

Fasta Summary Table

SUBMISSION PARAMETERS

Title	CCUG 52530	Database	empro
Sequence length	1398	Sequence type	n
Program	fasta	Version	3.4t25 Sept 2, 2005
Expectation upper value	10.0	Sequence range	1-
Number of scores	50	Number of alignments	50
Word size	6	Open gap penalty	-14
Gap extension penalty	-4	Histogram	false

Alignment	DB:ID	Source	Length	Identity%	Similar%	Overlap	E()
1	EM_PRO:AY707774	Aerococcus viridans strain 8	1419	95.986	95.986	1395	2.3e-215
2	EM_PRO:AY707779	Aerococcus viridans strain A	1419	95.986	95.986	1395	2.3e-215
3	EM_PRO:AY707773	Aerococcus viridans strain 3	1419	95.986	95.986	1395	2.3e-215
4	EM_PRO:AY707775	Aerococcus viridans strain 1	1419	95.914	95.914	1395	4.8e-215
5	EM_PRO:AY707781	Aerococcus viridans strain N	1419	95.914	95.914	1395	4.8e-215
6	EM_PRO:AY707776	Aerococcus viridans strain 1	1419	95.914	95.914	1395	4.8e-215
7	EM_PRO:AY707777	Aerococcus viridans strain A	1419	95.914	95.914	1395	4.8e-215
8	EM_PRO:AY707780	Aerococcus viridans strain M	1419	95.914	95.914	1395	4.8e-215
9	EM_PRO:AY707778	Aerococcus viridans strain A	1419	95.914	95.914	1395	4.8e-215
10	EM_PRO:AY707782	Aerococcus viridans strain R	1419	95.914	95.914	1395	4.8e-215
11	EM_PRO:D87677	Pediococcus urinaeequi DNA for 1	1419	95.842	95.842	1395	7.6e-215
12	EM_PRO:M58797	Aerococcus viridans 16S ribosoma	1554	95.497	95.497	1399	7.8e-215
13	EM_PRO:AY626833	Marine bacterium YTHM-17 16S	1403	95.677	95.677	1388	2e-212
14	EM_PRO:AJ276512	Aerococcus sp. CCUG43001 16S	1499	94.711	94.711	1399	1.4e-210
15	EM_PRO:AF318316	Aerococcus sanguinicola stra	1503	94.711	94.711	1399	1.8e-210

16	EM_PRO:AY837833	<i>Aerococcus sanguinicola</i> 16S	1488	94.711	94.711	1399	1.8e-210
17	EM_PRO:AY422717	<i>Aerococcus urinae</i> isolate DK	1545	94.564	94.707	1398	2.6e-210
18	EM_PRO:AF076639	<i>Aerococcus</i> sp. LV65.5:W1 16S	1407	95.827	95.900	1366	3e-210
19	EM_PRO:AY422711	<i>Aerococcus urinae</i> isolate NL	1545	94.635	94.635	1398	3.3e-210
20	EM_PRO:AY422712	<i>Aerococcus urinae</i> isolate CD	1545	94.635	94.635	1398	4.1e-210
21	EM_PRO:AY422713	<i>Aerococcus urinae</i> isolate NL	1545	94.635	94.635	1398	4.1e-210
22	EM_PRO:AY422714	<i>Aerococcus urinae</i> isolate NL	1545	94.635	94.635	1398	4.1e-210
23	EM_PRO:AY422715	<i>Aerococcus urinae</i> isolate CD	1545	94.635	94.635	1398	4.1e-210
24	EM_PRO:AY422716	<i>Aerococcus urinae</i> isolate CC	1545	94.635	94.635	1398	4.1e-210
25	EM_PRO:U64458	<i>Aerococcus urinae</i> 16S ribosomal	1544	94.635	94.635	1398	5.3e-210
26	EM_PRO:U64459	<i>Aerococcus urinae</i> 16S ribosomal	1503	94.492	94.492	1398	7.3e-210
27	EM_PRO:U64457	<i>Aerococcus urinae</i> 16S ribosomal	1484	94.496	94.568	1399	2.9e-209
28	EM_PRO:M77819	<i>Aerococcus urinae</i> 16S ribosomal	1481	93.996	93.996	1399	1.5e-206
29	EM_PRO:Y17318	<i>Aerococcus</i> sp. CCUG28826 16S rRN	1501	93.419	93.419	1398	3.3e-205
30	EM_PRO:AY879301	<i>Abiotrophia paradjacens</i> stra	1515	93.348	93.348	1398	1.9e-203
31	EM_PRO:AY879298	<i>Granulicatella paradiacens</i> s	1514	93.348	93.348	1398	1.9e-203
32	EM_PRO:AY879304	<i>Granulicatella adiacens</i> stra	1529	93.276	93.276	1398	3.8e-203
33	EM_PRO:AY879299	<i>Granulicatella adiacens</i> stra	1513	93.276	93.276	1398	3.8e-203
34	EM_PRO:AY879302	<i>Granulicatella adiacens</i> stra	1517	93.205	93.205	1398	7.8e-203
35	EM_PRO:AY879303	<i>Granulicatella adiacens</i> stra	1517	93.205	93.205	1398	7.8e-203
36	EM_PRO:AY879300	<i>Granulicatella adiacens</i> stra	1514	93.205	93.205	1398	7.9e-203
37	EM_PRO:AY879305	<i>Granulicatella adiacens</i> stra	1518	93.133	93.133	1398	1.6e-202
38	EM_PRO:Y17005	<i>Aerococcus christensenii</i> 16S rib	1451	92.704	92.704	1398	8.7e-202
39	EM_PRO:X87150	<i>L.pasteurii</i> 16S rRNA gene	1554	92.847	92.847	1398	2.7e-201
40	EM_PRO:AY570695	<i>Bacterium</i> sp. A15 16S riboso	1438	92.847	92.847	1398	2.9e-201
41	EM_PRO:AJ306611	<i>Trichococcus flocculiformis</i>	1438	92.847	92.847	1398	2.9e-201
42	EM_PRO:AF300974	<i>Bacterium</i> B16S 16S ribosomal	1528	93.847	93.847	1414	8.4e-201
43	EM_PRO:DQ173030	<i>Bacterium</i> TSBY-79 16S riboso	1520	92.704	92.704	1398	1.2e-200
44	EM_PRO:AJ306612	<i>Trichococcus collinsii</i> 16S r	1520	92.704	92.704	1398	1.2e-200
45	EM_PRO:AF255736	<i>Nostocoida limicola</i> I 16S ri	1514	92.704	92.704	1398	1.2e-200
46	EM_PRO:AF394926	<i>Lactosphaera</i> sp. PMagG1 16S	1483	92.704	92.704	1398	1.2e-200

47	EM_PRO:AJ312375	Granulicatella adiacens part	1456	92.765	92.908	1396	2.9e-200
48	EM_PRO:AM039968	Enterococcus termitidis 16S	1512	92.951	92.951	1376	2.2e-199
49	EM_PRO:AJ296179	Ruminococcus palustris 16S r	1520	92.899	92.899	1380	2.8e-199
50	EM_PRO:Y17301	Trichococcus flocculiformis 16S	1550	92.280	92.280	1399	4.1e-198

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