

Protein ID
Cluster of Class-3 porin protein (Fragment) OS=Neisseria meningitidis GN=porB PE=3 SV=1 (Q51147_NEIME)
Cluster of Outer membrane protein class 4 OS=Neisseria meningitidis NM2795 GN=ompM PE=3 SV=1 (A0A125WFE9_NEIME)
Cluster of Iron-regulated outer membrane protein (Fragment) OS=Neisseria meningitidis GN=IRA PE=3 SV=1 (Q84029_NEIME)
Cluster of Factor I binding protein (Fragment) OS=Neisseria meningitidis GN=ira1870 PE=4 SV=1 (Q30375_NEIME)
Cluster of Periplasmic iron-binding protein (Fragment) OS=Neisseria gonorrhoeae GN=fbp PE=4 SV=1 (Q50988_NEIGO)
Cluster of IgA-specific serine endopeptidase OS=Neisseria meningitidis (strain alpha14) GN=iga1 PE=4 SV=1 (C65SXX1_NEIML)
Cluster of PorA OS=Neisseria meningitidis GN=porA PE=3 SV=1 (QJ0P11_NEIME)
Cluster of Transferrin binding protein 1 (Fragment) OS=Neisseria meningitidis GN=tbpA PE=3 SV=1 (Q53136_NEIME)
Cluster of Thiol-disulfide interchange protein OS=Neisseria meningitidis serogroup C / serotype 2a (strain ATCC 700532 / DSM 15464 / FAM18) GN=NMCO490 PE=3 SV=1 (A1KSP9_NEIMF)
Cluster of Elongation factor Tu OS=Neisseria meningitidis LNP21362 GN=elf PE=3 SV=1 (A0A0AFA088_NEIME)
Cluster of Gamma-glutamyltransferase OS=Neisseria meningitidis LNP21362 GN=N875_01000 PE=4 SV=1 (A0A0A8F236_NEIME)
Cluster of Transferrin binding protein B (Fragment) OS=Neisseria meningitidis serogroup B / serotype 15 (strain H44/76) PE=4 SV=1 (A0A0E3SU61_NEIMH)
Cluster of Outer membrane protein assembly factor BAmA OS=Neisseria gonorrhoeae GN=ompB8 PE=3 SV=1 (P95359_NEIGO)
Cluster of Class 5c outer membrane protein (Fragment) OS=Neisseria meningitidis GN=opc PE=4 SV=1 (Q51228_NEIME)
Cluster of Type IV pilus biogenesis and competence protein PilQ OS=Neisseria meningitidis serogroup B / serotype 15 (strain H44/76) GN=pilQ PE=3 SV=2 (PLQ_NEIMH)
Cluster of Peptidylprolyl isomerase OS=Neisseria meningitidis GN=CNQ34_01945 PE=4 SV=1 (A0A2A3LV89_NEIME)
Cluster of PIIS cassette OS=Neisseria meningitidis (strain alpha14) GN=pilS2 PE=4 SV=1 (C6S9N9_NEIML)
Cluster of 50S ribosomal protein L2 OS=Neisseria meningitidis LNP21362 GN=rplB PE=3 SV=1 (A0A0A8FAA4_NEIME)
Uncharacterized protein OS=Neisseria meningitidis LNP21362 GN=N875_00970 PE=4 SV=1 (A0A0A8F509_NEIME)
Cluster of Putative lipoprotein OS=Neisseria meningitidis NM2795 GN=NMEN2795_1510 PE=4 SV=1 (A0A125WBX2_NEIME)
Cluster of Putative extracellular heme-binding protein OS=Neisseria meningitidis alpha275 GN=haeR PE=4 SV=1 (C6S154_NEIME)
Uncharacterized protein OS=Neisseria meningitidis LNP21362 GN=N875_00975 PE=4 SV=1 (A0A0A8F313_NEIME)
Cluster of Hemoglobin receptor (Fragment) OS=Neisseria meningitidis GN=hmbR PE=3 SV=1 (BDY09_NEIME)
Cluster of Uncharacterized protein OS=Neisseria meningitidis LNP21362 GN=N875_00075 PE=4 SV=1 (A0A0A8F313_NEIME)
Cluster of Uncharacterized protein OS=Neisseria meningitidis (strain alpha14) GN=NMO_1093 PE=4 SV=1 (C6S797_NEIML)
Cluster of Adhesin OS=Neisseria meningitidis LNP21362 GN=N875_02500 PE=4 SV=1 (A0A0A8F509_NEIME)
Cluster of Alcohol dehydrogenase OS=Neisseria meningitidis serogroup B (strain alpha170) GN=radA PE=3 SV=1 (E3D230_NEIMT)
Cluster of Cysteine synthase OS=Neisseria meningitidis alpha153 GN=cysK3 PE=4 SV=1 (C6S808_NEIME)
Cluster of Putative lipoprotein OS=Neisseria meningitidis NM2795 GN=NMEN2795_1176 PE=4 SV=1 (A0A125WD37_NEIME)
Cluster of 50S ribosomal protein L3 OS=Neisseria meningitidis LNP21362 GN=rplP PE=3 SV=1 (A0A0A8F829_NEIME)
Cluster of Peptidyl-prolyl cis-trans isomerase OS=Neisseria meningitidis LNP21362 GN=N875_08420 PE=4 SV=1 (A0A0A8F8Q5_NEIME)
Cluster of Phosphoglycerate kinase OS=Neisseria meningitidis LNP21362 GN=pgk PE=3 SV=1 (A0A0A8F8R2_NEIME)
Cluster of Putative serotype P-specific adhesin OS=Neisseria B (strain MC58) GN=NMB1669 PE=4 SV=1 (QJ0XM7_NEIMB)
Cluster of LysM domain protein OS=Neisseria meningitidis ATCC 13091 GN=HMPREF0602_2194 PE=4 SV=1 (E0NCQ2_NEIME)
Cluster of Lactoferrin-binding protein B OS=Neisseria meningitidis NM2795 GN=hbp PE=4 SV=1 (A0A125WC21_NEIME)
Cluster of Iron-regulated protein FrpA OS=Neisseria meningitidis serogroup C GN=frpA PE=4 SV=1 (FRPA_NEIMC)
Cluster of 50S ribosomal protein LNP21362 GN=N875_01000 PE=3 SV=1 (A0A0A8F3W7_NEIME)
