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FASTA Results

SUBMISSION PARAMETERS

Title	58157	Database	em_rel_pro
Sequence length	978	Sequence type	n
Program	fasta	Version	35.04 Mar. 8, 2009
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

<u>Alignment</u>	<u>DB:ID</u>	<u>Source</u>	<u>Length</u>	<u>Identity%</u>	<u>Similar%</u>	<u>Overlap</u>	<u>E()</u>
2	EM_PRO:AY362908	Haemophilus parainfluenzae s	1362	98.2	99.2	977	6.1e-217
1	EM_PRO:EU083530	Haemophilus parainfluenzae s	1505	98.6	99.0	977	5.8e-217
3	EM_PRO:AB098612	TerraHaemophilus aromatico	1502	98.2	98.6	979	4.5e-215
5	EM_PRO:AY365450	Haemophilus parainfluenzae s	1322	98.2	98.6	960	5.3e-211
4	EM_PRO:AJ290754	Haemophilus pittmaniae parti	1466	97.1	97.6	978	1.5e-211
6	EM_PRO:AJ290755	Haemophilus pittmaniae parti	1468	97.1	97.6	980	2.1e-210
7	EM_PRO:M75082	Haemophilus paraphrophilus 16S s	1479	96.3	96.7	977	1.2e-209
8	EM_PRO:M75081	Haemophilus parainfluenzae 16S s	1483	96.0	96.4	977	1.1e-208
9	EM_PRO:DQ851143	Haemophilus simiae strain RO	1540	95.9	96.3	976	1e-205
10	EM_PRO:DQ301927	Mannheimia varigena strain H	1503	95.4	95.8	976	1.4e-203
12	EM_PRO:AF224287	Pasteurella haemolytica CCUG	1519	95.3	95.7	976	3.7e-203
13	EM_PRO:DQ301926	Mannheimia varigena strain 3	1512	95.3	95.7	976	3.7e-203
27	EM_PRO:DQ211725	Actinobacillus rossii strain	1364	94.9	95.7	978	7.2e-202
29	EM_PRO:DQ211727	Actinobacillus rossii strain	1364	94.8	95.7	978	1e-201
30	EM_PRO:DQ211730	Actinobacillus rossii strain	1364	94.8	95.7	978	1e-201
31	EM_PRO:AY362895	Actinobacillus rossii strain	1364	94.8	95.7	978	1e-201
36	EM_PRO:DQ211721	Actinobacillus rossii strain	1364	94.5	95.7	978	1.9e-201
43	EM_PRO:DQ211722	Actinobacillus rossii strain	1362	94.3	95.7	978	3.7e-201
11	EM_PRO:EU909639	Haemophilus sp. 11p18 16S ri	1362	95.1	95.6	978	1.8e-203
15	EM_PRO:AF053901	Mannheimia granulomatis stra	1504	95.2	95.6	976	9.8e-203
26	EM_PRO:AF224289	Actinobacillus rossii 16S ri	1535	95.1	95.6	978	6.9e-202
33	EM_PRO:EU909645	Haemophilus sp. 65p13 16S ri	1362	93.5	95.6	978	1.4e-201
40	EM_PRO:DQ211720	Actinobacillus rossii strain	1364	94.7	95.6	978	2.7e-201
14	EM_PRO:U57072	Pasteurella haemolytica strain P	1472	95.2	95.5	977	8e-203
18	EM_PRO:AF053899	Mannheimia varigena strain C	1504	95.1	95.5	976	2.6e-202

45	EM_PRO:DQ666549	Actinobacillus capsulatus st	1362	94.0	95.5	976	3.7e-201
47	EM_PRO:AY425295	Mannheimia sp. BJ3956.1 16S	1356	95.2	95.5	970	3.7e-201
16	EM_PRO:U57076	Pasteurella CCUG16500 16S rRNA g	1476	95.0	95.4	976	1.9e-202
17	EM_PRO:AY362914	Mannheimia sp. NCTC 11313 16	1363	95.1	95.4	977	2.2e-202
19	EM_PRO:AF053898	Mannheimia sp. HPA121 16S ri	1505	95.1	95.4	976	3.6e-202
20	EM_PRO:EU909667	Haemophilus sp. CCUG 43251 1	1362	94.6	95.4	978	4.7e-202
25	EM_PRO:DQ301925	Mannheimia granulomatis stra	1670	95.0	95.4	976	6.6e-202
37	EM_PRO:AY362904	Bisgaard Taxon 7 strain CCUG	1362	94.7	95.4	976	1.9e-201
39	EM_PRO:AY431031	Actinobacillus rossii strain	1464	95.1	95.4	978	2.6e-201
21	EM_PRO:AF224307	Haemophilus quentini 16S rib	1535	95.1	95.3	976	5e-202
22	EM_PRO:EU909641	Haemophilus sp. 16N 16S ribo	1362	95.1	95.3	976	5.2e-202
23	EM_PRO:EU909640	Haemophilus sp. 11PS 16S rib	1362	95.1	95.3	976	5.2e-202
24	EM_PRO:AY962106	Haemophilus quentini strain	1335	95.1	95.3	976	5.3e-202
28	EM_PRO:AF053893	Mannheimia varigena strain 1	1504	95.0	95.3	976	9.6e-202
32	EM_PRO:AY465370	Mannheimia sp. BA597/9 16S r	1362	95.0	95.3	976	1e-201
41	EM_PRO:EU909646	Haemophilus sp. CCUG 11096 1	1362	93.5	95.3	978	3.3e-201
49	EM_PRO:AF224290	Pasteurella aerogenes MCCM 0	1535	95.0	95.3	978	6.7e-201
50	EM_PRO:DQ211726	Actinobacillus rossii strain	1364	95.0	95.3	978	7e-201
34	EM_PRO:X87979	H.influenzae DNA for ribosomal R	1442	95.0	95.2	975	1.7e-201
35	EM_PRO:U57071	Pasteurella haemolytica strain P	1471	94.9	95.2	976	1.9e-201
38	EM_PRO:AF053894	Mannheimia sp. R19.2 16S rib	1505	94.9	95.2	976	2.5e-201
42	EM_PRO:AF247720	Actinobacillus equuli strain	1461	94.9	95.1	976	3.6e-201
44	EM_PRO:EU909642	Haemophilus sp. 26E 16S ribo	1362	94.9	95.1	976	3.7e-201
46	EM_PRO:AY465359	Actinobacillus equuli strain	1362	94.9	95.1	976	3.7e-201
48	EM_PRO:U57070	Pasteurella haemolytica strain P	1471	94.8	95.1	976	4.9e-201