*OXA-1-like_BBE55408.1_K.p.*

α1    β1

1  10  20  30★  40★

| Name | Sequence |
|------|----------|
| OXA-1-like_BBE55408.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| 1M6K_beta-lactamase_OXA-1 | ..........................STDISTVASPLFEGTEGCFL |
| OXA-1-like_AVK27274.1_P.a. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AWE78522.1_P.a. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLA55054.1_P.a. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QKG33210.1_P.a. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QOC40829.1_K.p. | MKNTIHINFAIFFIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLF61683.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_ANP63432.1_P.a. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QII98525.1_P.a. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AXL78701.1_P.a. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QEL33624.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QCZ41047.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QPO60228.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_ATW95033.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QBB08468.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QFH57730.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QFI25362.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QHC80210.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QMT04262.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QPD73921.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QPO69359.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QPO94996.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QQA69926.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_ASP02997.1_E.spp. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AOF07832.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_APD44928.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AWF45003.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AWX26315.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_BBE66685.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_ARA49869.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_ART03029.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_ART14043.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_ATO08041.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_ATQ97012.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AUD20317.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AUD31186.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AUD36737.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AVF14577.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AWC05483.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QAZ99397.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QIF88052.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLF61156.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLF66737.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLG18431.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AIT04829.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_AKE78741.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
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| OXA-1-like_APM67903.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_APP44482.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_APQ29072.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_APR51020.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_ARM26101.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_ARM26250.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
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| OXA-1-like_ARZ96561.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_ASC31826.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QIF55803.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QIF55813.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_VAX91792.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_VAX91482.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QQL36550.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QPP73193.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QPJ20091.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QPB99042.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QPA13355.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QOU70610.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QOU65181.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QOU39182.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
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| OXA-1-like_QMS74616.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLQ67472.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLI33436.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLI28063.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
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| OXA-1-like_QLI17177.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLI11731.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLC16052.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLC05373.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
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| OXA-1-like_QLB84130.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASUPLFEGTEGCFL |
| OXA-1-like_QLB78789.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QLB73465.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QIR58461.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QHG08101.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QGZ85920.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QGZ80560.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QGT67290.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QFY31451.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QFH43884.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QER51630.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |
| OXA-1-like_QEG84620.1_K.p. | MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL |

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OXA-1-like_QBD88840.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_QBA36467.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_QAZ94541.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_BBE58940.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
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OXA-1-like_AZI07357.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_AZH52602.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_AZG68296.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
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OXA-1-like_AYD37605.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_AXZ62257.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
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OXA-1-like_AVO80920.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_AUX60365.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_ATR77794.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_ATR72263.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_ATR66734.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_ATR61171.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_ATR39902.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_ATR34406.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_ATR28842.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_ATR07462.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
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OXA-1-like_ASC42867.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_QKK74875.1_K.p.    MLAVKIKPFTKPILIMKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_ATM40508.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_ATM68467.1_K.p.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
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OXA-1-like_APV28290.1_K.p.    MLAVKIKPFTKPILIMKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
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OXA-1-like_ANF44129.1_K.p.    MLAVKIKPFTKPILIMKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_AIG86468.1_K.p.    MLAVKIKPFTKPILIMKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_AHI38983.1_K.p.    MLAVKIKPFTKPILIMKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_QCP74266.1_P.a.    ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
OXA-1-like_QDE47052.1_E.spp.  ...............MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFL
                         acc
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OXA-1-like subfamily

*OXA-1-like_BBE55408.1_K.p.*
β2  α2  η1  α3  η2
TTT  ←  llll  llllllll  lllllllllllllll  TT  ★  lll
50  60  70  80  90  100

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OXA-1-like_BBE55408.1_K.p.      LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
1M6K_beta-lactamase_OXA-1       LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
OXA-1-like_AVK27274.1_P.a.      LYDVSTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
OXA-1-like_AWE78522.1_P.a.      LYDVSTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
OXA-1-like_QLA55054.1_P.a.      LYDVSTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
OXA-1-like_QKG33210.1_P.a.      LYDVSTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
OXA-1-like_QOC40829.1_K.p.      LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
OXA-1-like_QLF61683.1_K.p.      LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
OXA-1-like_ANP63432.1_P.a.      LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
OXA-1-like_QII98525.1_P.a.      LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
OXA-1-like_AXL78701.1_P.a.      LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
OXA-1-like_QEL33624.1_E.spp.    LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
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OXA-1-like_QER51630.1_K.p.      LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIDQKTIFKWDKTPKGMEIW
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| OXA-1-like_QCX07927.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_QBH11436.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_QBF99632.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_QBD88840.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_QBA36467.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_QAZ94541.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_BBE58940.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_BBE06733.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AZI07357.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AZH52602.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AZG68296.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AZG62714.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AYD37605.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AXZ62257.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AXZ56762.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AXZ10749.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AWR57732.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AWQ55663.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AWQ55646.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
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| OXA-1-like_AWC00236.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
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| OXA-1-like_AUX60365.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
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| OXA-1-like_ATR72263.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_ATR66734.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_ATR61171.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
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| OXA-1-like_ATR34406.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
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| OXA-1-like_ATQ27236.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
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| OXA-1-like_ATM40508.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_ATM68467.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_ASC37513.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AQU64164.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AQU64317.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
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| OXA-1-like_APV22515.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_ANF44129.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AIG86468.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_AHI38983.1_K.p. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_QCP74266.1_P.a. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIW |
| OXA-1-like_QDE47052.1_E.spp. | LYDASTNAEIAQFNKAKCATQMAPDSTFKIALSLMAFDAEIMNQKTIFKWDKTPKGMEIW |

*acc*

OXA-1-like subfamily

*OXA-1-like_BBE55408.1_K.p.*

α4  α5  α6  α7

110  120  130  140  150  160

| Label | Sequence |
|---|---|
| OXA-1-like_BBE55408.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| 1M6K_beta-lactamase_OXA-1 | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AVK27274.1_P.a. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWE78522.1_P.a. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLA55054.1_P.a. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QKG33210.1_P.a. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QOC40829.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLF61683.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ANP63432.1_P.a. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QII98525.1_P.a. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AXL78701.1_P.a. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QEL33624.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QCZ41047.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QPO60228.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATW95033.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QBB08468.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QFH57730.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QFI25362.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QHC80210.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QMT04262.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QPD73921.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QPO69359.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QPO94996.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QQA69926.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ASP02997.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AOF07832.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_APD44928.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWF45003.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWX26315.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_BBE66685.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ARA49869.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ART03029.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ART14043.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATO08041.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATQ97012.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AUD20317.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AUD31186.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AUD36737.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AVF14577.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWC05483.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QAZ99397.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QIF88052.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLF61156.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLF66737.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLG18431.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AIT04829.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AKE78741.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ANK19483.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AOF02731.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_APB53840.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_APM47234.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_APM56693.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_APM62285.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_APM67903.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_APP44482.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_APQ29072.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_APR51020.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ARM26101.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ARM26250.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ART14424.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ARX22868.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ARZ96561.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ASC31826.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QIF55803.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QIF55813.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_VAX91792.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_VAX91482.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QQL36550.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QPP73193.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QPJ20091.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QPB99042.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QPA13355.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QOU70610.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QOU65181.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QOU39182.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QOT97178.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QOD80635.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QOD80603.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QNB32204.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QMS74616.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLQ67472.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLI33436.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLI28063.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLI22583.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLI17177.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLI11731.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLC16052.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLC05373.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLC00088.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLB94808.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLB89468.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLB84130.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLB78789.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QLB73465.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QIR58461.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QHG08101.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QGZ85920.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QGZ80560.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QGT67290.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QFY31451.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QFH43884.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QER51630.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QEG84620.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |

| Label | Sequence |
|---|---|
| OXA-1-like_QDZ35954.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QCX07927.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QBH11436.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QBF99632.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QBD88840.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QBA36467.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QAZ94541.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_BBE58940.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_BBE06733.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AZI07357.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AZH52602.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AZG68296.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AZG62714.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AYD37605.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AXZ62257.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AXZ56762.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AXZ10749.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWR57732.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWQ55663.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWQ55646.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWQ55645.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWO49662.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWO44383.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWO39120.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWC00236.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AWA71813.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AVO80920.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AUX60365.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATR77794.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATR72263.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATR66734.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATR61171.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATR39902.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATR34406.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATR28842.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATR07462.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATQ91613.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATQ27236.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ASC42867.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QKK74875.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATM40508.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ATM68467.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ASC37513.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AQU64164.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AQU64317.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_APV28290.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_APV22515.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_ANF44129.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AIG86468.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_AHI38983.1_K.p. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QCP74266.1_P.a. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |
| OXA-1-like_QDE47052.1_E.spp. | NSNHTPKTWMQFSVVWVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLE |

acc

α8    α9    β3    β4

*OXA-1-like_BBE55408.1_K.p.*

170    180    190    200    210    220

TT    TT

| Sequence | Alignment |
|---|---|
| OXA-1-like_BBE55408.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| 1M6K_beta-lactamase_OXA-1 | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AVK27274.1_P.a. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLENSTKLYGKTGAGFTANR |
| OXA-1-like_AWE78522.1_P.a. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLENSTKLYGKTGAGFTANR |
| OXA-1-like_QLA55054.1_P.a. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLENSTKLYGKTGAGFTANR |
| OXA-1-like_QKG33210.1_P.a. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QOC40829.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLF61683.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIENENTIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ANP63432.1_P.a. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QII98525.1_P.a. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AXL78701.1_P.a. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QEL33624.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QCZ41047.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QPO60228.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATW95033.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QBB08468.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QFH57730.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QFI25362.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QHC80210.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QMT04262.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QPD73921.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QPO69359.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QPO94996.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QQA69926.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ASP02997.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AOF07832.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_APD44928.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWF45003.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWX26315.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_BBE66685.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ARA49869.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ART03029.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ART14043.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATO08041.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATQ97012.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AUD20317.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AUD31186.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AUD36737.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AVF14577.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWC05483.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QAZ99397.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QIF88052.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLF61156.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLF66737.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLG18431.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AIT04829.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AKE78741.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ANK19483.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AOF02731.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_APB53840.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_APM47234.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_APM56693.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_APM62285.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_APM67903.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_APP44482.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_APQ29072.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_APR51020.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ARM26101.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ARM26250.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ART14424.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ARX22868.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ARZ96561.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ASC31826.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QIF55803.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QIF55813.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_VAX91792.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_VAX91482.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QQL36550.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QPP73193.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QPJ20091.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QPB99042.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QPA13355.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QOU70610.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QOU65181.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QOU39182.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QOT97178.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QOD80635.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QOD80603.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QNB32204.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QMS74616.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLQ67472.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLI33436.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLI28063.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLI22583.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLI17177.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLI11731.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLC16052.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLC05373.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLC00088.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLB94808.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLB89468.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLB84130.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLB78789.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QLB73465.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QIR58461.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QHG08101.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QGZ85920.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QGZ80560.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QGT67290.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QFY31451.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QFH43884.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QER51630.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QEG84620.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |

| Label | Sequence |
|---|---|
| OXA-1-like_QDZ35954.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QCX07927.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QBH11436.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QBF99632.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QBD88840.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QBA36467.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QAZ94541.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_BBE58940.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_BBE06733.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AZI07357.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AZH52602.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AZG68296.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AZG62714.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AYD37605.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AXZ62257.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AXZ56762.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AXZ10749.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWR57732.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWQ55663.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWQ55646.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWQ55645.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWO49662.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWO44383.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWO39120.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWC00236.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AWA71813.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AVO80920.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AUX60365.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATR77794.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATR72263.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATR66734.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATR61171.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATR39902.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATR34406.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATR28842.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATR07462.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATQ91613.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATQ27236.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ASC42867.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QKK74875.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATM40508.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ATM68467.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ASC37513.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AQU64164.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AQU64317.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_APV28290.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_APV22515.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_ANF44129.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AIG86468.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_AHI38983.1_K.p. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QCP74266.1_P.a. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |
| OXA-1-like_QDE47052.1_E.spp. | SSLKISPEEQIQFLRKIINHNLPVKNSAIEN.TIENMYLQDLDNSTKLYGKTGAGFTANR |

acc

OXA-1-like subfamily

OXA-1-like_BBE55408.1_K.p.