Cluster of Peptidase OS=Neisseria meningitidis GN=iga_2 PE=4 SV=1 (A0A0Y6SUJ0_NEIME)
Cluster of Amino acid ABC transporter substrate-binding protein OS=Neisseria meningitidis LNP21362 GN=N875_03055 PE=3 SV=1 (A0A0A8F2T0_NEIME)
Cluster of Phosphate acetyltransferase OS=Neisseria meningitidis alpha522 GN=pta PE=3 SV=1 (ME795_NEIME)
Cluster of 50S ribosomal protein L15 OS=Neisseria meningitidis LNP21362 GN=rplO PE=3 SV=1 (A0A0A8F489_NEIME)
Cluster of Type IV pilus biogenesis/stability protein PilW OS=Neisseria meningitidis NM2795 GN=pilF PE=4 SV=1 (A0A125WD38_NEIME)
Cluster of Hemagglutinin-hemolysin-related protein TpsA2 OS=Neisseria meningitidis 93004 GN=NMEN53004_1151 PE=4 SV=1 (J8VLH5_NEIME)
Cluster of Peptidyl-prolyl isomerase OS=Neisseria C (strain NZ-05/33) GN=NMCC_1470 PE=3 SV=1 (A9M0Z7_NEIMD)
Cluster of Possible lipoprotein OS=Neisseria meningitidis alpha153 GN=yaP PE=4 SV=1 (C6S6E8_NEIME)
Cluster of NlpB (Fragment) OS=Neisseria meningitidis GN=nlpB PE=4 SV=1 (A7LA07_NEIME)
Cluster of 50S ribosomal protein L24 OS=Neisseria meningitidis GN=rplX PE=3 SV=1 (A0A2A3LQ61_NEIME)
Cluster of 60 chaperonin OS=Neisseria meningitidis LNP21362 GN=groEL PE=3 SV=1 (A0A0A8F8D9_NEIME)
Cluster of Lipoprotein OS=Neisseria meningitidis GN=gna1946 PE=3 SV=1 (XSEP09_NEIME)
Cluster of Carboxy-terminal processing protease CtpA OS=Neisseria meningitidis NM2795 GN=ctpA2 PE=3 SV=1 (A0A125WDK9_NEIME)
Cluster of LPS assembly protein OS=Neisseria meningitidis B (strain MC58) GN=nmrD PE=1 SV=1 (LPTD_NEIME)
Cluster of Thiolate-binding protein OS=Neisseria meningitidis alpha522 GN=tpa PE=4 SV=1 (MEK21_NEIME)
Cluster of Protein Crib OS=Neisseria meningitidis GN=ERS514410_00524 PE=4 SV=1 (A0A0Y5KUX7_NEIME)
Cluster of Uncharacterized protein OS=Neisseria meningitidis LNP21362 GN=N875_05160 PE=4 SV=1 (A0A0A8F3W2_NEIME)
Cluster of Elongation factor Tu OS=Neisseria meningitidis LNP21362 GN=efa PE=3 SV=1 (A0A0A8F479_NEIME)
Cluster of IgA-specific serine endopeptidase OS=Neisseria meningitidis alpha153 GN=pilC PE=4 SV=1 (C6S2Q6_NEIME)
Cluster of Phosphoenolpyruvate synthase OS=Neisseria meningitidis 93004 GN=ppsA PE=3 SV=1 (J8WMZ9_NEIME)
Cluster of Uncharacterized protein OS=Neisseria meningitidis ATCC 13091 GN=HMPREF0602_0818 PE=4 SV=1 (E0N7Y8_NEIME)
Cluster of 50S ribosomal protein LNP21362 GN=rplU PE=3 SV=1 (A0A0A8F8F2_NEIME)
Cluster of Uncharacterized protein OS=Neisseria meningitidis LNP21362 GN=N875_01760 PE=3 SV=1 (A0A0A8F7M1_NEIME)
Cluster of Succinate-CoA ligase [ADP-forming] subunit beta OS=Neisseria meningitidis (strain alpha14) GN=sucC PE=3 SV=1 (C6S6M6_NEIML)
Cluster of Putative outer membrane protein OS=Neisseria meningitidis 93004 GN=NMEN53004_0126 PE=4 SV=1 (BVE1E1_NEIME)
Cluster of TPR repeat-containing protein OS=Neisseria meningitidis M0579 GN=M0579_00675 PE=4 SV=1 (A0A0A8BYD5_NEIME)
Cluster of Multifunctional fusion protein OS=Neisseria meningitidis GN=nmrB PE=3 SV=1 (A0A1V3SQW6_NEIME)
Cluster of 50S ribosomal protein L4 OS=Neisseria meningitidis serogroup B (strain NZ-05/33) GN=rplD PE=3 SV=1 (A0A0E0TN23_NEIMN)
Cluster of 2,3-bisubstrate-dependent dehydrogenase OS=Neisseria meningitidis LNP21362 GN=N875_04030 PE=3 SV=1 (N875_04030_NEIME)
Cluster of Ketol-acid reductoisomerase (NADP(+)) OS=Neisseria meningitidis NM2795 GN=ilvC PE=3 SV=1 (A0A125WC25_NEIME)
Cluster of Putrescine-binding periplasmic protein OS=Neisseria meningitidis GN=pod1 PE=3 SV=1 (XSFBA2_NEIME)
Cluster of Succinate-CoA ligase [ADP-forming] subunit alpha OS=Neisseria meningitidis GN=sucC PE=3 SV=1 (A0A0Y6U395_NEIME)
Cluster of Chaperone SurA OS=Neisseria meningitidis 93004 GN=surA PE=4 SV=1 (J8VH0Q_NEIME)
Cluster of 30S ribosomal protein S12 OS=Neisseria meningitidis serogroup B (strain NZ-05/33) GN=psl PE=3 SV=1 (A0A0E0TMP2_NEIMN)
Cluster of Glyceraldehyde-3-phosphate dehydrogenase OS=Neisseria meningitidis serogroup B (strain MC58) GN=gapA-2 PE=3 SV=1 (QJ0X95_NEIME)
Trypan OS=Sus scrofa PE=1 SV=1
Cluster of Chaperone protein DnaK OS=Neisseria meningitidis serogroup B / serotype 15 (strain H44/76) GN=dnaK PE=2 SV=1 (ESMV09_NEIMH)
Cluster of ABC transporter substrate-binding protein OS=Neisseria meningitidis LNP21362 GN=N875_06710 PE=4 SV=1 (A0A0A8FA15_NEIME)
