

OXA-204 Carbapenemase in Clinical Isolate of *Pseudomonas guariconensis*, Tunisia

Appendix

LUT-1 -----MNVILNRRTFLLASAVVSASYSILGTLAGANRDDASFQDRLAKLEQQLNGLGVCAIDTANGAQLGYRANERFAMNSTFKVMLASAFLARSQDEPGILLEERLTYTRADLVVTS	112
GUA-1 -----MKPTFSRRLIQQAMGGSV----MLATPAWASDPVHQRQLQHLEQALGTLGLWALDTGSRRRLSYRSEARAAALCSTFKVVLAAAALQQDQAEPGLLERRIAYTAEQLVVYS	107
FUL-1 ---MGSAHVKPTIRRRQQLLQIGCTS---LLALPSWAADPPLQRQLRERSADRTLGLWALDTGSQQQLAWRAEERFAFCSTFKAILAGAILHRSQAEPGLLERRIAYGAEQLVVYS	112
SOL-1 -----MTPTLSRRLVQGALCST---LLALPTWADDPLRKALQQLEATTGSTIGLWALDTGSGRQLAWRGEERFAFCSTFKAVLAGAVLHRNEAEPGLLARRIAYGAEQLVVYS	107
MOS-1 -----MISTLSRRRVLQGALCSA---LLTLPWAHDPLRQAMQQLEAASGSTIGLWALDTGSRQLAWRGEERFAFCSTFKAVLAGAVLHRNEAEPGLLARRIAYGAEQLVVYS	107
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LUT-1 PVTEKHVSTGMTVAELCAAGIQYSDNTAANVLMKKLGGPEAVTAFAKSIGDTHFR <u>LDRWE</u> TELNSAIPGDP <u>RDTSTP</u> QAMAMSLQRLALGDMIAADKQHQQLQAWLKGNNTGGKRIRAGVP	232
GUA-1 PVTEKHVATGMSVAELCAAQYSDNTAANLLLGVGGPSKLTAYARSIGDPA <u>RLD</u> RYEPALNSALPN <u>DPRDTST</u> PLAMGVTLQRLLTDGLPAAARLRLQDWLKGNNTGDTRIRAGLP	227
FUL-1 PVTQKHVGAGMTVAELCAAQYSDNTAGNLLLEQVGGPAGLTAFA <u>RLGDPV</u> LRLDRYEPELNSALPGDPRDTSSPLAMGATLERLLLT <u>DGLP</u> TAAQQQLQAWLKGNNTGATRIRAGLP	232
SOL-1 PVTEKHVGAGMTVAGLCAAQYSDNTAGNLLLEVGGPSGLTAFTRSISLGDPT <u>FR</u> LDRNEPTLN <u>TA</u> LP <u>GDPRDT</u> TTPLAMGVTLQRLLGDGLAAERQQQLQDWLKGNNTGDARIRAGVP	227
MOS-1 PVTEKHVGPGMTVAELCAAQYQSDNTAGNLLLEVGGPG <u>LTA</u> V <u>RGLDPT</u> F <u>RLDR</u> NEPTLN <u>TA</u> LP <u>GDPRDT</u> TTPLAMGVTLQRLLTDGLPAGAQQLQAWLKGNNTGDARIRAGLP	227
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LUT-1 AGWQVGDK <u>TGT</u> GDYGSANDVAILWP <u>PR</u> APV <u>V</u> LALYSALENQQAEARN <u>D</u> VLA <u>A</u> ARIVAEWV <u>TG</u>	296
GUA-1 EGWVVGDK <u>TG</u> GDYGVANDV <u>G</u> IWPP <u>C</u> QAP <u>W</u> I <u>AI</u> YTR <u>DR</u> DKT <u>S</u> W <u>R</u> DS <u>V</u> IA <u>E</u> AT <u>RV</u> VE <u>AW</u> K <u>V</u>	291
FUL-1 EGWTVGDK <u>TG</u> GDYGVANDV <u>G</u> IVW <u>PP</u> GRAP <u>W</u> I <u>LV</u> V <u>S</u> RGTK <u>AD</u> FW <u>N</u> SET <u>I</u> AA <u>AT</u> RV <u>VE</u> AW <u>KG</u>	296
SOL-1 KDWVVGDK <u>TG</u> GDYGVANDV <u>G</u> AVIWP <u>K</u> GRAP <u>W</u> I <u>VV</u> V <u>S</u> RGTK <u>AD</u> FW <u>N</u> SET <u>I</u> AA <u>AT</u> RV <u>VE</u> AW <u>KG</u>	291
MOS-1 KDWVVGDK <u>TG</u> GDYGVANDV <u>G</u> AVIWP <u>K</u> GRAP <u>W</u> I <u>LV</u> V <u>S</u> RGTK <u>QD</u> FW <u>N</u> SET <u>I</u> AA <u>AT</u> RV <u>VE</u> AW <u>KG</u>	291
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Appendix Figure. Amino acid sequence alignment of Gua-1 from *P. guariconensis* 65411 and *P. guariconensis* (UQM99659.1), Ful from *P. fulva* (MBF8778391.1), Sol from *P. soli* (UXZ44560.1), Mos from *P. mosselii* (WP_186595297.1) and Lut-1 from *P. luteola* (AAU10324.1). Conserved motives among class A beta-lactamases are highlighted in grey. The Omega loop is underlined.

Appendix References

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