β5 → TT → β6 → α10

230    240★   250    ★ 260    270

| Sequence | Alignment |
|---|---|
| OXA-1-like_BBE55408.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| 1M6K_beta-lactamase_OXA-1 | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AVK27274.1_P.a. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWE78522.1_P.a. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLA55054.1_P.a. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QKG33210.1_P.a. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QOC40829.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLF61683.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ANP63432.1_P.a. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QII98525.1_P.a. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AXL78701.1_P.a. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QEL33624.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QCZ41047.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QPO60228.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATW95033.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QBB08468.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QFH57730.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QFI25362.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QHC80210.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QMT04262.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QPD73921.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QPO69359.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QPO94996.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QQA69926.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ASP02997.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AOF07832.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_APD44928.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWF45003.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWX26315.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_BBE66685.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ARA49869.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ART03029.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ART14043.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATO08041.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATQ97012.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AUD20317.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AUD31186.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AUD36737.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AVF14577.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWC05483.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QAZ99397.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QIF88052.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLF61156.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLF66737.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLG18431.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AIT04829.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AKE78741.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ANK19483.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AOF02731.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_APB53840.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_APM47234.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_APM56693.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_APM62285.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_APM67903.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_APP44482.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_APQ29072.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_APR51020.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ARM26101.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ARM26250.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ART14424.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ARX22868.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ARZ96561.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ASC31826.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QIF55803.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QIF55813.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_VAX91792.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_VAX91482.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QQL36550.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QPP73193.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QPJ20091.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QPB99042.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QPA13355.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QOU70610.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QOU65181.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QOU39182.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QOT97178.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QOD80635.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QOD80603.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QNB32204.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QMS74616.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLQ67472.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLI33436.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLI28063.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLI22583.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLI17177.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLI11731.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLC16052.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLC05373.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLC00088.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLB94808.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLB89468.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLB84130.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLB78789.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QLB73465.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QIR58461.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QHG08101.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QGZ85920.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QGZ80560.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QGT67290.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QFY31451.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QFH43884.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QER51630.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QEG84620.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |

| | |
|---|---|
| OXA-1-like_QDZ35954.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QCX07927.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QBH11436.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QBF99632.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QBD88840.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QBA36467.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QAZ94541.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_BBE58940.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_BBE06733.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AZI07357.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AZH52602.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AZG68296.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AZG62714.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AYD37605.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AXZ62257.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AXZ56762.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AXZ10749.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWR57732.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWQ55663.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWQ55646.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWQ55645.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWO49662.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWO44383.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWO39120.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWC00236.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AWA71813.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AVO80920.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AUX60365.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATR77794.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATR72263.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATR66734.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATR61171.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATR39902.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATR34406.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATR28842.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATR07462.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATQ91613.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATQ27236.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ASC42867.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QKK74875.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATM40508.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ATM68467.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ASC37513.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AQU64164.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AQU64317.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_APV28290.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_APV22515.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_ANF44129.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AIG86468.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_AHI38983.1_K.p. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QCP74266.1_P.a. | TLQNGWFEGFIISKSGHKYVFVSALTGSLGSNLTSSIKAKKNAITILNTLNL |
| OXA-1-like_QDE47052.1_E.spp. | TLQNGWFEGFIISKSGHKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL |

*acc*

1K38_1_Beta-lactamase

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1K38_1_Beta-lactamase          . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_VAV79013.1_E.spp.   . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_AXX42030.1_A.b.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_APV34059.1_K.p.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_ARG86543.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_AVK22308.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_SZD41030.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_ASA15328.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_ASA17138.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_ASD12027.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_AYZ57805.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_QPV55945.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_AXS74056.1_P.a.     MSRAPVRDRRISVRKSRPACCCPHCCSTRRRSTRVCRAFDPPVCGGVLLQRCVLGGLVR
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1K38_1_Beta-lactamase

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1K38_1_Beta-lactamase          . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_VAV79013.1_E.spp.   . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_AXX42030.1_A.b.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_APV34059.1_K.p.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_ARG86543.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_AVK22308.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_SZD41030.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_ASA15328.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_ASA17138.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_ASD12027.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_AYZ57805.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_QPV55945.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . .
OXA-2-like_AXS74056.1_P.a.     CAKCAASGRRCSSGKSRRALGSWSRMHRIRLRHSTYQQRKHLGAPGGWIHGGCSSTLLPE
                          acc
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1K38_1_Beta-lactamase

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1K38_1_Beta-lactamase          . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_VAV79013.1_E.spp.   . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_AXX42030.1_A.b.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_APV34059.1_K.p.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_ARG86543.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_AVK22308.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_SZD41030.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_ASA15328.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_ASA17138.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_ASD12027.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_AYZ57805.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_QPV55945.1_P.a.     . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . . MAIRIFAIL
OXA-2-like_AXS74056.1_P.a.     TVVRGAALVPNNSFKPNLLRYTKAMAERACHGFGSTTQVGLTQQLGIKEKLMAIRIFAIL
                          acc
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β1    η1    α1              β2           β3    α2
1K38_1_Beta-lactamase

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                                       1         10        20        30      ★ 40
1K38_1_Beta-lactamase          . . . . . . . . . . QEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_VAV79013.1_E.spp.   FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_AXX42030.1_A.b.     FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_APV34059.1_K.p.     FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_ARG86543.1_P.a.     FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_AVK22308.1_P.a.     FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_SZD41030.1_P.a.     FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_ASA15328.1_P.a.     FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_ASA17138.1_P.a.     FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_ASD12027.1_P.a.     FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_AYZ57805.1_P.a.     FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_QPV55945.1_P.a.     FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
OXA-2-like_AXS74056.1_P.a.     FSIFSLATFAHAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFDPVRSKKRYS
                          acc
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η2         α3              TT                          α4          α5
1K38_1_Beta-lactamase

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                         50        60        70        80        90      ★ 100
1K38_1_Beta-lactamase          PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_VAV79013.1_E.spp.   PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_AXX42030.1_A.b.     PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_APV34059.1_K.p.     PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_ARG86543.1_P.a.     PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_AVK22308.1_P.a.     PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_SZD41030.1_P.a.     PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_ASA15328.1_P.a.     PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_ASA17138.1_P.a.     PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_ASD12027.1_P.a.     PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_AYZ57805.1_P.a.     PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_QPV55945.1_P.a.     PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
OXA-2-like_AXS74056.1_P.a.     PASTFKPHTLFALDAGAVRDEFQIFRWDGVNRGFAGHNQDQDLRSAMRNSTVWVYELFA
                          acc
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OXA-2-like subfamily

**1K38_1_Beta-lactamase** (α6 ... α7)

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                              α6                                              α7
                    110       120       130       140       150       160
1K38_1_Beta-lactamase        KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_VAV79013.1_E.spp. KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_AXX42030.1_A.b.   KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_APV34059.1_K.p.   KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_ARG86543.1_P.a.   KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_AVK22308.1_P.a.   KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_SZD41030.1_P.a.   KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_ASA15328.1_P.a.   KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_ASA17138.1_P.a.   KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_ASD12027.1_P.a.   KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_AYZ57805.1_P.a.   KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_QPV55945.1_P.a.   KEIGDDKARRYLKKIDYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
OXA-2-like_AXS74056.1_P.a.   KEIGDDKARRYLKKIDYGNVVPSTSNGDYWIEGSLAISAQEQIAFLRKLYRNELPFRVEH
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acc

**1K38_1_Beta-lactamase** (α8 β4 TT β5 β6 TT β7 η3)

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                    α8        β4   TT   β5          β6    TT   β7        η3
                    170       180       190       200       210       220
1K38_1_Beta-lactamase        QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_VAV79013.1_E.spp. QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_AXX42030.1_A.b.   QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_APV34059.1_K.p.   QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_ARG86543.1_P.a.   QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_AVK22308.1_P.a.   QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_SZD41030.1_P.a.   QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_ASA15328.1_P.a.   QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_ASA17138.1_P.a.   QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_ASD12027.1_P.a.   QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_AYZ57805.1_P.a.   QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_QPV55945.1_P.a.   QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
OXA-2-like_AXS74056.1_P.a.   QRLVKDLMIVEAGRNWILRAKTGYEGRMGWWVGWVEWPTGSVFFALNIDTPNRMDDLFKR
```
acc

**1K38_1_Beta-lactamase** (α9)

```
                    α9
                    230       240       250
1K38_1_Beta-lactamase        EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_VAV79013.1_E.spp. EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_AXX42030.1_A.b.   EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_APV34059.1_K.p.   EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_ARG86543.1_P.a.   EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_AVK22308.1_P.a.   EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_SZD41030.1_P.a.   EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_ASA15328.1_P.a.   EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_ASA17138.1_P.a.   EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_ASD12027.1_P.a.   EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_AYZ57805.1_P.a.   EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_QPV55945.1_P.a.   EAIVRAILRSIEALPPNPAVNSDAAR
OXA-2-like_AXS74056.1_P.a.   EAIVRAILRSIEALPPNPAVNSDAAR
```
acc

OXA-5-like subfamily

OXA-5-like_AXN26200.1_P.a.

β1  η1  α1  β2  β3  α2