Cluster of ABC transporter substrate-binding protein OS=Neisseria meningitidis LNP21362 GN=N875_09885 PE=3 SV=1 (A0A0A8F7F8_NEIME)
Cluster of Elongation factor Tu OS=Neisseria meningitidis serogroup B / serotype 15 (strain H44/76) GN=efa PE=3 SV=1 (ESM2J4_NEIMF)
Cluster of Peptidyl-prolyl cis-trans isomerase OS=Neisseria meningitidis LNP21362 GN=N875_03035 PE=3 SV=1 (A0A0A8F859_NEIME)
Cluster of Transketolase OS=Neisseria meningitidis 81858 GN=kt PE=3 SV=1 (ROU033_NEIME)
Cluster of Porin OS=Neisseria meningitidis GN=porB PE=3 SV=1 (B7S1T2_NEIME)
Cluster of Thioredoxin OS=Neisseria meningitidis GN=trxB PE=4 SV=1 (A0A1V0G5E8_NEIME)
Keratin, type I cytoskeletal 1 OS=Homo sapiens GN=KRT1 PE=1 SV=6
Keratin, type I cytoskeletal 2 epidermal OS=Homo sapiens GN=KRT2 PE=1 SV=2
Uncharacterized protein OS=Neisseria meningitidis GN=A6J54_02205 PE=4 SV=1
Cluster of Putative membrane protein OS=Neisseria meningitidis (strain alpha14) GN=NMO_0724 PE=4 SV=1 (C6S699_NEIML)
Cluster of TonB-dependent receptor OS=Neisseria meningitidis LNP21362 GN=N875_08420 PE=3 SV=1 (A0A0A8F536_NEIME)
Cluster of ATP synthase subunit alpha OS=Neisseria meningitidis GN=atpA PE=3 SV=1 (A0A121T6Z3_NEIME)
Cluster of Endolase OS=Neisseria meningitidis LNP21362 GN=eno PE=3 SV=1 (A0A0A8F703_NEIME)
Cluster of Glutamate dehydrogenase OS=Neisseria meningitidis LNP21362 GN=N875_09320 PE=3 SV=1 (A0A0A8F716_NEIME)
Cluster of Thiol-disulfide interchange protein OS=Neisseria meningitidis NM2795 GN=dsbA3 PE=3 SV=1 (A0A0A125WFY0_NEIME)
Cluster of Thioredoxin OS=Neisseria meningitidis serogroup B PE=3 SV=1 (A0A0A8Q426_NEIM)
Cluster of 50S ribosomal protein L7/L12 OS=Neisseria meningitidis LNP21362 GN=rplL PE=3 SV=1 (A0A0A8FA86_NEIME)
Cluster of ComE operon protein 1-related protein OS=Neisseria meningitidis serogroup B (strain MC58) GN=NMB1657 PE=4 SV=1 (QJ0YB8_NEIMB)
Cluster of Pyruvate dehydrogenase E1 component OS=Neisseria meningitidis (strain alpha14) GN=pacE PE=4 SV=1 (C6S7H6_NEIML)
30S ribosomal protein S21 OS=Neisseria meningitidis LNP21362 GN=rpsL PE=3 SV=1
50S ribosomal protein L14 OS=Neisseria meningitidis LNP21362 GN=rplN PE=3 SV=1
Cluster of 47 lipoprotein OS=Neisseria meningitidis PE=4 SV=1 (QJ0MP9_NEIME)
Cluster of 50S ribosomal protein L1 OS=Neisseria meningitidis LNP21362 GN=rplA PE=3 SV=1 (A0A0Y6WE99_NEIME)
Cluster of Putative hydrolase OS=Neisseria meningitidis (strain alpha14) GN=NMO_1967 PE=4 SV=1 (C6S184_NEIM)
Cluster of Trigger factor OS=Neisseria meningitidis serogroup B (strain NZ-05/33) GN=trg PE=3 SV=1 (A0A0E0TQY5_NEIMN)
LPS-assembly lipoprotein Lpe OS=Neisseria meningitidis LNP21362 GN=lpe PE=3 SV=1
30S ribosomal protein S4 OS=Neisseria meningitidis LNP21362 GN=rpsD PE=3 SV=1
50S ribosomal protein L27 OS=Neisseria meningitidis LNP21362 GN=rplM PE=3 SV=1
Cluster of Lipoprotein NlpD, putative OS=Neisseria meningitidis GN=nlpD PE=4 SV=1 (A0A0Y5C793_NEIME)
Cluster of NADH-ubiquinone oxidoreductase OS=Neisseria meningitidis LNP21362 GN=N875_03675 PE=4 SV=1 (A0A0A8FA41_NEIME)
Cluster of Protein-export protein SecE OS=Neisseria meningitidis LNP21362 GN=secE PE=3 SV=1 (A0A0A8FAU2_NEIME)
Cluster of Uncharacterized protein OS=Neisseria meningitidis NM2795 GN=NMEN2795_1566 PE=4 SV=1 (A0A125WCH0_NEIME)
50S ribosomal protein L29 OS=Neisseria meningitidis LNP21362 GN=rpmC PE=3 SV=1
Cluster of Heme ABC exporter, ATP-binding protein CmaA OS=Neisseria meningitidis NM2795 GN=NMEN2795_0410 PE=4 SV=1 (A0A125WFA8_NEIME)
Cluster of Outer membrane protein assembly factor BamD OS=Neisseria meningitidis LNP21362 GN=bamD PE=3 SV=1 (A0A0A8F481_NEIME)
Cluster of Outer-membrane lipoprotein LolB OS=Neisseria meningitidis alpha153 GN=lob PE=3 SV=1 (C6SCD3_NEIME)
Cluster of Pyruvate kinase OS=Neisseria meningitidis serogroup B (strain NZ-05/33) GN=pyk PE=3 SV=1 (A0A0E0TMU0_NEIMN)
Cluster of Serine hydroxymethyltransferase OS=Neisseria meningitidis LNP21362 GN=glyA PE=3 SV=1 (A0A0A8F7J0_NEIME)

Accession Number	Alternate ID	Total Spectrum Count			
		MW (kDa)	WT	NORM.VOL	NORM-Empty
Q51147_NEIME [83]	porB	34	362	1445	1225
A0A125WFE9_NEIME [8]	mpm	26	936	433	485
Q84029_NEIME [37]	fbA	74	31	360	301
Q30375_NEIME [74]	gna1870	23	35	110	82
