

Fasta **Summary Table**

SUBMISSION PARAMETERS			
Title	CCUG 46407	Database	empro
Sequence length	467	Sequence type	n
Program	fasta	Version	3.4t23 March 18, 2004
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

Show Annotation	Fasta Result	MView	VisualFasta	XML	SUBMIT ANOTHER JOB
Show Alignments	Clear all	Check all	Invert selection	Reset	

Alignment	DB:ID	Source	Length	Identity%	Ungapped%	Overlap	E()
1 <input checked="" type="checkbox"/>	EM_PRO:ASP535636	Arthrobacter sp. SB28 16S	1487	94.444	94.850	468	9.8e-63
2 <input checked="" type="checkbox"/>	EM_PRO:ASP535637	Arthrobacter sp. SB30 16S	1467	94.444	94.850	468	9.9e-63
3 <input checked="" type="checkbox"/>	EM_PRO:AY636111	Kazacoccus imtecheus strain	1488	94.231	95.043	468	1.2e-62
4 <input checked="" type="checkbox"/>	EM_PRO:DSP244775	Dermatophilus like sp. V4.	1452	94.231	94.635	468	2e-62
5 <input checked="" type="checkbox"/>	EM_PRO:AY382898	Serinicoccus marinus JC1078	1421	94.255	95.064	470	2.8e-62
6 <input checked="" type="checkbox"/>	EM_PRO:AS16SR483	Arthrobacter sp. 16S ribosom	1484	93.843	94.850	471	7.3e-62
7 <input checked="" type="checkbox"/>	EM_PRO:AS16SR537	Arthrobacter sp. 16S ribosom	1484	93.843	94.850	471	7.3e-62
8 <input checked="" type="checkbox"/>	EM_PRO:AS16SR445	Arthrobacter cumminsi 16S r	1483	93.843	94.850	471	7.3e-62
9 <input checked="" type="checkbox"/>	EM_PRO:BC16SRDNA	B.caseolytica 16S rRNA gene	1476	93.803	94.206	468	7.9e-62
10 <input checked="" type="checkbox"/>	EM_PRO:AB070602	Arthrobacter sp. JCM 1339 g	1440	93.590	93.991	468	1.6e-61
11 <input checked="" type="checkbox"/>	EM_PRO:AF409020	Arthrobacter sp. Ellin178 1	1432	93.590	93.991	468	1.6e-61
12 <input checked="" type="checkbox"/>	EM_PRO:AY244783	Arthrobacter cumminsi 16S	1160	93.418	94.421	471	1.7e-61
13 <input checked="" type="checkbox"/>	EM_PRO:AJ717366	Citrococcus sp. AC7 16S rRN	1506	93.376	93.777	468	3.1e-61
14 <input checked="" type="checkbox"/>	EM_PRO:AY376164	Citricoccus alkalitolerans	1480	93.376	93.777	468	3.2e-61
15 <input checked="" type="checkbox"/>	EM_PRO:MSP244665	Micrococcus sp. V4.MO.20 p	1449	93.376	93.777	468	3.2e-61
16 <input checked="" type="checkbox"/>	EM_PRO:DQ108399	Citrococcus sp. Tibet-ITa3	1436	93.376	93.777	468	3.2e-61
17 <input checked="" type="checkbox"/>	EM_PRO:AY572475	Arthrobacter sp. R33S 16S r	1520	93.590	94.397	468	4.6e-61
18 <input checked="" type="checkbox"/>	EM_PRO:ASP243421	Arthrobacter albus partial	1501	93.023	94.421	473	5.4e-61
19 <input checked="" type="checkbox"/>	EM_PRO:AF256196	Arthrobacter keyseri 12B 16	1452	93.617	94.421	470	6.4e-61
20 <input checked="" type="checkbox"/>	EM_PRO:AB127968	Micrococcus sp. YGJ1 gene f	1439	93.617	94.421	470	6.5e-61
21 <input checked="" type="checkbox"/>	EM_PRO:AB117515	Arthrobacter dextranolyticu	1414	93.162	93.562	468	6.5e-61
22 <input checked="" type="checkbox"/>	EM_PRO:AY572482	Arthrobacter sp. R36S 16S r	1521	93.376	94.181	468	9.2e-61
23 <input checked="" type="checkbox"/>	EM_PRO:AF197021	Arthrobacter sp. 'SMCC G919	1331	93.191	93.991	470	9.3e-61
24 <input checked="" type="checkbox"/>	EM_PRO:AF235090	Arthrobacter methylotrophus	1459	93.376	94.181	468	9.4e-61
25 <input checked="" type="checkbox"/>	EM_PRO:ASP491107	Arthrobacter gandensis par	1450	93.376	94.181	468	9.5e-61