```
OXA-5-like_AXN26200.1_P.a.
OXA-5-like_QQV63761.1_P.a.
OXA-5-like_AWR71556.1_E.spp.
OXA-5-like_AXM02354.1_E.spp.
1FOF_BETA
                          acc
```

OXA-5-like_AXN26200.1_P.a.

β4  η2  α3  TT  η3  α4

```
OXA-5-like_AXN26200.1_P.a.
OXA-5-like_QQV63761.1_P.a.
OXA-5-like_AWR71556.1_E.spp.
OXA-5-like_AXM02354.1_E.spp.
1FOF_BETA
                          acc
```

OXA-5-like_AXN26200.1_P.a.

α5  β5  α6

```
OXA-5-like_AXN26200.1_P.a.
OXA-5-like_QQV63761.1_P.a.
OXA-5-like_AWR71556.1_E.spp.
OXA-5-like_AXM02354.1_E.spp.
1FOF_BETA
                          acc
```

OXA-5-like_AXN26200.1_P.a.

α7  β6  β7  β8  TT  β9

```
OXA-5-like_AXN26200.1_P.a.
OXA-5-like_QQV63761.1_P.a.
OXA-5-like_AWR71556.1_E.spp.
OXA-5-like_AXM02354.1_E.spp.
1FOF_BETA
                          acc
```

OXA-5-like_AXN26200.1_P.a.

η4  α8