Q50988_NEIGO [10]	tpa	34	32	283	383
C6S5X1_NEIML [29]	iga1	203	20	417	498
QJ0P11_NEIME [196]	porA	42	69	301	310
Q53136_NEIME [26]	tbpA	69	44	110	82
A1KSP9_NEIMF [10]	NMC0490	31	39	76	93
A0A0AFA088_NEIME [10]	tmf	43	90	192	248
A0A0A8F236_NEIME [10]	n875_01600	85	90	94	86
A0A0E3SU61_NEIMH [30]	tpbB	72	4	61	51
P95359_NEIGO [12]	omp85	88	126	88	86
Q51228_NEIME [8]	opc	28	4	17	19
PLQ_NEIMH [3]	pilQ	83	172	93	131
A0A2A3LV89_NEIME [5]	CNQ34_01945	31	64	71	81
C6S9N9_NEIML [35]	pilS2	12	252	22	94
A0A0A8FAA4_NEIME [10]	rplB	30	169	9	28
A0A0A8F533_NEIME (+2)	N875_00970	14	61	31	48
A0A125WBX2_NEIME [2]	NMEN2795_1510	14	134	64	47
C6S154_NEIME [14]	haeR	29	53	66	44
N82XV6_NEIME [10]	LbpA	105	1	45	29
BDY09_NEIME [8]	hmbR	87	6	30	37
N875_00975		13	30	18	34
C6S797_NEIML [4]	NMO_1093	11	82	46	59
A0A0A8F509_NEIME [3]	adnA	13	120	13	32
E3D230_NEIMT [9]	adnA	37	0	84	97
C6S808_NEIME [7]	cysK3	5	12	91	121
A0A125WD37_NEIME [17]	nmC	24	40	34	38
A0A0A8F829_NEIME [14]	N875_08420	33	23	38	41
A0A0A8F8Q5_NEIME [4]	pgk	44	36	41	56
A0A0A8F8R2_NEIME [10]	nmB1969	74	114	63	63
N875_04030	E0NCQ2_NEIME [5]	50	85	71	90
HMPREF0602_2194	lbpB	83	3	39	47
A0A125WC21_NEIME [11]	FRPA_NEIMC [13]	122	0	107	102
FRPA_NEIMC [13]	adnA	5	4	19	19
A0A0A8F3W7_NEIME [17]	iga_2	114	8	55	71
A0A0A8F2T0_NEIME [4]	N875_03055	29	63	18	32
ME795_NEIME [10]	pta	72	12	35	92
A0A0A8F489_NEIME [14]	pilF	28	35	20	15
A0A125WD38_NEIME [7]	N875_08420	33	23	38	41
J8VLH5_NEIME [14]	NMEN3004_1151	255	1	15	32
NM0ZT_1470	NM0ZT_1470	15	0	13	27
C6S6E8_NEIME [16]	yaP	19	26	28	32
ATL0A7_NEIME [14]	nlpB	37	106	38	34
A0A2A3LQ61_NEIME [2]	gna1	12	16	0	13
A0A0A8F8D9_NEIME [15]	groL	57	34	24	43
XSEP09_NEIME [5]	cpa1946	31	163	13	24
A0A125WDK9_NEIME [8]	tpa2	53	15	52	38
LPTD_NEIME [8]	MEK21_NEIME [4]	88	28	17	24
ERS514410_00524	ERS514410_00524	17	0	16	18
N875_05160	N875_05160	19	19	22	21
A0A0A8F479_NEIME [11]	pic1	159	11	97	109
C6S2Q6_NEIME [17]	ppsA	87	14	10	29
J8WMZ9_NEIME [5]	E0N7Y8_NEIME [2]	11	76	14	22
HMPREF0602_0618	A0A0A8F8F2_NEIME [16]	20	5	30	58
N875_01760	A0A0A8F7M1_NEIME [9]	20	5	30	58
C6S6M6_NEIML [5]	sucC	41	8	25	27
BVE1E1_NEIME [5]	NMEN53004_0126	55	1	11	26
M0579_00675	NM0579_00675	5	17	23	6
A0A1V3SQW6_NEIME [8]	mriB	58	39	21	32
A0A0E0TN23_NEIMN [5]	nmrD	23	58	1	26
N875_04030	A0A0A8F4M9_NEIME [3]	38	1	28	39
A0A125WC25_NEIME [5]	ilvC	36	91	2	21
XSFBA2_NEIME [4]	pod1	44	83	7	21
A0A0Y6U395_NEIME [3]	sucD	31	4	18	23
B7S1T2_NEIME [26]	surA	46	163	2	12
A0A0E0TMP2_NEIMN [17]	rpsL	14	35	7	22
QJ0X95_NEIME [5]	gapA-2	36	79	8	10
TRYP_PG		24	10	9	9
ESMV09_NEIMH [7]	dnaK	67	21	18	37
A0A0A8FA15_NEIME [7]	N875_06710	36	2	32	27
A0A0A8F7F8_NEIME [7]	N875_09885	29	44	4	12
ESM2J4_NEIMF [5]	trf	30	6	19	36
A0A0A8F859_NEIME [20]	N875_03035	19	4	50	33
ROU033_NEIME [8]	kt	72	4	13	24
B7S1T2_NEIME [26]	porB	38	12	24	22
A0A1V0G5E8_NEIME [3]	trxB	18	2	15	23
K2C1_HUMAN	KRT1	66	11	10	13
K2Z2_HUMAN	KRT2	66	5	10	4
A0A1R1N790_NEIME (+2)	ALJL54_02205	9	11	8	13
C6S699_NEIML [3]	NMO_0724	24	5	5	10
A0A0A8F536_NEIME [3]	N875_08420	77	0	9	6
A0A121T6Z3_NEIME [5]	atpA	55	76	2	8
A0A0A8F703_NEIME [7]	eno	46	163	2	12
A0A0A8F716_NEIME [10]	N875_09320	48	78	6	15
A0A125WFY0_NEIME [5]	dsbA3	26	47	0	9
A0A0H5Q426_NEIM [10]	trxB	105	1	45	29
A0A0A8FA86_NEIME [20]	nplL	13	47	11	15
QJ0YB8_NEIMB [3]	NMB1657	22	11	1	8
C6S7H6_NEIML [5]	aceE	100	8	14	37
A0A0A8F509_NEIME (+16)	rplN	13	45	1	0
QJ0MP9_NEIME [7]	QJ0MP9_NEIME [7]	42	34	9	16
A0A0Y6WE99_NEIME [3]	rgIA	24	11	2	19
C6S184_NEIM	NMO_1967	19	14	22	19
A0A0E0TQY5_NEIMN [4]	lpe	48	10	26	51
A0A0A8FA45_NEIME (+5)	tg	18	12	9	7
A0A0A8FAC5_NEIME (+16)	rplD	23	24	0	11
A0A0A8F8N4_NEIME (+15)	rpmA	10	5	1	11
A0A0Y5C793_NEIME [3]	nlpD	42	33	17	22
A0A0A8FA41_NEIME [7]	N875_03675	46	2	14	26
A0A0A8FAU2_NEIME [7]	secE	16	12	8	16
A0A125WCH0_NEIME [8]	nmC	17	4	37	44
A0A0A8FAA9_NEIME (+18)	rpmC	7	6	0	8
NMEN2795_0410	NMEN2795_0410	62	0	6	25
A0A0A8F481_NEIME	band	34	12	8	8
C6SCD3_NEIME [5]	lob	25	14	0	5
A0A0E0TMU0_NEIMN [8]	pyk	52	3	6	6
A0A0A8F7J0_NEIME [10]	glyA	45	3	11	5

Cluster of Superoxide dismutase [Cu-Zn] OS=Neisseria meningitidis GN=sodC PE=3 SV=1 (A0A0Y5YZ13_NEIME)
Cluster of Uncharacterized protein OS=Neisseria meningitidis alpha275 GN=NMW_2164 PE=4 SV=1 (CS6SM5_NEIME)
