000	25.2	<a href="#">1121888.4</a>	85.8	80.3	59.0
Flavobacterium sp. Root186		LMHW00000000	26.1	<a href="#">1736485.4</a>	85.5	81.6	59.6
Flavobacterium sp. MEB061		JXQR00000000	26.6	<a href="#">1587524.5</a>	85.4	83.1	60.6
Flavobacterium sp. VMW		LHVT00000000	26.2	<a href="#">6666666.1</a>	85.2	82.2	59.7
Flavobacterium hibernum	DSM 12611	JPRK00000000	26.7	<a href="#">17252.6</a>	85.1	77.3	56.0
Flavobacterium sp. Root935		LMJL00000000	25.9	<a href="#">1736610.4</a>	84.9	82.4	59.4
Flavobacterium sp. Root901		LMJB00000000	24.9	<a href="#">1736605.4</a>	84.8	80.8	58.1
Flavobacterium reichenbachii	LMG 25512	JPRL00000000	25.3	<a href="#">362418.7</a>	84.2	81.5	57.8
Flavobacterium sp. WG21		AMYW00000000	25	<a href="#">1229487.6</a>	84.1	70.9	50.1
Flavobacterium johnsoniae	UW101	CP000685	25	<a href="#">376686.16</a>	83.7	83.7	58.6
Flavobacterium sp. CF136		AKJZ00000000	25.7	<a href="#">1144313.7</a>	83.5	78.9	52.2
Flavobacterium sp. F52		AKJZ00000000	25.7	<a href="#">1202532.8</a>	83.3	79.7	55.2
Flavobacterium sp. ABG		LEKS00000000	24.8	<a href="#">1423322.8</a>	82.5	72.4	49.3
Flavobacterium chungangense	LMG 26729	JASY00000000	24.8	<a href="#">1453505.5</a>	82.4	76.7	52.1
Flavobacterium sp. JRM	DSM 2063	JSYO00000000	22.3	<a href="#">1566022.5</a>	78.4	71.3	43.9
Flavobacterium hydatis		JPRM00000000	22.5	<a href="#">991.1</a>	78.3	71.7	43.9
Flavobacterium sp. KMS		JSYP00000000	22.2	<a href="#">1566023.5</a>	78.1	73.0	44.5
Flavobacterium sp. JGI 0001001-D01		AUOT00000000	24.9	<a href="#">1235980.5</a>	78.1	25.8	15.7
Flavobacterium glaciei		from JGI-IMG	20.9	<a href="#">386300.3</a>	75.3	52.7	29.9
Flavobacterium sp. 83		JQMS00000000	20.9	<a href="#">1131812.4</a>	75.0	58.6	32.9
Flavobacterium sp. ACAM 123		AJXL00000000	19.9	<a href="#">1189620.8</a>	73.0	54.1	28.9
Flavobacterium frigidis	PS16618	AHKF00000000	20.3	<a href="#">1086011.9</a>	72.8	58.0	30.8
Flavobacterium sp. EM1321		JNCA00000000	21	<a href="#">1492738.4</a>	72.7	57.9	30.6
Flavobacterium frigidarium	DSM 17623	AUDO00000000	20	<a href="#">1121890.4</a>	72.2	56.2	29.3
Flavobacterium succinicans	LMG 10402	JATV00000000	20.2	<a href="#">1450525.7</a>	71.9	57.5	29.7
Flavobacterium daejeonense	RCH33	AUDK00000000	20.6	<a href="#">1121887.4</a>	71.1	58.5	29.5
Flavobacterium soli	DSM 19725	AUGO00000000	20.9	<a href="#">1121897.6</a>	68.5	58.0	27.2
Flavobacterium aquatile	LMG 4008	JRHH00000000	19.6	<a href="#">1451498.1</a>	67.9	53.5	24.6
Flavobacterium saliperosum	S13 FSS13T	AVFO00000000	20	<a href="#">1341155.4</a>	67.4	46.9	21.3
Flavobacterium suncheonense	DSM 17707	AUCZ00000000	19.1	<a href="#">1121899.4</a>	67.1	45.9	20.7