```
OXA-5-like_AXN26200.1_P.a.
OXA-5-like_QQV63761.1_P.a.
OXA-5-like_AWR71556.1_E.spp.
OXA-5-like_AXM02354.1_E.spp.
1FOF_BETA
                          acc
```

OXA-9-like subfamily

α1     α2        β1

TT                         TT

1    10      20      30      40      50

| | |
|---|---|
| OXA-9-like_AHE47441.1_K.p. | .........MKDTLMKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AWX31296.1_K.p. | .........MKDTLMKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_QPP43211.1_P.a. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_QNU96141.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_QFP85352.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_QEI39179.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AXZ29898.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AXZ16965.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AWC00377.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AVS24357.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AVO98723.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AVF15061.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AUU98532.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AUH99537.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AUA54226.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_ATO26466.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_ASC42960.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_ASC31839.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_ASC14689.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_ASB15563.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_ARX47977.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_ARM26190.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_ARA50435.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_ARA44412.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_APQ29078.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_APM79139.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_APM73509.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AOF02440.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AIG86711.1_K.p. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_QDE47049.1_E.spp. | ..............MKKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_AXL27566.1_K.p. | MHLLSAQRSNEGYLDEKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| OXA-9-like_QCX08049.1_K.p. | MHLLSAQRSNEGYLDEKILLLHMLVFVSATLPISSVASDEVETLKCTIIADAITGNTLYE |
| 6NHU_Beta-lactamase | ...........................SNAQQQPQAFECTLVTSIETGAIINQ |

acc
1

η1   α3              η2    α4

TT                      TT

60      70      80      90    100    110

| | |
|---|---|
| OXA-9-like_AHE47441.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AWX31296.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_QPP43211.1_P.a. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_QNU96141.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_QFP85352.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_QEI39179.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AXZ29898.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AXZ16965.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AWC00377.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AVS24357.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AVO98723.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AVF15061.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AUU98532.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AUH99537.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AUA54226.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_ATO26466.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_ASC42960.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_ASC31839.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_ASC14689.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_ASB15563.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_ARX47977.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_ARM26190.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_ARA50435.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_ARA44412.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_APQ29078.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_APM79139.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_APM73509.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AOF02440.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AIG86711.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_QDE47049.1_E.spp. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_AXL27566.1_K.p. | TGECARRVSPCSSFKLPLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| OXA-9-like_QCX08049.1_K.p. | TGECARRVSPFSSFKLTLAIMGFDSGILQSPKSPTWELKPEYNPSPRDRTYKQVYPALWQ |
| 6NHU_Beta-lactamase | QGACDQRVAPASTFKVPLALIGFDAGILQDGKTPAWDWKPGTEARASDR..KTVDPTIWE |

acc
1

α5     α6     α7

Positions: 120   130   140   150   160   170

| Sequence | Alignment |
|---|---|
| OXA-9-like_AHE47441.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AWX31296.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_QPP43211.1_P.a. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_QNU96141.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_QFP85352.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_QEI39179.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AXZ29898.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AXZ16965.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AWC00377.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AVS24357.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AVO98723.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AVF15061.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AUU98532.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AUH99537.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AUA54226.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_ATO26466.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_ASC42960.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_ASC31839.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_ASC14689.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_ASB15563.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_ARX47977.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_ARM26190.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_ARA50435.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_ARA44412.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_APQ29078.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_APM79139.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_APM73509.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AOF02440.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AIG86711.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_QDE47049.1_E.spp. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_AXL27566.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| OXA-9-like_QCX08049.1_K.p. | SDSVVWFSQQLTSRLGVDRFTEYVKKFFYGNDVSGDSGKHNGLTQSWLMSSLTISPKEQ |
| 6NHU_Beta-lactamase | QDSVLWYSREITRRLGPEKFAAYVKRLGYGNDVSGEPGKNNGLTHSWLGASLTISPVEQ |

acc

α8     α9     β2     β3    TT    β4

Positions: 180   190   200   210   220   230

| Sequence | Alignment |
|---|---|
| OXA-9-like_AHE47441.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AWX31296.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_QPP43211.1_P.a. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_QNU96141.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_QFP85352.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_QEI39179.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AXZ29898.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AXZ16965.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AWC00377.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AVS24357.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AVO98723.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AVF15061.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AUU98532.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AUH99537.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AUA54226.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_ATO26466.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_ASC42960.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_ASC31839.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_ASC14689.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_ASB15563.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_ARX47977.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_ARM26190.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_ARA50435.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_ARA44412.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_APQ29078.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_APM79139.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_APM73509.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AOF02440.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AIG86711.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_QDE47049.1_E.spp. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYRAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_AXL27566.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYQAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| OXA-9-like_QCX08049.1_K.p. | IQFLLRFVAHKLPVSEAAYDMAYATIPQYRAAEGWAVHGKSGSGWLRDNNGKINESRPQG |
| 6NHU_Beta-lactamase | VGFLRRLLGGNLPFSRDAQAKTRAIMPVFDAPESWAVHGKTGCGYMRDEKGNPDRNRPFG |

acc

OXA-9-like subfamily

OXA-10-like subfamily

OXA-10-like_QQV64094.I_P.a.

β1   η1   α1          β2        β3      α2

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1          10          20          30          40          50          60
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OXA-10-like_QQV64094.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QIP22830.1_E.spp.  MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AZN49265.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AZN03320.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QFI25219.1_E.spp.  MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QQW19467.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QLF44358.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QBC10196.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AZN61489.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AZN16107.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AZN09071.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AYR16166.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AYQ84126.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AXN28660.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QGQ07367.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QGQ07350.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QGQ00675.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AYZ81322.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AYZ81321.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QCC69644.1_P.a.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QRC68437.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QQL36827.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QQL36820.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QOD80660.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QIV52637.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QFH43880.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_QDF35708.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_BBR80767.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AZI07531.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AZH95660.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AXS22507.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AXR54228.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AXR54222.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_ASC26206.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_ASC20336.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_ASC14736.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_ASC14731.1_K.p.    MKTFAAYVIIACLSSTALAGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
1K6R_Beta-lactamase            ................MGSITENTSWNKEFSAEAVNGVFVLCKSSSKSCATNDLARAS
OXA-10-like_AUA92950.1_P.a.    MKTFAAYVITACLSSTALASSITENTFWNKEFSAEAVNGVFVLCKSSSKSCATNNLARAS
OXA-10-like_AUA86857.1_P.a.    MKTFAAYVITACLSSTALASSITENTFWNKEFSAEAVNGVFVLCKSSSKSCATNNLARAS
OXA-10-like_QQW12184.1_P.a.    MKTFAAYVITACLSSTALASSITENTFWNKEFSAEAVNGVFVLCKSSSKSCATNNLARAS
OXA-10-like_QEK44114.1_P.a.    MKTFAAYVITACLSSTALASSITENTFWNKEFSAEAVNGVFVLCKSSSKSCATNNLARAS
OXA-10-like_QPP37890.1_P.a.    MKTFAAYVITACLSSTALASSITENTFWNKEFSAEAVNGVFVLCKSSSKSCATNNLARAS

acc

OXA-10-like_QQV64094.I_P.a.

η2   α3        TT        η3      α4

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70          80          90         100         110         120
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OXA-10-like_QQV64094.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_QIP22830.1_E.spp.  KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_AZN49265.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAFPVF
OXA-10-like_AZN03320.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_QFI25219.1_E.spp.  KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_QQW19467.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_QLF44358.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_QBC10196.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_AZN61489.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_AZN16107.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
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OXA-10-like_AYR16166.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_AYQ84126.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_AXN28660.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
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OXA-10-like_AYZ81321.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_QCC69644.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
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OXA-10-like_BBR80767.1_K.p.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_AZI07531.1_K.p.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_AZH95660.1_K.p.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_AXS22507.1_K.p.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_AXR54228.1_K.p.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_AXR54222.1_K.p.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_ASC26206.1_K.p.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_ASC20336.1_K.p.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_ASC14736.1_K.p.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_ASC14731.1_K.p.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
1K6R_Beta-lactamase            KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLTLRGAIQVSAVPVF
OXA-10-like_AUA92950.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLSLRGAIQVSAVPVF
OXA-10-like_AUA86857.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLSLRGAIQVSAVPVF
OXA-10-like_QQW12184.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLSLRGAIQVSAVPVF
OXA-10-like_QEK44114.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLSLRGAIQVSAVPVF
OXA-10-like_QPP37890.1_P.a.    KEYLPASTFKIPNAIIGLETGVIKNEHQVFKWDGKPRAMKQWERDLSLRGAIQVSAVPVF

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OXA-10-like_QQV64094.1_P.a.

α5　　α6　　　　　α7　　　α8
130　　140　　150　　160　　170　　180

OXA-10-like_QQV64094.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QIP22830.1_E.spp.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AZN49265.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AZN03320.1_P.a.　QQIAREVGEVGMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QFI25219.1_E.spp.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QQW19467.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QLF44358.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QBC10196.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AZN61489.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AZN16107.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AZN09071.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AYR16166.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AYQ84126.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AXN28660.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QGQ07367.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QGQ07350.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QGQ00675.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AYZ81322.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AYZ81321.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QCC69644.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QRC68437.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QQL36827.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QQL36820.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QOD80660.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QIV52637.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QFH43880.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_QDF35708.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_BBR80767.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AZI07531.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AZH95660.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AXS22507.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AXR54228.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AXR54222.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_ASC26206.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_ASC20336.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_ASC14736.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_ASC14731.1_K.p.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
1K6R_Beta-lactamase　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLYLNKLSA
OXA-10-like_AUA92950.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLFLNKLSA
OXA-10-like_AUA86857.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLFLNKLSA
OXA-10-like_QQW12184.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLFLNKLSA
OXA-10-like_QEK44114.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLFLNKLSA
OXA-10-like_QPP37890.1_P.a.　QQIAREVGEVPMQKYLKKFSYGNQNISGGIDKFWLEGQLRISAVNQVEFLESLFLNKLSA

acc

OXA-10-like_QQV64094.1_P.a.

α9　　β4　　β5　　TT　　β6　　β7
190　　200　　210　　220　　230　　240

OXA-10-like_QQV64094.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QIP22830.1_E.spp.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AZN49265.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AZN03320.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QFI25219.1_E.spp.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QQW19467.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QLF44358.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QBC10196.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AZN61489.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AZN16107.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AZN09071.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AYR16166.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AYQ84126.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AXN28660.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QGQ07367.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QGQ07350.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QGQ00675.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AYZ81322.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AYZ81321.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QCC69644.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QRC68437.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QQL36827.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QQL36820.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QOD80660.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QIV52637.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QFH43880.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_QDF35708.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_BBR80767.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AZI07531.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AZH95660.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AXS22507.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AXR54228.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AXR54222.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_ASC26206.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_ASC20336.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_ASC14736.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_ASC14731.1_K.p.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
1K6R_Beta-lactamase　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKETEVYFFAFNMD
OXA-10-like_AUA92950.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKGTEVYFFAFNMD
OXA-10-like_AUA86857.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKGTEVYFFAFNMD
OXA-10-like_QQW12184.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKGTEVYFFAFNMD
OXA-10-like_QEK44114.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKGTEVYFFAFNMD
OXA-10-like_QPP37890.1_P.a.　SKENQLIVKEALVTEAAPEYLVHSKTGTSGVGTESNPGVAWWVGWVEKGTEVYFFAFNMD

acc

η4　　　　　α10

OXA-10-like_QQV64094.1_P.a.　▶　ᵒᵒᵒ　ᵒᵒᵒᵒᵒᵒᵒᵒᵒᵒᵒᵒ
　　　　　　　　　　　　　　　　　250　　　　260

| | |
|---|---|
| OXA-10-like_QQV64094.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QIP22830.1_E.spp. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AZN49265.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AZN03320.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QFI25219.1_E.spp. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QQW19467.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QLF44358.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QBC10196.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AZN61489.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AZN16107.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AZN09071.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AYR16166.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AYQ84126.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AXN28660.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QGQ07367.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QGQ07350.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QGQ00675.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AYZ81322.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AYZ81321.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QCC69644.1_P.a. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QRC68437.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QQL36827.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QQL36820.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QOD80660.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QIV52637.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QFH43880.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_QDF35708.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_BBR80767.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AZI07531.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AZH95660.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AXS22507.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AXR54228.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AXR54222.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_ASC26206.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_ASC20336.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_ASC14736.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_ASC14731.1_K.p. | IDNESKLPLRKSIPTKIMGSEGIIGG |
| 1K6R_Beta-lactamase | IDNESKLPLRKSIPTKIMGSEGIIGG |
| OXA-10-like_AUA92950.1_P.a. | IDNENKLPLRKSIPTKIMASEGIIGG |
| OXA-10-like_AUA86857.1_P.a. | IDNENKLPLRKSIPTKIMASEGIIGG |
| OXA-10-like_QQW12184.1_P.a. | IDNENKLPLRKSIPTKIMASEGIIGG |
| OXA-10-like_QEK44114.1_P.a. | IDNENKLPLRKSIPTKIMASEGIIGG |
| OXA-10-like_QPP37890.1_P.a. | IDNENKLPLRKSIPTKIMASEGIIGG |

acc

6XJ3_Beta-lactamase

β1  η1  α1      β2        β3
1        10      20      30      40

6XJ3_Beta-lactamase         ....................SNAQEGTLERSDWRKFFSEFQAKGTIVVADERQADRAMLVFD
OXA-20-like_ACC55538.1_A.b. MIIRFLALLFSAVVLVSLGHAQEKTHESSNWGKYFSDFNAKGTIVVVDERTNGNSTSVYN
acc

α2        η2          α3           TT        η3        α4
50          60          70      80        90        100

6XJ3_Beta-lactamase         PVRSKKRYSPASTFKIPHTLFALDAGAVRDEFQLFRWDGVNRGFAGHNQDQDLRSAMRNS
OXA-20-like_ACC55538.1_A.b. ESRAQQRYSPASTFKIPHTLFALDAGAVRDEFHVFRWDGAKRSFAGHNQDQNLRSAMRNS
acc

α5          α6                        α7
110        120        130        140        150        160

6XJ3_Beta-lactamase         TVWVYELFAKEIGDDKARRYLKKIYGNADPSTSNGDYWIEGSLAISAQEQIAFLRKLYR
OXA-20-like_ACC55538.1_A.b. TVWVYQLFAKEIGENKARSYLEKLNYGNADPSTKSGDYWIDGNLAISANEQISILKKLYR
acc

α8      β4    TT    β5        β6    TT    β7
170      ★180      190        200      210        220