Keratin, type I cytoskeletal 10 OS=Homo sapiens GN=KRT10 PE=1 SV=6
Uncharacterized protein OS=Neisseria meningitidis serogroup B (strain NZ-0533) GN=NMNBZ0533_1106 PE=4 SV=1
Cluster of 5-methyltetrahydropteroylglutamate--homocysteine methyltransferase OS=Neisseria meningitidis (strain alpha14) GN=metE PE=3 SV=1 (CS6SL6_NEIML)
Cluster of 50S ribosomal protein L25 OS=Neisseria meningitidis GN=rlsA PE=3 SV=1 (E0N4S3_NEIME)
Cluster of Acetate kinase OS=Neisseria meningitidis alpha275 GN=fackA PE=3 SV=1 (CS6SHZ_NEIME)
Cluster of Adenylate kinase OS=Neisseria meningitidis LNP21362 GN=radK PE=3 SV=1 (A0A0AF325_NEIME)
Cluster of Bacterioferritin OS=Neisseria meningitidis LNP21362 GN=N875_00990 PE=3 SV=1 (A0A0AF157_NEIME)
Cluster of Polyribonucleotase OS=Neisseria meningitidis ATCC 13091 GN=ppn PE=3 SV=1 (E0NA11_NEIME)
Cluster of Putative peroxiredoxin/glutathione family protein OS=Neisseria meningitidis alpha275 GN=NMW_2538 PE=4 SV=1 (CS6KNK_NEIME)
Cluster of Tetrahydropteroyl repeat protein OS=Neisseria meningitidis serogroup B (strain NZ-0533) GN=NMNBZ0533_0916 PE=4 SV=1 (A0A0E0TPL8_NEIMN)
Cluster of Thiol disulfide interchange protein DnaD OS=Neisseria meningitidis GN=dnad PE=3 SV=1 (A0A2AAL231_NEIME)
Cluster of 50S ribosomal protein L28 OS=Neisseria meningitidis LNP21362 GN=ppmB PE=3 SV=1 (A0A0AF5E8_NEIME)
Cluster of Membrane protein OS=Neisseria meningitidis LNP21362 GN=N875_00130 PE=4 SV=1 (A0A0AF2K7_NEIME)
Cluster of NADPH-dependent FMN reductase OS=Neisseria meningitidis serogroup B (strain NZ-0533) GN=NMNBZ0533_0530 PE=4 SV=1 (A0A0E0TNZ7_NEIMN)
Cluster of Putrescine-binding periplasmic protein OS=Neisseria meningitidis serogroup B PE=3 SV=1 (A0A0H5QV38_NEIME)
Cluster of Sec-independent protein translocase protein YaaA OS=Neisseria meningitidis LNP21362 GN=yaaA PE=3 SV=1 (A0A0AF8390_NEIME)
Cluster of Transaldolase OS=Neisseria meningitidis LNP21362 GN=tal PE=3 SV=1 (A0A0AF6R0_NEIME)
Elongation factor P OS=Neisseria meningitidis LNP21362 GN=efp PE=3 SV=1
Transferin binding protein B (Fragment) OS=Neisseria meningitidis PE=4 SV=1
Cluster of Uncharacterized protein OS=Neisseria meningitidis LNP21362 GN=N875_02780 PE=4 SV=1 (A0A0AF2N5_NEIME)
Cluster of Outer membrane protein OS=Neisseria meningitidis (strain alpha14) GN=mspA PE=4 SV=1 (CS6SS9_NEIML)
Cluster of Lamrinin subunit alpha Lamrinin A chain OS=Neisseria meningitidis alpha522 GN=alpA PE=4 SV=1 (ME754_NEIME)
Cluster of Putative lipoprotein OS=Neisseria meningitidis NM2795 GN=NMENZ795_0105 PE=4 SV=1 (A0A125WFK9_NEIME)
Cluster of Transglycosylase OS=Neisseria meningitidis GN=sat PE=4 SV=1 (A0A0Y5YPU4_NEIME)
30S ribosomal protein S19 OS=Neisseria meningitidis LNP21362 GN=psl PE=3 SV=1
Cluster of 50S ribosomal protein N2795 GN=ppB PE=3 SV=1 (A0A125WAS9_NEIME)
Cluster of 50S ribosomal protein L11 OS=Neisseria meningitidis GN=ppK PE=3 SV=1 (A0A121QVP4_NEIME)
Cluster of 50S ribosomal protein L9 OS=Neisseria meningitidis NM2795 GN=ppL PE=3 SV=1 (A0A125WD48_NEIME)
Cluster of DNA-directed RNA polymerase subunit beta' OS=Neisseria meningitidis GN=ppoc PE=3 SV=1 (A0A1V3SMAK8_NEIME)
Cluster of Glutamine synthetase OS=Neisseria meningitidis LNP21362 GN=gsaA PE=3 SV=1 (A0A0AF6U2_NEIME)
Acyl carrier protein OS=Neisseria meningitidis LNP21362 GN=acpP PE=3 SV=1
Cluster of 3-oxoacyl-CoA dehydrogenase OS=Neisseria meningitidis GN=leuB PE=3 SV=1 (A0A1V3SR58_NEIME)
Cluster of 3-oxoacyl-CoA dehydrogenase OS=Neisseria meningitidis (strain alpha14) GN=leuB PE=4 SV=1 (CS6SS0_NEIML)
Cluster of 50S ribosomal protein L13 OS=Neisseria meningitidis serogroup B / serotype 15 (strain H4476) GN=ppM PE=3 SV=1 (E0M2P5_NEIMH)
Cluster of Aminomethyltransferase OS=Neisseria meningitidis LNP21362 GN=gcvT PE=3 SV=1 (A0A0AF4N6_NEIME)
Cluster of Bifunctional purine biosynthesis protein PurH OS=Neisseria meningitidis serogroup B GN=ppuH PE=3 SV=1 (A0A0H5QDY1_NEIME)
Cluster of Isocitrate--RNA ligase OS=Neisseria / serotype 4A (strain Z2491) GN=hisA PE=3 SV=1 (SV1_NEIME)
Cluster of Outer membrane ferric siderophore receptor OS=Neisseria meningitidis (strain alpha14) GN=NMO_0332 PE=3 SV=1 (CS6S79_NEIML)
Cluster of Phosphoribosylformylglycinamide cyclo-ligase OS=Neisseria meningitidis LNP21362 GN=ppuM PE=3 SV=1 (A0A0AF228_NEIME)
