

Fusobacterium sp HMT 248 homd HMT 248

Items order: Presence absence (D: Euclidean; L: Ward) | Current view: gene_cluster_presence_absence | Samples order: gene_cluster frequencies

	Gene cluster	Sample
1	1	1
2	2	2
3	3	3
4	4	4
5	5	5
6	6	6
7	7	7
8	8	8
9	9	9
10	10	10
11	11	11
12	12	12
13	13	13
14	14	14
15	15	15
16	16	16
17	17	17
18	18	18
19	19	19
20	20	20
21	21	21
22	22	22
23	23	23
24	24	24
25	25	25
26	26	26
27	27	27
28	28	28
29	29	29
30	30	30
31	31	31
32	32	32
33	33	33
34	34	34
35	35	35
36	36	36
37	37	37
38	38	38
39	39	39
40	40	40
41	41	41
42	42	42
43	43	43
44	44	44
45	45	45
46	46	46
47	47	47
48	48	48
49	49	49
50	50	50
51	51	51
52	52	52
53	53	53
54	54	54
55	55	55
56	56	56
57	57	57
58	58	58
59	59	59
60	60	60
61	61	61
62	62	62
63	63	63
64	64	64
65	65	65
66	66	66
67	67	67
68	68	68
69	69	69
70	70	70
71	71	71
72	72	72
73	73	73
74	74	74
75	75	75
76	76	76
77	77	77
78	78	78
79	79	79
80	80	80
81	81	81
82	82	82
83	83	83
84	84	84
85	85	85
86	86	86
87	87	87
88	88	88
89	89	89
90	90	90
91	91	91
92	92	92
93	93	93
94	94	94
95	95	95
96	96	96
97	97	97
98	98	98
99	99	99
100	100	100

Fusobacterium sp HMT 248 str Marseille P2749 id GCA 900095705 1
Fusobacterium sp HMT 248 str MGYG HGUT 01555 id GCA 902376175 1
COG20 FUNCTION
KOfam
Pfam

CAZyme

UNKNOWN (1568) KNOWN (40)

COG20 CATEGORY

KNOWN (1294) UNKNOWN (314)

COG20 FUNCTION

KNOWN (1294) UNKNOWN (314)

COG20 PATHWAY

UNKNOWN (1180) KNOWN (428)

KEGG BRITE

KNOWN (1033) UNKNOWN (575)

KEGG Class

UNKNOWN (1362) KNOWN (246)

KEGG Module

UNKNOWN (1362) KNOWN (246)

KOfam

KNOWN (1036) UNKNOWN (572)

Pfam

KNOWN (1351) UNKNOWN (257)