6XJ3_Beta-lactamase         NELPFRVEHQRLVKDLMIVEAGRNWILRLKTGNEGRMGWWVGWVEWPTGSVFFALNIDTP
OXA-20-like_ACC55538.1_A.b. NELPFRVEHQRLVKDLMIVEAKRDWILRLKTGNDGQMGWWVGWVEWPTGPVFFALNIDTP
acc

η4    α9
230        240        250

6XJ3_Beta-lactamase         NRMDDLFKREAIVRAILRSIEALPPNPAVNSDAAR
OXA-20-like_ACC55538.1_A.b. NRMEDLHKREAIARAILQSVNALPPN.........
acc

α1
ℓℓℓℓℓℓℓℓℓ

OXA-23-like_QQE03160.1_A.b.

1        10        20        30        40

OXA-23-like_QQE03160.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QQE03411.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QQD99609.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QQD99362.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QQD95815.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QQD95566.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QQD93073.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QQD92824.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QQB67579.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QQB65684.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QPO03745.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QPO00239.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QLG80230.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QLA71954.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJP32450.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJH23301.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJH19267.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJH16482.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJH15745.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJH15717.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJH08361.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJH12331.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJH05456.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJH00715.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJG95773.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJG96932.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJG96957.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QJH00689.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AOX88011.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AOX91849.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AOX95708.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_APM50633.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_APM50651.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AQU56756.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AQU56788.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARF92486.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARF92497.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARF95290.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARF96125.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARG05487.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARG10150.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARG12999.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARG19192.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARG19219.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARG28873.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARG28898.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARG30155.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ARG35250.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_ASF49900.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AUG13479.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AVN25815.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AVN25876.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AVN25910.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AWW79127.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AWW83877.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AWW86703.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AYY88859.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_BAP68327.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QAS95332.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
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OXA-23-like_QCP27215.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
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OXA-23-like_QCP43779.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
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OXA-23-like_QDR92294.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
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OXA-23-like_QEE57744.1_A.b. ..................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
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OXA-23-like_AMC14310.1_A.b.  ...................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AMC14301.1_A.b.  ...................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
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OXA-23-like_AGQ10499.1_A.b.  ...................MNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
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OXA-23-like_SBS23848.1_A.b.  MSYLFLSCTELFSIDLVFKMNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_AOP62462.1_A.b.  MSYLFLSCTELFSIDLVFKMNKYFTCYVVASLFLSGCTVQHNLINETPSQIVQGHNQVIH
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OXA-23-like_QCP18237.1_A.b.  ......................MWLLLFFFLSGCTVQHNLINETPSQIVQGHNQVIH
OXA-23-like_QRN19109.1_A.b.  ......................MWLLLFF.LSGCTVQHNLINETPSQIVQGHNQVIH
                        acc
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# OXA-23-like subfamily



Multiple sequence alignment of the OXA-23-like subfamily. Reference sequence OXA-23-like_QQE03160.1_A.b. with secondary structure annotation (β1, β2, η1, η2, α2) and residue numbering 50, 60, 70, 80, 90, 100. Conserved consensus sequence: QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKMLNALIGLENQKTDINEIF

Aligned sequences:

OXA-23-like_QQE03160.1_A.b.
OXA-23-like_QQE03411.1_A.b.
OXA-23-like_QQD99609.1_A.b.
OXA-23-like_QQD99362.1_A.b.
OXA-23-like_QQD95815.1_A.b.
OXA-23-like_QQD95566.1_A.b.
OXA-23-like_QQD93073.1_A.b.
OXA-23-like_QQD92824.1_A.b.
OXA-23-like_QQB67579.1_A.b.
OXA-23-like_QQB65684.1_A.b.
OXA-23-like_QPO03745.1_A.b.
OXA-23-like_QPO00239.1_A.b.
OXA-23-like_QLG80230.1_A.b.
OXA-23-like_QLA71954.1_A.b.
OXA-23-like_QJP32450.1_A.b.
OXA-23-like_QJH23301.1_A.b.
OXA-23-like_QJH19267.1_A.b.
OXA-23-like_QJH16482.1_A.b.
OXA-23-like_QJH15745.1_A.b.
OXA-23-like_QJH15717.1_A.b.
OXA-23-like_QJH08361.1_A.b.
OXA-23-like_QJH12331.1_A.b.
OXA-23-like_QJH05456.1_A.b.
OXA-23-like_QJH00715.1_A.b.
OXA-23-like_QJG95773.1_A.b.
OXA-23-like_QJG96932.1_A.b.
OXA-23-like_QJG96957.1_A.b.
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OXA-23-like_AOX88011.1_A.b.
OXA-23-like_AOX91849.1_A.b.
OXA-23-like_AOX95708.1_A.b.
OXA-23-like_APM50633.1_A.b.
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OXA-23-like_AYY88859.1_A.b.
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OXA-23-like_QAS95332.1_A.b.
OXA-23-like_QAT02983.1_A.b.
OXA-23-like_QBY15339.1_A.b.
OXA-23-like_QCH32733.1_A.b.
OXA-23-like_QCP17255.1_A.b.
OXA-23-like_QCP24705.1_A.b.
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OXA-23-like_QCP40778.1_A.b.
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OXA-23-like_QEE57736.1_A.b.
OXA-23-like_QEE57740.1_A.b.
OXA-23-like_QEE57744.1_A.b.
OXA-23-like_QEY29291.1_A.b.
OXA-23-like_QEY29320.1_A.b.
OXA-23-like_QEY29496.1_A.b.
OXA-23-like_QIX29121.1_A.b.
OXA-23-like_QIX32951.1_A.b.
OXA-23-like_QIX36912.1_A.b.
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OXA-23-like_QJG70485.1_A.b.
OXA-23-like_QJG71265.1_A.b.
OXA-23-like_QJG74213.1_A.b.
OXA-23-like_QJG81833.1_A.b.
OXA-23-like_QJG84680.1_A.b.
OXA-23-like_QJG89564.1_A.b.
OXA-23-like_QJG89584.1_A.b.
OXA-23-like_QJG90447.1_A.b.
OXA-23-like_QJG93290.1_A.b.
OXA-23-like_ADX93021.1_A.b.
OXA-23-like_AXG83856.1_A.b.
OXA-23-like_AML68547.1_A.b.
OXA-23-like_ARG03241.1_A.b.
OXA-23-like_AOX84133.1_A.b.
OXA-23-like_AOX82595.1_A.b.
OXA-23-like_AOX75920.1_A.b.
OXA-23-like_AOX75888.1_A.b.

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OXA-23-like_AOX68171.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AMC14310.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AMC14301.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_ALY00722.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_ALX98871.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
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OXA-23-like_AGQ05635.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_ACJ39972.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_QKK76748.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_QKK76707.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
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OXA-23-like_AYC00505.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_ARG39357.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_ARG23739.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_ARG23713.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_ARG03186.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AEP06450.2_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AML75286.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AML75277.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AML71526.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AML69469.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AML66436.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AML65160.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AML65128.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AMN00602.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AMN00587.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AMN00578.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AMN00569.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_QJF37605.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_QJF33717.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_QJH05102.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_QJH23846.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_QCH35083.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_QAT07029.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_QAS99734.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AVE56824.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AVE44374.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_ATR89604.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_ARG29802.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_APF45746.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_APQ95067.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_APQ87413.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_ATR88802.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_ATR88943.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_SBS23848.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_AOP62462.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
4JF4_Beta-lactamase            QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_QCP18237.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
OXA-23-like_QRN19109.1_A.b.    QYFDEKNTSGVLVIQTDKKINLYGNALSRANTEYVPASTFKILNALIGLENQKTDINEIF
                         acc
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OXA-23-like subfamily — multiple sequence alignment

Reference: OXA-23-like_QQE03160.1_A.b.

Secondary structure elements: η3, α3, α4, α5

Position markers: 110, 120, 130, 140, 150, 160

| Sequence ID | Alignment |
|---|---|
| OXA-23-like_QQE03160.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QQE03411.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QQD99609.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QQD99362.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QQD95815.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QQD95566.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QQD93073.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QQD92824.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QQB67579.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QQB65684.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QPO03745.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QPO00239.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QLG80230.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QLA71954.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJP32450.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJH23301.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJH19267.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJH16482.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJH15745.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJH15717.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJH08361.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJH12331.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJH05456.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJH00715.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG95773.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG96932.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG96957.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJH00689.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AOX88011.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AOX91849.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AOX95708.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_APM50633.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_APM50651.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AQU56756.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AQU56788.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARF92486.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARF92497.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARF95290.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARF96125.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARG05487.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARG10150.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARG12999.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARG19192.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARG19219.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARG28873.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARG28898.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARG30155.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARG35250.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ASF49900.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AUG13479.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AVN25815.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AVN25876.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AVN25910.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AWW79127.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AWW83877.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AWW86703.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AYY88859.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_BAP68327.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QAS95332.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QAT02983.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QBY15339.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QCH32733.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QCP17255.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QCP24705.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QCP27215.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QCP29558.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QCP40778.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QCP43779.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QCP44095.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QCP47140.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QCR88532.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QDR92294.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QDR95367.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QEE57736.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QEE57740.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QEE57744.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QEY29291.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QEY29320.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
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| OXA-23-like_QIX29121.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QIX32951.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QIX36912.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QIX40711.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QIX44509.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG69304.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG70467.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG70485.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG71265.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG74213.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG81833.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG84680.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG89564.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG89584.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG90447.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_QJG93290.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ADX93021.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AXG83856.1_A.b. | KWKGEKRLFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AML68547.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_ARG03241.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AOX84133.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AOX82595.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AOX75920.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |
| OXA-23-like_AOX75888.1_A.b. | KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV |

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OXA-23-like_AOX72443.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
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OXA-23-like_AMC14310.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AMC14301.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_ALY00722.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_ALX98871.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AKA30333.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AGQ10499.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AGQ05635.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_ACJ39972.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_QKK76748.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_QKK76707.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_QKK76393.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_QFX72557.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_QFZ56693.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AYC00513.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AYC00505.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_ARG39357.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_ARG23739.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
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OXA-23-like_ARG03186.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AEP06450.2_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AML75286.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AML75277.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AML71526.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AML69469.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AML66436.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AML65160.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AML65128.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
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OXA-23-like_AMN00578.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
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OXA-23-like_QAT07029.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_QAS99734.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AVE56824.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AVE44374.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_ATR89604.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_ARG29802.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
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OXA-23-like_ATR88943.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_SBS23848.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKVSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_AOP62462.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
4JF4_Beta-lactamase           KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_QCP18237.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
OXA-23-like_QRN19109.1_A.b.   KWKGEKRSFTAWEKDMTLGEAMKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQV
                        acc
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OXA-23-like_QQE03160.1_A.b.

α6　　　　α7　　　β3　　　β4

170　　　180　　　190　　　200　　　210　　　220

| Label | Sequence |
|---|---|
| OXA-23-like_QQE03160.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QQE03411.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QQD99609.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QQD99362.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QQD95815.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QQD95566.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QQD93073.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QQD92824.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QQB67579.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QQB65684.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QPO03745.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QPO00239.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QLG80230.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QLA71954.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJP32450.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJH23301.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJH19267.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJH16482.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJH15745.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJH15717.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJH08361.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJH12331.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJH05456.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJH00715.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG95773.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG96932.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG96957.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_AOX91849.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AOX95708.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_APM50633.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_APM50651.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AQU56756.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AQU56788.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_ARF92497.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_ARF96125.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_ARG10150.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ARG12999.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ARG19192.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_ARG35250.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_AVN25876.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AVN25910.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_AWW83877.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AWW86703.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_QCP17255.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_QCP27215.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_QCR88532.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_QEY29320.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
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| OXA-23-like_QIX29121.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QIX32951.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QIX36912.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QIX40711.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QIX44509.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG69304.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG70467.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG70485.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG71265.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG74213.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG81833.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG84680.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG89564.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG89584.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG90447.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJG93290.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ADX93021.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AXG83856.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AML68547.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ARG03241.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AOX84133.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AOX82595.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AOX75920.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AOX75888.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |

| | |
|---|---|
| OXA-23-like_AOX72443.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AOX68171.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AMC14310.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AMC14301.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ALY00722.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ALX98871.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AKA30333.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AGQ10499.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AGQ05635.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ACJ39972.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QKK76748.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QKK76707.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QKK76393.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QFX72557.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QFZ56693.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AYC00513.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AYC00505.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ARG39357.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ARG23739.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ARG23713.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ARG03186.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AEP06450.2_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AML75286.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AML75277.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AML71526.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AML69469.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AML66436.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AML65160.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AML65128.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AMN00602.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AMN00587.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AMN00578.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AMN00569.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJF37605.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJF33717.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJH05102.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QJH23846.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QCH35083.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QAT07029.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QAS99734.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AVE56824.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AVE44374.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ATR89604.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ARG29802.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_APF45746.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_APQ95067.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_APQ87413.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ATR88802.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_ATR88943.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_SBS23848.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_AOP62462.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| **4JF4_Beta-lactamase** | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QCP18237.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |
| OXA-23-like_QRN19109.1_A.b. | DNFWLVGPLKVTPIQEVEFVSQLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAM |

*acc*

OXA-23-like subfamily



OXA-23-like_QQE03160.1_A.b.

β5  β6  α8

★230  240  250  260★  270

| Sequence | Alignment |
|---|---|
| OXA-23-like_QQE03160.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QQE03411.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QQD99609.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QQD99362.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QQD95815.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QQD95566.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QQD93073.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QQD92824.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QQB67579.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QQB65684.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QPO03745.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QPO00239.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QLG80230.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QLA71954.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJP32450.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH23301.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH19267.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH16482.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH15745.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH15717.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH08361.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH12331.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH05456.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH00715.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG95773.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG96932.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG96957.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH00689.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AOX88011.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AOX91849.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AOX95708.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_APM50633.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_APM50651.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AQU56756.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AQU56788.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARF92486.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARF92497.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARF95290.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARF96125.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG05487.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG10150.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG12999.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG19192.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG19219.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG28873.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG28898.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG30155.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG35250.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ASF49900.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AUG13479.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AVN25815.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AVN25876.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AVN25910.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AWW79127.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AWW83877.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AWW86703.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AYY88859.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_BAP68327.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QAS95332.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QAT02983.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QBY15339.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCH32733.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCP17255.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCP24705.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCP27215.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCP29558.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCP40778.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCP43779.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCP44095.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCP47140.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCR88532.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QDR92294.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QDR95367.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QEE57736.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QEE57740.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QEE57744.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QEY29291.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QEY29320.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QEY29496.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QIX29121.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QIX32951.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QIX36912.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QIX40711.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QIX44509.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG69304.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG70467.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG70485.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG71265.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG74213.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG81833.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG84680.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG89564.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG89584.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG90447.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJG93290.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ADX93021.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AXG83856.1_A.b. | NIKSQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AML68547.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG03241.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AOX84133.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AOX82595.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AOX75920.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AOX75888.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |

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| OXA-23-like_AOX72443.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AOX68171.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AMC14310.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AMC14301.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ALY00722.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ALX98871.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AKA30333.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AGQ10499.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AGQ05635.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ACJ39972.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QKK76748.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QKK76707.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QKK76393.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QFX72557.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QFZ56693.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AYC00513.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AYC00505.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG39357.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG23739.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG23713.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG03186.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AEP06450.2_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AML75286.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AML75277.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AML71526.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AML69469.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AML66436.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AML65160.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AML65128.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AMN00602.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AMN00587.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AMN00578.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AMN00569.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJF37605.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJF33717.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH05102.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QJH23846.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCH35083.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QAT07029.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QAS99734.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AVE56824.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AVE44374.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ATR89604.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ARG29802.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_APF45746.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_APQ95067.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_APQ87413.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ATR88802.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_ATR88943.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_SBS23848.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_AOP62462.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| 4JF4_Beta-lactamase | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QCP18237.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |
| OXA-23-like_QRN19109.1_A.b. | DIKPQVGWLTGWVEQPDGKIVAFALNMEMRSEMPASIRNELLMKSLKQLNII |

*acc*

OXA-24-like subfamily



Protein sequence alignment of OXA-24-like subfamily. Sequences: OXA-24-like_QAB42530.1_A.b., OXA-24-like_QRN23785.1_A.b., OXA-24-like_QRN23770.1_A.b., OXA-24-like_QER77259.1_A.b., OXA-24-like_QBA07845.1_A.b., OXA-24-like_QBA07837.1_A.b., OXA-24-like_AXX58331.1_A.b., OXA-24-like_AXX50759.1_A.b., OXA-24-like_AXX43427.1_A.b., 4WM9_1_Beta-lactamase, OXA-24-like_AWO18633.1_A.b., OXA-24-like_QLB37632.1_A.b.

OXA-24-like subfamily

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                                                      α1            β1          β2
OXA-33-like_AWT34042.1_P.a.                       ooooooooo      ──────►   TTT   ──────►
                                          1       10        20        30★     40★    50         60
                                          |        |         |         |       |     |           |
OXA-33-like_AWT34042.1_P.a.   MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFLLYDVSTNAEIAQFNK
OXA-33-like_AXZ95524.1_P.a.   MKNTIHINFAIFLIIANIIYSSASASTDISTVASPLFEGTEGCFLLYDVSTNAEIAQFNK
1M6K_beta-lactamase           .........................STDISTVASPLFEGTEGCFLLYDASTNAEIAQFNK
                      acc
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       α2              η1          α3              TT               η2        α4
OXA-33-like_AWT34042.1_P.a.   ooo          oo   oooooooo                    ooo    oooooooooo
                                   70        80        90       100       110       120
                                    |         |         |         |         |         |
OXA-33-like_AWT34042.1_P.a.   AKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIWNSNHTPKTWMQFSVV
OXA-33-like_AXZ95524.1_P.a.   AKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIWNSNHTPKTWMQFSVV
1M6K_beta-lactamase           AKCATQMAPDSTFKIALSLMAFDAEIIDQKTIFKWDKTPKGMEIWNSNHTPKTWMQFSVV
                      acc
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           α5               α6                      α7                   α8
OXA-33-like_AWT34042.1_P.a.   oooooooooo   oooooooooooo          ooooooo          oooooooooo
                                    130       140       150       160       170       180
                                     |         |         |         |         |         |
OXA-33-like_AWT34042.1_P.a.   WVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLESSLKISPEEQIQFLR
OXA-33-like_AXZ95524.1_P.a.   WVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLESSLKISPEEQIQFLR
1M6K_beta-lactamase           WVSQEITQKIGLNKIKNYLKDFDYGNQDFSGDKERNNGLTEAWLESSLKISPEEQIQFLR
                      acc
```

```
      ooo              α9       β3      TT      β4      TT       β5        TT
OXA-33-like_AWT34042.1_P.a.      ★   ooooooooooo  ──────►        ──────►         ──────►
                                    190       200       210       220       230       240
                                     |         |         |         |         |         |
OXA-33-like_AWT34042.1_P.a.   KIINHNLPVKNSAIENTIENMYLQDLENSTKLYGKTGAGFTANRTLQNGWFEGFIISKSG
OXA-33-like_AXZ95524.1_P.a.   KIINHNLPVKNSAIENTIENMYLQDLENSTKLYGKTGAGFTANRTLQNGWFEGFIISKSG
1M6K_beta-lactamase           KIINHNLPVKNSAIENTIENMYLQDLDNSTKLYGKTGAGFTANRTLQNGWFEGFIISKSG
                      acc
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```
               β6                         α10
OXA-33-like_AWT34042.1_P.a.   ──────►           oooooooooooooooo
                              ★       250     ★   260       270
                                      |           |         |
OXA-33-like_AWT34042.1_P.a.   HKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL
OXA-33-like_AXZ95524.1_P.a.   HKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL
1M6K_beta-lactamase           HKYVFVSALTGNLGSNLTSSIKAKKNAITILNTLNL
                      acc
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OXA-50-like subfamily

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                                        TTT TTT      η1        α1            β1       β2      α2
                                       1      10   llll     lllllllll     ___>     ___>    lllll
                                       .       .   20        30            40        50        .
OXA-50-like_QQW22444.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QGG00241.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QQW19268.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QQW06309.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QQV61463.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QPP39331.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QNE62162.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QLF44164.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QLF18710.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QEK46239.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QJA41232.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AYZ58090.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QDR20381.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AYW69566.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AYZ45913.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AYK26584.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_ASM88695.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AUA80490.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AUB05120.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AVR86641.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AXS75147.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_ALS13221.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_ALY73377.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_ANA70339.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_APB68390.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_ARI94830.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_ARJ01367.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_ASA19039.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AUA74232.1_P.a.   MRPLLFSALLLL.SGHAQASEWNNSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_QCC69374.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSRAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_ATH06157.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSQAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AZZ14888.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSQAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_VDK95206.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSQAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_VDL11569.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSQAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_VDL54838.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSQAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_VDL56340.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSQAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_VDL57163.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSQAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_ALZ29437.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSQAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AON10100.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSQAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
OXA-50-like_AON16083.1_P.a.   MRPLLFSALLLL.SGHAQASEWNDSQAVDKLFGAAGVKGTFVLYDVQRQRYVGHDRERAE
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OXA-50-like_ALZ28576.1_P.a.  TRFVPASTYKVANSLIGLSTGAVRSADEVLPYGGKPQRFKAWEHDMSLREAIKASNVPVY
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acc
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α5 α6 α7

120 130 140 150 160 170

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OXA-50-like subfamily

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α9

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OXA-50-like_QDR14302.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_ALZ16475.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QKF04533.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QNO22317.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QQW12925.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_AYZ81447.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QKL16662.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QQW00026.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_ALV75159.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_ALY89963.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_ALY96872.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_AMX88486.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_ARC82831.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_ASD14027.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_AYZ75988.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QFZ61919.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QKP90385.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QKR47436.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QKR53224.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QKR59118.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QLF25267.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QPP21637.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QQA85373.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QRG72678.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_AYY40220.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_ALZ28576.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QNQ04301.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_ALZ94285.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_AMU04106.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_AOT37986.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_ASD20357.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_AVZ22774.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_AXC20842.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_AYN84711.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_CRZ33166.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QBF65101.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QEF81401.1_P.a.    GGEADIGKRVELGKASLKALGILP
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OXA-50-like_QGM44151.1_P.a.    GGEADIGKRVELGKASLKALGILP
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OXA-50-like_VDL04843.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QCP80115.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QEK88934.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_ALU47404.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_APC75958.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QPZ59753.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QFZ03142.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_QGG01915.1_P.a.    GGEADIGKRVELGKASLKALGILP
OXA-50-like_AUA86524.1_P.a.    GGEADIGKRVELSKASLKALGILP
OXA-50-like_AUA92615.1_P.a.    GGEADIGKRVELSKASLKALGILP
4JF4_Beta-lactamase            .SEMPASIRNELLMKSLKQLNII.
                          acc
```