Cluster of Ribonucleoside-diphosphate reductase OS=Neisseria meningitidis GN=A6J54_08400 PE=3 SV=1 (A0A1V0GT79_NEIME)
Cluster of Uncharacterized protein OS=Neisseria meningitidis C (strain 033442) GN=NMOC_548 PE=4 SV=1 (ADMTE0_NEIME)
Cluster of UPP2010 protein N875_09670 OS=Neisseria meningitidis LNP21362 GN=N875_09670 PE=3 SV=1 (A0A0AF7B8_NEIME)
Peptide deformylase OS=Neisseria meningitidis LNP21362 GN=def PE=3 SV=1
Transferin binding protein B (Fragment) OS=Neisseria meningitidis PE=4 SV=1
Uncharacterized protein OS=Neisseria meningitidis ATCC 13091 GN=HMPREF0602_1882 PE=4 SV=1
Cluster of Uncharacterized protein OS=Neisseria meningitidis GN=ERS514729_00095 PE=4 SV=1 (A0A0Y6E3F5_NEIME)
Cluster of Uncharacterized protein OS=Neisseria meningitidis GN=A6J54_08695 PE=4 SV=1 (A0A1V0GB02_NEIME)
Cluster of Uncharacterized protein OS=Neisseria meningitidis GN=A6J27_06600 PE=4 SV=1 (A0A1B1ZDL3_NEIME)
Peptidase M23 OS=Neisseria meningitidis LNP21362 GN=N875_09105 PE=4 SV=1
30S ribosomal protein S13 OS=Neisseria meningitidis GN=ppM PE=3 SV=1
Alkyl hydroxide reductase AhrpD OS=Neisseria meningitidis serogroup B (strain NZ-0533) GN=NMNBZ0533_1569 PE=3 SV=1
Cluster of 30S ribosomal protein S11 OS=Neisseria meningitidis LNP21362 GN=ppsk PE=3 SV=1 (A0A0AF4X2_NEIME)
Cluster of Adenylosuccinate synthetase OS=Neisseria meningitidis GN=ppuA PE=3 SV=1 (A0A1V0GX3_NEIME)
Cluster of Aminotransferase OS=Neisseria meningitidis NM2795 GN=tyb2 PE=3 SV=1 (A0A125WEM1_NEIME)
Cluster of Argininosuccinate lyase OS=Neisseria meningitidis NM2795 GN=argG PE=3 SV=1 (A0A125WAV9_NEIME)
Cluster of Branched-chain amino acid aminotransferase OS=Neisseria meningitidis 93004 GN=ivE PE=4 SV=1 (J0T6C2_NEIME)
Cluster of Chemotaxis protein CheY OS=Neisseria meningitidis LNP21362 GN=N875_04075 PE=4 SV=1 (A0A0AF8Q07_NEIME)
Cluster of Electron transfer flavoprotein FAD-binding domain protein OS=Neisseria meningitidis ATCC 13091 GN=mta PE=4 SV=1 (E0NC51_NEIME)
Cluster of Glutamate--cysteine ligase OS=Neisseria meningitidis LNP21362 GN=gluA PE=3 SV=1 (A0A1Z5WAV9_NEIME)
Cluster of Leucine--RNA ligase OS=Neisseria meningitidis NM2795 GN=leuS PE=3 SV=1 (A0A125WBF2_NEIME)
Cluster of Malate phosphorylase OS=Neisseria meningitidis (strain alpha14) GN=mapA PE=4 SV=1 (CS6SR9_NEIML)
Cluster of Oligopeptidase A OS=Neisseria meningitidis NM2795 GN=pp1 PE=3 SV=1 (A0A125WFR0_NEIME)
Cluster of Putative large exoprotein involved in heme utilization or adhesion of Shiga/HecA/FlaA family OS=Neisseria meningitidis serogroup B PE=4 SV=1 (A0A0H5QAE5_NEIML)
Cluster of Putative peptidase OS=Neisseria meningitidis alpha275 GN=nlpD1 PE=4 SV=1 (CS6H79_NEIME)
Cluster of Ribonuclease E OS=Neisseria meningitidis serogroup A / serotype 4A (strain Z2491) GN=rnaE PE=3 SV=1 (A0A0U1RG59_NEIMA)
Cluster of Valine--RNA ligase OS=Neisseria meningitidis LNP21362 GN=ivB PE=3 SV=1 (A0A0AF691_NEIME)
Lactoferrin-binding protein B OS=Neisseria meningitidis GN=B1A92_10920 PE=4 SV=1
Putative cytochrome oxidase subunit OS=Neisseria meningitidis serogroup A / serotype 4A (strain Z2491) GN=NMA1979 PE=4 SV=1
30S ribosomal protein S17 OS=Neisseria meningitidis LNP21362 GN=ppaD PE=3 SV=1
30S ribosomal protein OS=Neisseria meningitidis LNP21362 GN=ppE PE=3 SV=1
50S ribosomal protein L22 OS=Neisseria meningitidis LNP21362 GN=ppV PE=3 SV=1
50S ribosomal protein L36 OS=Neisseria meningitidis serogroup B (strain NZ-0533) GN=ppuL PE=3 SV=1
Cell division protein Pz2 OS=Neisseria meningitidis LNP21362 GN=fbz PE=3 SV=1
Cluster of 2,3,4,5-tetrahydroxydine-2,6-dicarboxylate N-succinyltransferase OS=Neisseria meningitidis GN=dapD PE=3 SV=1 (A0A0Y6TGX3_NEIME)
Cluster of 23S rRNA (guanosine-2'-O-)-methyltransferase RlmB OS=Neisseria meningitidis ATCC 13091 GN=rmbB PE=3 SV=1 (E0NK92_NEIME)
Cluster of 50S ribosomal protein L10 OS=Neisseria meningitidis LNP21362 GN=ppJ PE=3 SV=1 (A0A0AF458_NEIME)
Cluster of Arylsulfatase OS=Neisseria meningitidis ATCC 13091 GN=HMPREF0602_0390 PE=4 SV=1 (E0NTB0_NEIME)
Cluster of Aspartokinase OS=Neisseria meningitidis ATCC 13091 GN=lysC PE=3 SV=1 (E0NTV4_NEIME)
Cluster of Carbamoyl-phosphate synthase large chain OS=Neisseria meningitidis serogroup B (strain MC58) GN=carB PE=3 SV=1 (CARB_NEIMB)