Flavobacterium branchiophilum	FL 15	FQ859183	21.2	<a href="#">1034807.1</a>	67.1	47.4	21.3
Flavobacterium cauense	R2A-7	AVBI00000000	19.8	<a href="#">1341154.4</a>	67.0	49.5	22.2
Flavobacterium sp. Leaf359		LMPW00000000	20.5	<a href="#">1736351.4</a>	66.8	54.8	24.5
Flavobacterium antarcticum	DSM 19726	ATTM00000000	19.5	<a href="#">1111730.6</a>	66.8	49.5	22.1
Flavobacterium enshiense	DK69	AVCS00000000	20.1	<a href="#">1107311.4</a>	66.4	49.5	21.8
Flavobacterium limnosediminis	JC2902 FLJC2902T	AVGG00000000	20.7	<a href="#">1341181.6</a>	66.1	49.9	21.8
Flavobacterium tegetincola	DSM 22377	AUDN00000000	20.3	<a href="#">1121901.4</a>	65.9	51.7	22.5
Flavobacterium filum	DSM17961	AUDM00000000	20.2	<a href="#">1121889.4</a>	65.8	46.6	20.2
Flavobacterium sasangense	DSM 21067	JMLU00000000	19.9	<a href="#">1121896.4</a>	65.5	46.9	20.1
Flavobacterium columnare	94-081	CP003222	19.8	<a href="#">1041826.3</a>	64.2	46.3	19.1
Flavobacterium sp. 316		JYGZ00000000	19.9	<a href="#">1603293.5</a>	63.8	52.2	21.3
Flavobacterium gelidilacus	DSM 15343	AUGN00000000	19.7	<a href="#">1121891.5</a>	63.8	50.0	20.3

Flavobacterium indicum	GPTSA100-9T	HE774682	20.5	<a href="#">1094466.1</a>	63.7	47.2	19.2
Flavobacterium rivuli	DSM 21788	ARKJ00000000	18.6	<a href="#">1121895.6</a>	63.6	59.1	23.9
Flavobacterium subsaxonicum	DSM 21790	AUGP00000000	18.6	<a href="#">1121898.4</a>	63.4	62.6	25.2
Flavobacterium beibuense	RSKm HC5	JRLV00000000	18.2	<a href="#">1406840.4</a>	63.4	55.7	22.4
Flavobacterium akiainvivens	IK-1	LIYD00000000	20.6	<a href="#">1202724.4</a>	61.8	59.8	22.8
Mesoflavibacter zeaxanthinifaciens	DSM 18436	AULQ00000000	19.1	<a href="#">1121225.3</a>	58.8	45.7	15.8
Myroides odoratus		CM001437	20.4	<a href="#">929704.5</a>	57.5	47.6	15.7
Croceibacter atlanticus	HTCC2559T	CP002046	18.3	<a href="#">216432.3</a>	57.1	45.0	14.7
Gillisia limnaea	DSM 15749	AHKR00000000	19	<a href="#">865937.4</a>	56.9	49.5	16.0
Cellulophaga lytica	LIM-21	CP002534	19.5	<a href="#">867900.5</a>	56.7	51.2	16.5
Flavobacterium sp. SCGC AAA160-P02		ARTD00000000	18.7	<a href="#">1009412.6</a>	56.2	37.6	11.9
Leeuwenhoekiella blandensis	MED217	AANC00000000	18.6	<a href="#">398720.6</a>	55.8	54.5	17.0
Flavobacterium sp. B17		BACY00000000	18.8	<a href="#">95618.6</a>	51.1	50.4	13.2
Chryseobacterium gleum	ATCC 35910	ACKQ00000000	18.7	<a href="#">525257.8</a>	50.4	56.4	14.3
Flavobacterium sp. SCGC AAA536-P05		ARQV00000000	31.1	<a href="#">1009712.6</a>	46.1	10.7	2.3
Flavobacterium sp. MS220-5C		AOTS00000000	47.1	<a href="#">926447.6</a>	46.1	10.7	2.3