**4ZDX_Beta-lactamase**

α1   β1   TT

```
                          1        10        20        30       40        50       60
                          |         .         .         .        .         .        |
4ZDX_Beta-lactamase       MNIKTLLLITSAIFISACSPYIVTANPNHSASKSDEKAEKIKNLFNEVHTTGVLVIQQGQ
OXA-51-like_QLB35675.1_A.b.  MNIKTLLLITSAIFISACSPYIVTANPNHSASKSDEKAEKIKNLFNEVHTTGVLVIQQGQ
OXA-51-like_AUT37227.1_A.b.  MNIKTLLLITSAIFISACSPYIVTANPNHSASKSDEKGEKIKNLFNEAHTTGVLVIQQGQ
OXA-51-like_QJF39460.1_A.b.  MNIKTLLLITSAIFISACSPYIVTANPNHSASKSDEKAEKIKNLFNEVHTTGVLVIQQGQ
OXA-51-like_QLI40100.1_A.b.  MNIKTLLLITSAIFISACSPYIVTANPNHSASKSDEKAEKIKNLFNEVHTTGVLVIQQGQ
OXA-51-like_ARG36573.1_A.b.  MNIKALLLITSAIFISACSPYIVTANPNHSASKSDEKGEKIKNLFNEAHTTGVLVIQQGQ
OXA-51-like_ASO71619.1_A.b.  MNIKALLLITSAIFISACSPYIVTANPNHSASKSDEKGEKIKNLFNEAHTTGVLVIQQGQ
OXA-51-like_AVP34890.1_A.b.  MNIKALLLITSAIFISACSPYIVTANPNHSASKSDEKGEKIKNLFNEAHTTGVLVIQQGQ
OXA-51-like_QCR89674.1_A.b.  MNIKALLLITSAIFISACSPYIVTANPNHSASKSDEKGEKIKNLFNEAHTTGVLVIQQGQ
OXA-51-like_QJH20957.1_A.b.  MNIKALLLITSAIFISACSPYIVTANPNHSASKSDEKGEKIKNLFNEAHTTGVLVIQQGQ
OXA-51-like_QMS82274.1_A.b.  MNIKALLLITSAIFISACSPYIVTANPNHSASKSDEKGEKIKNLFNEAHTTGVLVIQQGQ
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OXA-51-like_QCP28577.1_A.b.  MNIKALLLITSAIFISACSPYIVSANPNHSASKSDEKAEKIKNLFNEAHTTGVLVIQQGQ
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OXA-51-like subfamily

OXA-51-like subfamily

4ZDX_Beta-lactamase

β2    η1    η2    α2    η3

4ZDX_Beta-lactamase
OXA-51-like_QLB35675.1_A.b.
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OXA-51-like_QJF39460.1_A.b.
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*acc*

OXA-51-like subfamily

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OXA-51-like_ARG25110.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AYC01674.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ATU49292.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ATU52990.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ATU56628.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
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OXA-51-like_AZK41733.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QAS39909.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QAS33473.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
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OXA-51-like_AMC15222.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ANB88975.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AOX69607.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AOX73369.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AOX77482.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AOX81661.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AOX85074.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AOX88942.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AOX92779.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AOX96637.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_APF43961.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_APJ23527.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_APM48724.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_APQ85553.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_APQ93188.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AQU55455.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ARF91818.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ARF95229.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ARG04169.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ARG09260.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ARG14122.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ARG20812.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ARG26815.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
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OXA-51-like_AUG11400.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
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OXA-51-like_AVE90526.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
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OXA-51-like_AWO15631.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AWW77770.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AWW85353.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AXW91373.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AXX48901.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AXX55862.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QBA07478.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QBB77223.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QCH33275.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QCP15970.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
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OXA-51-like_QCP45479.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QDR93660.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QEY27958.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QIX30833.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QIX34674.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QIX38631.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QIX42205.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
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OXA-51-like_QJG69125.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QJG72958.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QJG88231.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QJG91989.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QJG95593.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QJG99403.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QJH07093.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QJH10219.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
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OXA-51-like_QLA73304.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QNV12531.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWVVGPLKITPQQEAQ
OXA-51-like_QAS93855.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWVVGPLKITHQQEAQ
OXA-51-like_QAT01315.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWVVGPLKITPQQEAQ
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OXA-51-like_QEE57116.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ADX03652.1_A.b.   GDAMKASAIQVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QCO82814.1_A.b.   GDAMKASAIQVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AJF81571.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ATP87403.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AKA32028.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
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OXA-51-like_ARG31555.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_ASF77094.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AVN14185.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AVN25165.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_AXB14467.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QLG79253.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QQB68144.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_SQI55469.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QJH03197.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSKEVKRVGYGNADIGTQVDNFWLVGPLKITPQQEAQ
OXA-51-like_QHB90233.1_A.b.   GDAMKASAIPVYQDLARRIGLELMSNEVKRIGYGNADIGTQVDNFWLVGPLKITPQQEAQ
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OXA-51-like subfamily

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OXA-51-like_ALX99780.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
OXA-51-like_AMC15222.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
OXA-51-like_ANB88975.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
OXA-51-like_AOX69607.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
OXA-51-like_AOX73369.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
OXA-51-like_AOX77482.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
OXA-51-like_AOX81661.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
OXA-51-like_AOX85074.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
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OXA-51-like_ARG14122.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
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OXA-51-like_ARG26815.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
OXA-51-like_ATR88137.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
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OXA-51-like_AXX48901.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
OXA-51-like_AXX55862.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
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OXA-51-like_QCH33275.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
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OXA-51-like_QCP19079.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
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OXA-51-like_QEY27958.1_A.b. FAYKLANKTLPFSQKVQDEVQSMLFIEEKNGNKIYAKSGAGWDVNPQVGWLTGWVVQPQG
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OXA-51-like subfamily

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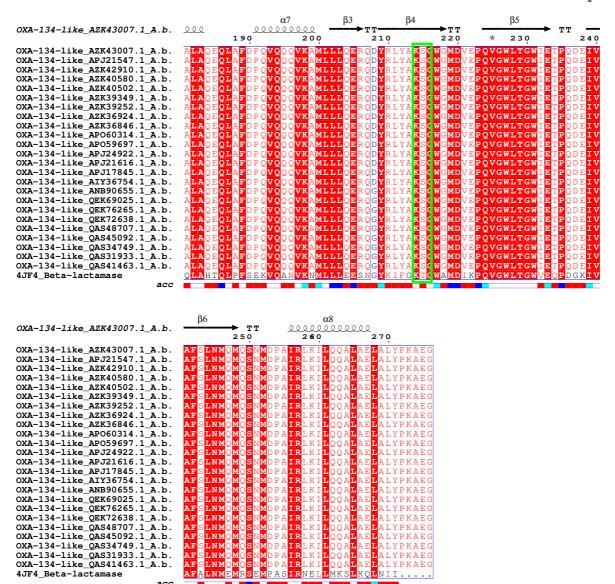
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| OXA-51-like_ATR88137.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_ATU22923.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AUG11400.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AVE45033.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AVE90526.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AVO88307.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AWO15631.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AWW77770.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AWW85353.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AXW91373.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AXX48901.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AXX55862.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QBA07478.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QBB77223.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QCH33275.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QCP15970.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QCP19079.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QCP45479.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QDR93660.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QEY27958.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QIX30833.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QIX34674.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QIX38631.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QIX42205.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QIX46166.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJG69125.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJG72958.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJG88231.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJG91989.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJG95593.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJG99403.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJH07093.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJH10219.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJH14483.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJH17776.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJP31173.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QLA73304.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QNV12531.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QAS93855.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QAT01315.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QAS97806.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QAT04873.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QEE57116.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_ADX03652.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QCO82814.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AJF81571.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_ATP87403.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AKA32028.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_ALJ88113.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_ARG31555.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_ASF77094.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AVN14185.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AVN25165.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_AXB14467.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QLG79253.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QQB68144.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_SQI55469.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QJH03197.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |
| OXA-51-like_QHB90233.1_A.b. | NIVAFSLNLEMKKGISSSVRKEITYRSLEQLGIL |
| OXA-51-like_BBR73296.1_A.b. | NIVAFSLNLEMKKGIPSSVRKEITYKSLEQLGIL |

acc

OXA-58-like subfamily

*4OH0_OXA-58-like_A.b.*

α1   β1

```
         1        10        20        30        40      50        60
```

| Name | Sequence |
|---|---|
| 4OH0_OXA-58-like_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_ANC38777.1_A.b. | .......................MSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_AXW92580.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_AVE88633.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_AVN12783.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_QBY16302.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_QMS84120.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_QJP39652.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_QJF29755.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_QOJ62286.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_QNB01599.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_QBY91612.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_QJF41203.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_QFH47329.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |
| OXA-58-like_AXB17680.1_A.b. | MKLLKILSLVCLSISIGACAEHSMSRAKTSTIPQVNNSIIDQNVQALFNEISADAVFVTY |

acc

*4OH0_OXA-58-like_A.b.*

β2   η1   η2   α2   η3

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              70        80        90       100      110       120
```

| Name | Sequence |
|---|---|
| 4OH0_OXA-58-like_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_ANC38777.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_AXW92580.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_AVE88633.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_AVN12783.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_QBY16302.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_QMS84120.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_QJP39652.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_QJF29755.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_QOJ62286.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_QNB01599.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_QBY91612.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_QJF41203.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_QFH47329.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFFKAWDKD |
| OXA-58-like_AXB17680.1_A.b. | DGQNIKKYGTHLDRAKTAYIPASTFKIANALIGLENHKATSTEIFKWDGKPRFLKAWDKD |

acc

*4OH0_OXA-58-like_A.b.*

α3   α4   α5   α6

```
             130       140       150       160       170       180
```

| Name | Sequence |
|---|---|
| 4OH0_OXA-58-like_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_ANC38777.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_AXW92580.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_AVE88633.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_AVN12783.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_QBY16302.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_QMS84120.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_QJP39652.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_QJF29755.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_QOJ62286.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_QNB01599.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_QBY91612.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_QJF41203.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_QFH47329.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQMGTEVDQFWLKGPLTITPIQ |
| OXA-58-like_AXB17680.1_A.b. | FTLGEAMQASTVPVYQELARRIGPSLMQSELQRIGYGNMQIGTEVDQFWLKGPLTITPIQ |

acc

*4OH0_OXA-58-like_A.b.*

α7   α8   β3   TT   β4   β5

```
             190       200       210       220       230       240
```

| Name | Sequence |
|---|---|
| 4OH0_OXA-58-like_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_ANC38777.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_AXW92580.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_AVE88633.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_AVN12783.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_QBY16302.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_QMS84120.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_QJP39652.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_QJF29755.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_QOJ62286.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_QNB01599.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_QBY91612.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_QJF41203.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_QFH47329.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |
| OXA-58-like_AXB17680.1_A.b. | EVKFVYDLAQGQLPFKPEVQQQVKEMLYVERRGENRLYAKSGWGMAVDPQVGWYVGFVEK |

acc

*4OH0_OXA-58-like_A.b.*

TT   β6   TT   α9

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             250       260       270       280
```

| Name | Sequence |
|---|---|
| 4OH0_OXA-58-like_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_ANC38777.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_AXW92580.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_AVE88633.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_AVN12783.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_QBY16302.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_QMS84120.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_QJP39652.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_QJF29755.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_QOJ62286.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_QNB01599.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_QBY91612.1_A.b. | ADGQVVAFALNMQMKDGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_QJF41203.1_A.b. | ADGQVVAFALNMQMKDGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_QFH47329.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |
| OXA-58-like_AXB17680.1_A.b. | ADGQVVAFALNMQMKAGDDIALRKQLSLDVLDKLGVFHYL |

acc

# OXA-134-like subfamily

OXA-134-like_AZK43007.1_A.b.