Cluster of Cysteine desulfurase IcsB OS=Neisseria meningitidis serogroup B (strain MC58) GN=icsB PE=3 SV=1 (ICSB_NEIMB)
Cluster of Efflux pump membrane transporter OS=Neisseria meningitidis LNP21362 GN=N875_09290 PE=3 SV=1 (A0A0AF772_NEIME)
Cluster of Glucokinase OS=Neisseria meningitidis M0579 GN=glk_2 PE=3 SV=1 (A0A0G4BV0C_NEIME)
Cluster of Hemolysin secretion/activation protein, ShB family OS=Neisseria meningitidis serogroup B (strain NZ-0533) GN=NMNBZ0533_1736 PE=4 SV=1 (A0A0E0TRV2_NEIMN)
Cluster of Nitrogen regulatory protein P-2 OS=Neisseria meningitidis (strain H4476) GN=nmrB_0187 PE=3 SV=1 (E0NMV1_NEIMH)
Cluster of Outer membrane protein assembly factor BamE OS=Neisseria meningitidis (strain alpha14) GN=bamE PE=3 SV=1 (CS6995_NEIML)
Cluster of Peptidase OS=Neisseria meningitidis LNP21362 GN=N875_07960 PE=4 SV=1 (A0A0AF6D7_NEIME)
Cluster of Pilus assembly protein PilP OS=Neisseria meningitidis LNP21362 GN=N875_04710 PE=4 SV=1 (A0A0AF925_NEIME)
Cluster of Protein translocase subunit SecY OS=Neisseria meningitidis GN=sacA_1 PE=3 SV=1 (A0A0Y5FM4_NEIME)
Cluster of Putative iron-sulfur cluster insertion protein ErpA OS=Neisseria meningitidis serogroup B (strain NZ-0533) GN=erpa PE=3 SV=1 (A0A0E0TQ68_NEIMN)
Cluster of Single-stranded DNA-binding protein OS=Neisseria meningitidis serogroup B (strain NZ-0533) GN=ssb PE=3 SV=1 (A0A0E0TS6_NEIMN)
Cluster of Site-determining factor OS=Neisseria meningitidis LNP21362 GN=N875_07400 PE=3 SV=1 (A0A0AF4X0_NEIME)
Hemagglutinin/hemolysin family protein OS=Neisseria meningitidis GN=haB_1 PE=4 SV=1
Lactoferrin-binding protein B OS=Neisseria meningitidis 93004 GN=lbpB PE=4 SV=1
Transferin-binding protein 1 (Fragment) OS=Neisseria meningitidis LNP21362 GN=hisA PE=3 SV=1
Universal stress protein OS=Neisseria meningitidis LNP21362 GN=N875_10440 PE=3 SV=1
1-(5-phosphoribosyl)-5-[5-(phosphoribosylamino)methylideneamino]imidazole-4-carboxamide isomerase OS=Neisseria meningitidis LNP21362 GN=hisA PE=3 SV=1
1-acyl-glycerol-3-phosphate acyltransferase OS=Neisseria meningitidis serogroup B PE=3 SV=1
30S ribosomal protein S14 OS=Neisseria meningitidis LNP21362 GN=ppM PE=3 SV=1
50S ribosomal protein L25 OS=Neisseria meningitidis M0579 GN=ppM PE=3 SV=1
50S ribosomal protein L36 OS=Neisseria meningitidis LNP21362 GN=ppM PE=3 SV=1
Acetylactate synthase OS=Neisseria meningitidis LNP21362 GN=N875_10110 PE=4 SV=1
Argininosuccinate synthetase OS=Neisseria meningitidis serogroup A / serotype 4A (strain Z2491) GN=argG PE=3 SV=1
Cluster of 3-hydroxyacid dehydrogenase OS=Neisseria meningitidis NM2795 GN=mmB PE=4 SV=1 (A0A125WC35_NEIME)
Cluster of 3-oxoacyl-CoA carrier-protein synthase 2 OS=Neisseria meningitidis LNP21362 GN=N875_07730 PE=3 SV=1 (A0A0AF8FP2_NEIME)
Cluster of 50S ribosomal protein L17 OS=Neisseria meningitidis 93004 GN=ppQ PE=3 SV=1 (J8WTP2_NEIME)
Cluster of 50S ribosomal protein L20 OS=Neisseria meningitidis alpha153 GN=ppL PE=3 SV=1 (CS6AQ2_NEIME)
Cluster of Antibiotic biosynthesis monooxygenase OS=Neisseria meningitidis LNP21362 GN=N875_10115 PE=4 SV=1 (A0A0AFBM2_NEIME)
Cluster of Beta-phosphoglucomutase OS=Neisseria meningitidis serogroup B / serotype 15 (strain H4476) GN=ppmB PE=4 SV=1 (E0M2X5_NEIMH)
Cluster of D-alanyl-D-alanine carboxypeptidase-D-alanyl-D-alanine-endopeptidase OS=Neisseria meningitidis 81658 GN=dacB PE=4 SV=1 (R0TXR8_NEIME)
Cluster of Delta-aminolevulinic acid dehydratase OS=Neisseria meningitidis serogroup A / serotype 4A (strain Z2491) GN=NMA1011 PE=3 SV=1 (A0A0U1IRB2_NEIMA)
Cluster of DNA gyrase subunit A OS=Neisseria meningitidis GN=gyrA PE=3 SV=1 (A0A076U6M4_NEIME)
Cluster of dTDP-4-dehydrothiamine 3,5-epimerase OS=Neisseria meningitidis GN=rbc PE=3 SV=1 (Q51155_NEIME)
Cluster of FeaB (Fragment) OS=Neisseria meningitidis GN=feaB PE=4 SV=1 (REVERSE_W6AQ02_NEIME-DECOY)

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A0A0AF325_NEIME [6] adk 23 2 14 19 7
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CS6SS9_NEIML nspA 18 0 3 11 5
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A0A0AF5L3_NEIME (+18) rpsL 14 8 1 6 5
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A0A125WD48_NEIME [6] rplI 16 43 0 15 5