26	<input checked="" type="checkbox"/>	EM_PRO:ASP491108	Arthrobacter gandensis par	1445	93.376	94.181	468	9.5e-61
27	<input checked="" type="checkbox"/>	EM_PRO:AB017650	Arthrobacter sp. gene for 1	1445	93.376	94.181	468	9.5e-61
28	<input checked="" type="checkbox"/>	EM_PRO:ASP316140	Arthrobacter gandensis 16S	1445	93.376	94.181	468	9.5e-61
29	<input checked="" type="checkbox"/>	EM_PRO:AY436791	Arthrobacter sp. C4 16S rib	1444	93.376	94.181	468	9.5e-61
30	<input checked="" type="checkbox"/>	EM_PRO:AF479328	Glacial ice bacterium Trans	1434	93.376	94.181	468	9.5e-61
31	<input checked="" type="checkbox"/>	EM_PRO:AY452082	Arthrobacter sp. AS19 16S r	1426	93.376	94.181	468	9.6e-61
32	<input checked="" type="checkbox"/>	EM_PRO:AY177353	Arthrobacter sp. I4 16S rib	1358	93.376	94.181	468	9.9e-61
33	<input checked="" type="checkbox"/>	EM_PRO:AY748914	Arthrobacter sp. NIPHL09090	1349	93.376	94.181	468	9.9e-61
34	<input checked="" type="checkbox"/>	EM_PRO:U96186	Rathayibacter rathayi 16S ribos	1272	93.376	94.181	468	1e-60
35	<input checked="" type="checkbox"/>	EM_PRO:AY628691	Arthrobacter sp. AD25 16S r	1491	92.812	94.206	473	1.1e-60
36	<input checked="" type="checkbox"/>	EM_PRO:AY628690	Arthrobacter sp. AD12 16S r	1491	92.812	94.206	473	1.1e-60
37	<input checked="" type="checkbox"/>	EM_PRO:AURDNA16	A.ureafaciens 16S rDNA	1479	92.812	94.206	473	1.1e-60
38	<input checked="" type="checkbox"/>	EM_PRO:DQ117533	Bacterium #WM-A3 16S riboso	1436	92.812	94.206	473	1.1e-60
39	<input checked="" type="checkbox"/>	EM_PRO:AY561572	Arthrobacter sp. RG-5 16S r	1004	93.376	94.181	468	1.2e-60
40	<input checked="" type="checkbox"/>	EM_PRO:AY561590	Arthrobacter sp. RG-24 16S	998	93.376	94.181	468	1.2e-60
41	<input checked="" type="checkbox"/>	EM_PRO:AY561579	Arthrobacter sp. RG-13 16S	994	93.376	94.181	468	1.2e-60
42	<input checked="" type="checkbox"/>	EM_PRO:AY188937	Micrococcus sp. P_wp0222 16	1490	93.177	93.777	469	1.3e-60
43	<input checked="" type="checkbox"/>	EM_PRO:ASP315070	Arthrobacter monumenti 16S	1484	93.404	94.206	470	1.3e-60
44	<input checked="" type="checkbox"/>	EM_PRO:AY561618	Actinobacterium RG-59 16S r	922	93.376	94.181	468	1.3e-60
45	<input checked="" type="checkbox"/>	EM_PRO:AF102267	Arthrobacter chlorophenolic	1446	92.949	93.348	468	1.3e-60
46	<input checked="" type="checkbox"/>	EM_PRO:AY327845	Streptomyces sp. MSU 2110 1	912	93.376	94.181	468	1.3e-60
47	<input checked="" type="checkbox"/>	EM_PRO:AF501368	Kocuria kristinae strain HA	810	93.376	94.181	468	1.4e-60
48	<input checked="" type="checkbox"/>	EM_PRO:AF323746	Kocuria kristinae 16S ribos	768	93.376	94.181	468	1.5e-60
49	<input checked="" type="checkbox"/>	EM_PRO:AB211026	Actinomycetales bacterium S	1529	92.979	93.777	470	1.7e-60
50	<input checked="" type="checkbox"/>	EM_PRO:AY345527	Bacterium W8 16S ribosomal	1517	93.162	93.966	468	1.8e-60

[Return to site](#)