```
                                          α1              β1      β2
                                    ΩΩΩΩΩΩΩΩΩΩΩΩΩΩ      →     TT
        1        10        20        30        40        50        60
```

| | |
|---|---|
| OXA-134-like_AZK43007.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_APJ21547.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_AZK42910.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_AZK40580.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_AZK40502.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_AZK39349.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_AZK39252.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_AZK36924.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_AZK36846.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_APO60314.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_APO59697.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_APJ24922.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_APJ21616.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_APJ17845.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_AIY36754.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_ANB90655.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_QEK69025.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_QEK76265.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_QEK72638.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_QAS48707.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_QAS45092.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_QAS34749.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_QAS31933.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| OXA-134-like_QAS41463.1_A.b. | MKTLILLPLLSCLSLTACSLPVSNSSSQITSTQSIQTIAKLFDQAQSSGVLVIQRGPHLQ |
| 4JF4_Beta-lactamase | ..........................QIVQGHN.QVIHQYFDEKNTSGVLVIQTDKKIN |

acc

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                →            η1          η2        α2              TT                  η3          α3
                           ΩΩΩΩΩ     ΩΩΩΩΩΩΩΩΩΩΩΩΩ                                    ΩΩΩ        ΩΩΩΩ
                              70        80        90        100       110       120
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| | |
|---|---|
| OXA-134-like_AZK43007.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_APJ21547.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_AZK42910.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_AZK40580.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_AZK40502.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_AZK39349.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_AZK39252.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_AZK36924.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_AZK36846.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_APO60314.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_APO59697.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_APJ24922.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_APJ21616.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_APJ17845.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_AIY36754.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_ANB90655.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_QEK69025.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_QEK76265.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_QEK72638.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_QAS48707.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_QAS45092.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_QAS34749.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_QAS31933.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| OXA-134-like_QAS41463.1_A.b. | VYGNDLSRAHTEYIPASTFKILNALIGLQHGKATTNEIFKWDGKKRSFAAWEKDMTLGQA |
| 4JF4_Beta-lactamase | LYGNALSRANTEYVPASTFKMLNALIGLENQKTDINEIFKWKGEKRSFTAWEKDMTLGEA |

acc

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                ΩΩΩ        α4              α5                            α6
                         ΩΩΩΩΩΩΩΩΩΩΩΩΩΩ   ΩΩΩΩΩΩΩΩΩΩ                    ΩΩΩΩΩΩΩΩΩΩ
                          130 ★         140       150       160       170       180
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| | |
|---|---|
| OXA-134-like_AZK43007.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_APJ21547.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_AZK42910.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_AZK40580.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_AZK40502.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_AZK39349.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_AZK39252.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_AZK36924.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_AZK36846.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_APO60314.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_APO59697.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_APJ24922.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_APJ21616.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_APJ17845.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_AIY36754.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_ANB90655.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_QEK69025.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_QEK76265.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_QEK72638.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_QAS48707.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_QAS45092.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_QAS34749.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_QAS31933.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| OXA-134-like_QAS41463.1_A.b. | MQASAVPVYQELARRIGLELMQQEVQRIRFGNQQIGQHIDNFWLVGPLKITPEQEVEFAS |
| 4JF4_Beta-lactamase | MKLSAVPVYQELARRIGLDLMQKEVKRIGFGNAEIGQQVDNFWLVGPLKVTPIQEVEFVS |

acc

OXA-134-like_AZK43007.1_A.b.

α7    β3    β4    β5
190    200    210    220    230    240

| | |
|---|---|
| OXA-134-like_AZK43007.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_APJ21547.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_AZK42910.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_AZK40580.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_AZK40502.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_AZK39349.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_AZK39252.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_AZK36924.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_AZK36846.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_APO60314.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_APO59697.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_APJ24922.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_APJ21616.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_APJ17845.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_AIY36754.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQDYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_ANB90655.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQGYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_QEK69025.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQGYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_QEK76265.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQGYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_QEK72638.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQGYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_QAS48707.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQGYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_QAS45092.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQGYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_QAS34749.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQGYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_QAS31933.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQGYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| OXA-134-like_QAS41463.1_A.b. | ALAQEQLAFDPQVQQQVKAMLLLQERQGYRLYAKSGWGMDVEPQVGWLTGWIETPQDEIV |
| 4JF4_Beta-lactamase | QLAHTQLPFSEKVQANVKNMLLLEESNGYKIFGKTGWAMDIKPQVGWLTGWVEQPDGKIV |

acc

β6    α8
250    260    270

| | |
|---|---|
| OXA-134-like_AZK43007.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_APJ21547.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_AZK42910.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_AZK40580.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_AZK40502.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_AZK39349.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_AZK39252.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_AZK36924.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_AZK36846.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_APO60314.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_APO59697.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_APJ24922.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_APJ21616.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_APJ17845.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_AIY36754.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_ANB90655.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_QEK69025.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_QEK76265.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_QEK72638.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_QAS48707.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_QAS45092.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_QAS34749.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_QAS31933.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| OXA-134-like_QAS41463.1_A.b. | AFSLNMQMQSNMDPAIRLKILQQALAELALYPKAEG |
| 4JF4_Beta-lactamase | AFALNMEMRSEMPASIRNELLMKSLKQLNII..... |

acc

OXA-213-like subfamily

Sequence alignment of 4ZDX_1_Beta-lactamase with OXA-213-like_QLF06531.1_A.b. and OXA-213-like_QNB02485.1_A.b.

OXA-48-like_QRC22072.1_K.p.

β1   η1   α1        β2        β3   α2

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                              1        10        20        30        40        50        60
OXA-48-like_QRC22072.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_ASG42064.1_E.spp. MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QRC13711.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_VAY01821.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_VAX91950.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_VAX91959.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QPP62592.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QOU38957.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QOT97232.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QLI33314.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QLI27956.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QLI22474.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QLC16065.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QLC05384.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QLC00099.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QLB94822.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QLB89481.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QLB84144.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QLB78805.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QLB73478.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QIJ23388.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QIJ17785.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QND07059.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QET12150.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QCS89961.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QCT00720.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_QDF35925.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_AZJ02288.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_AZI07598.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_AXZ24690.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_AXS22834.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_AXR54472.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_AWR57773.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
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OXA-48-like_ART03346.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_APQ28988.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_APP73786.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_APP67917.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
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OXA-48-like_APP44504.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_APP38728.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_APM79240.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_APM73588.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_APM67987.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_APM62354.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
OXA-48-like_APM47124.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
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OXA-48-like_QIR58822.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
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OXA-48-like_BBO98227.1_K.p.   MRVLALSAVFLVASIIGMPAVAKEWQENKSWNAHFTEHKSQGVVVLWNENKQQGFTNNLK
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acc

OXA-48-like subfamily

OXA-48-like_QRC22072.1_K.p.

η2   α3                TT           η3      α4
70          80         90      100    110        120

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| OXA-48-like_ASG42064.1_E.spp. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
| OXA-48-like_QRC13711.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
| OXA-48-like_VAY01821.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
| OXA-48-like_VAX91950.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
| OXA-48-like_VAX91959.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
| OXA-48-like_QPP62592.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
| OXA-48-like_QOU38957.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
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| OXA-48-like_QLI22474.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
| OXA-48-like_QLC16065.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
| OXA-48-like_QLC05384.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
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| OXA-48-like_ART03346.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
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| OXA-48-like_QIR58822.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIATWNRDHNLITAMKYSVV |
| 5OE0_1_Beta-lactamase | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIAAWNRDHDLITAMKYSVV |
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| OXA-48-like_QLF61822.1_K.p. | RANQAFLPASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIAAWNRDHDLITAMKYSVV |

acc

6NLW_Beta-lactamase

β1  η1  α1  β2  β3  α2  β4

1  10  20  30  40

6NLW_Beta-lactamase        ..............SNANEWQEKPSWNTHFSEHKAQGVIVLWNENKQQGFTNNLKRANQAFL
OXA-198-like_AYF71930.1_P.a. MLKSTLLAFGLFIALSARAENQAIAKLFLRAGVDGTIVIESLTTGQRLVHNDPRAQQRYP

acc

η2  α3  TT  η3  α4  α5

50  60  70  80  90  100

6NLW_Beta-lactamase        PASTFKIPNSLIALDLGVVKDEHQVFKWDGQTRDIAAWNRDHDLITAMKYSVVPVYQEFA
OXA-198-like_AYF71930.1_P.a. AASTFKVLNTLIALEEGAISGENQIFHWNGTQYSIANWNQDQTLDSAFKVSCVWCYQQIA

acc

α6  α7  β5  α8

110  120  130  140  150  160

6NLW_Beta-lactamase        RQIGQARMSKMLHAFDYGNEDISGNLDSFWLDGGIRISATEQVAFLRKLYHNKLHVSERS
OXA-198-like_AYF71930.1_P.a. LRVGALKYPAYIQQTNYGHLLEPFNGTEFWLDGSLTISAEEQVAFLRQVVERKLPFKASS

acc

α9  β6  β7  β8  β9  η4

170  180  190  200  210  220

6NLW_Beta-lactamase        QRIVKQAMLTEANSDYIIRAKTCYSTRIEPQIGWWVGWVELDDNVWFFAMNMDMPTADGL
OXA-198-like_AYF71930.1_P.a. YDSLKKVMFADENAQYRLYAKTGWATRMTPSVGWYVGYVEAKDDVWLFALNLATRDANDL

acc

α10

230  240

6NLW_Beta-lactamase        GLRQAITKEVLKQEKIIP..
OXA-198-like_AYF71930.1_P.a. PLRTQIAKDALKAIGAFPTK

acc

OXA-48-like subfamily — multiple sequence alignment (residues ~125–180), α5–α8 helices

OXA-48-like_QRC22072.1_K.p.

α9  β4  β5  β6  β7
190   200   210   220   230   240

| Sequence | Alignment |
|---|---|
| OXA-48-like_QRC22072.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_ASG42064.1_E.spp. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QRC13711.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_VAY01821.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_VAX91950.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_VAX91959.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QPP62592.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QOU38957.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QOT97232.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLI33314.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLI27956.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLI22474.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLC16065.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLC05384.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLC00099.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLB94822.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLB89481.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLB84144.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLB78805.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLB73478.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QIJ23388.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QIJ17785.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QND07059.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QET12150.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QCS89961.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QCT00720.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QDF35925.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AZJ02288.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AZI07598.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AXZ24690.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AXS22834.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AXR54472.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AWR57773.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_ART03415.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_ART03346.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APQ28988.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APP73786.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APP67917.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APP56266.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APP50376.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APP44504.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APP38728.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APM79240.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APM73588.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APM67987.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APM62354.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APM47124.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_APG80363.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AOF13070.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QPQ02212.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QPQ02194.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QIJ34285.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QIJ09605.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QIR58822.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTGIEPKIGWWVGWVELDDNVWFFAMNMD |
| 5OE0_1_Beta-lactamase | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_ANE69785.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_ANE71804.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_ANE72730.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AOA98603.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_CEP69754.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_CEP70686.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_CEP72746.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_ATX28768.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_ATX29640.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AVO14768.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AVO15618.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AWD00385.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AWD98100.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AWS85277.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QEG84876.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QIR47572.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QIR53202.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QOQ50577.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_BBO98227.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_BBO99129.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QDE47101.1_E.spp. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTRIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AIG86783.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTSIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_ALH88756.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTSIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AOF02466.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTSIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_ATO26616.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTSIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AZB79861.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTSIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_AZV17592.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTSIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QBG05392.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTSIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QBH46740.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTSIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QBH61269.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTSIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLF61798.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTSIEPKIGWWVGWVELDDNVWFFAMNMD |
| OXA-48-like_QLF61822.1_K.p. | LHVSERSQRIVKQAMLTEANGDYIIRAKTGYSTSIEPKIGWWVGWVELDDNVWFFAMNMD |

acc

OXA-48-like subfamily

OXA-48-like_QRC22072.1_K.p.

η4          α10

250          260

| | |
|---|---|
| OXA-48-like_QRC22072.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_ASG42064.1_E.spp. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QRC13711.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_VAY01821.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_VAX91950.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_VAX91959.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QPP62592.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QOU38957.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QOT97232.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLI33314.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLI27956.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLI22474.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLC16065.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLC05384.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLC00099.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLB94822.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLB89481.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLB84144.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLB78805.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLB73478.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QIJ23388.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QIJ17785.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QND07059.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QET12150.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QCS89961.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QCT00720.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QDF35925.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AZJ02288.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AZI07598.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AXZ24690.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AXS22834.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AXR54472.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AWR57773.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_ART03415.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_ART03346.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APQ28988.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APP73786.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APP67917.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APP56266.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APP50376.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APP44504.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APP38728.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APM79240.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APM73588.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APM67987.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APM62354.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APM47124.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_APG80363.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AOF13070.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QPQ02212.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QPQ02194.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QIJ34285.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QIJ09605.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QIR58822.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| 5OE0_1_Beta-lactamase | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_ANE69785.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_ANE71804.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_ANE72730.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AOA98603.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_CEP69754.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_CEP70686.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_CEP72746.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_ATX28768.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_ATX29640.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AVO14768.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AVO15618.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AWD00385.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AWD98100.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AWS85277.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QEG84876.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QIR47572.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QIR53202.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QOQ50577.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_BBO98227.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_BBO99129.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QDE47101.1_E.spp. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AIG86783.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_ALH88756.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AOF02466.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_ATO26616.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AZB79861.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_AZV17592.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QBG05392.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QBH46740.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QBH61269.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLF61798.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |
| OXA-48-like_QLF61822.1_K.p. | MPTSDGLGLRQAITKEVLKQEKIIP |

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