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A0A0AF6U2_NEIME (+11) glnA 52 68 11 8 5
A0A0AF569_NEIME (+18) acpP 9 3 7 6 4
leuB 39 2 7 9 4
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A0A1V0GB02_NEIME A6J54_08695 13 6 3 4 3
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A0A1V0GX3_NEIME purA 46 0 4 5 3
tyb2 44 4 5 9 3
A0A125WAV9_NEIME [3] rpsA 40 1 1 2 3
J0T6C2_NEIME [6] iivE 36 3 0 0 3
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CARB_NEIMB [7] carB_NEIMB [7] 117 0 3 16 2
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bame 15 10 8 5 5
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A0A0AF691_NEIME (+12) N875_10440 19 2 12 22 2
A0A0AF4F1_NEIME (+2) hisA 26 0 4 1 1
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A0A0G4BXC1_NEIME (+7) glnA 44 13 0 10 1
rmpJ 5 13 0 0 1
N875_10110 18 0 4 3 1
REVERSE_A0Y1NEIM-DECOY (+1) acpP 49 1 0 0 1
rmmB 30 0 10 8 1
N875_07730 43 2 6 10 1
J8WTP2_NEIME [7] rplQ 14 19 0 0 1
CS6AQ2_NEIME [4] rIT 9 11 0 0 1
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gyrA 102 0 0 6 1
rbc 38 0 1 6 1
feaA 34 0 0 0 1

Cluster of FHbp (Fragment) OS=Neisseria meningitidis GN=fHbp PE=4 SV=1 (REVERSE_W6AQX7_NEIME-DECOY)
Cluster of Fumarate hydratase class II OS=Neisseria meningitidis LNP21362 GN=aspA PE=3 SV=1 (A0A0A8F7U0_NEIME)
Cluster of Glucose-6-phosphate 1-dehydrogenase OS=Neisseria meningitidis GN=zwf PE=3 SV=1 (A0A0Y6G459_NEIME)
Cluster of IgA-specific serine endopeptidase autotransporter OS=Neisseria meningitidis 81858 GN=iga PE=4 SV=1 (REVERSE_ROWOC4_NEIME-DECOY)
Cluster of Na(+)-translocating NADH-quinone reductase subunit C OS=Neisseria meningitidis LNP21362 GN=nqrC PE=3 SV=1 (A0A0A8F8T4_NEIME)
Cluster of Outer-membrane lipoprotein carrier protein OS=Neisseria meningitidis GN=ioiA PE=3 SV=1 (XSEF49_NEIME)
Cluster of Periplasmic protein OS=Neisseria meningitidis GN=ERS514729_00530 PE=4 SV=1 (A0A0Y5STW7_NEIME)
Cluster of Phosphoribosylglycinamide formyltransferase OS=Neisseria meningitidis LNP21362 GN=N875_10555 PE=4 SV=1 (A0A0A8F7P6_NEIME)
Cluster of Prolyl oligopeptidase family protein OS=Neisseria meningitidis 81858 GN=NMB1858_0911 PE=4 SV=1 (R0LU4U3_NEIME)
Cluster of Putative phospholipase A1 OS=Neisseria meningitidis M0579 GN=M0579_00529 PE=4 SV=1 (A0A0G4BWG3_NEIME)
Cluster of Succinate dehydrogenase flavoprotein subunit OS=Neisseria meningitidis serogroup C / serotype 2a (strain ATCC 700532 / DSM 15464 / FAM18) GN=sdhA PE=3 SV=1 (A1KTL8_NEIMF)
Cluster of Threonine synthase OS=Neisseria meningitidis serogroup B (strain NZ-05/33) GN=thrC PE=4 SV=1 (A0A0E0TQR4_NEIMN)
Cluster of Transcription elongation factor GnaA OS=Neisseria meningitidis serogroup B GN=gnaA PE=3 SV=1 (A0A0H5QFJ9_NEIMI)
Cluster of Transcription termination/antitermination protein NusG OS=Neisseria meningitidis GN=nusG PE=3 SV=1 (A0A1V3SNN1_NEIME)
Cluster of Uncharacterized protein OS=Neisseria meningitidis alpha 153 GN=NME_1515 PE=4 SV=1 (REVERSE_C6SDZ7_NEIME-DECOY)
Cluster of Uncharacterized protein OS=Neisseria meningitidis serogroup B PE=4 SV=1 (A0A0H5QT51_NEIMI)
Lactoferrin-binding protein B OS=Neisseria meningitidis serogroup B (strain NZ-05/33) GN=lfbpB PE=4 SV=1
Nucleoid-associated protein NMBNZ0533_1423 OS=Neisseria meningitidis serogroup B (strain NZ-05/33) GN=NMBNZ0533_1423 PE=3 SV=1
Transcriptional regulator OS=Neisseria meningitidis LNP21362 GN=N875_00875 PE=3 SV=1
Uncharacterized protein OS=Neisseria meningitidis alpha 153 GN=NME_0322 PE=4 SV=1
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REVERSE_ROWOC4_NEIME-DECOY	iga	171	0	0	1	1
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N875_10555	N875_10555	30	34	0	2	1
NMB1858_0911	NMB1858_0911	23	0	1	2	1
M0579_00529	M0579_00529	42	0	0	0	1
A1KTL8_NEIMF [3]	sdhA	65	6	0	4	1
A0A0E0TQR4_NEIMN [2]	thrC	52	0	4	4	1
R0LU4U3_NEIME	gnaA	19	0	4	6	1
A0A0G4BWG3_NEIME [7]	nusG	21	3	3	5	1
A0A1V3SNN1_NEIME [13]	NME_1515	6	0	0	0	1
REVERSE_C6SDZ7_NEIME-DECOY [2]	A0A0H5QT51_NEIMI [6]	24	0	3	5	1
A0A0E0TSR5_NEIMN	lfbpB	78	1	0	3	1
A0A0E0TSH6_NEIMN (+16)	NMBNZ0533_1423	12	0	0	1	1
A0A0A8F318_NEIME (+16)	N875_00875	9	4	2	6	1
C6SAL1_NEIME	NME_0322	6	1	1	1	1
A0A0A8F773_NEIME (+19)	N875_08665	10	0	2	2	1