

Feature	Reference
Amino acid sequence	
S182T mutation makes the strain resistant to carbapenems	Abbasi et al., 2020
AEVGTTGYG sequence at N-terminal is conserved.	Novovic, et al., 2015
Group I, II and III: each group has two variable regions VRs at the N-	
terminal and two hypervariable regions in between 135–180, and 203–	
240 amino acid positions.	
19 – 160 amino acids are most immunogenic	Tohidinia et al., 2019
Amino acid deletion at 133 and insertions at 140-141 and 154-156	Zhu et al., 2019
complement with imipenem resistance.	
Transcriptional and post transcriptional regulation of CarO	
CarO downregulates at osmotic stress and membrane bound CarO	Hood <i>et al.</i> , 2009
releases into supernatant.	
Sub-MIC concentration of Tetracycline releases CarO into supernatant	Yun et al., 2008
but does not alter transcription.	
Absence of CarO in OMPs despite the presence of gene is found in	Mussi et al., 2005
imipenem resistant A. baumannii.	
In the absence of RNA chaperone Hfq, mRNA levels of CarO rise.	Kuo <i>et al.</i> , 2017
CarO variants and functional aspects	
K178 of CarO interacts with the periplasmic Oxa-23 protein to	Wu et al., 2016
facilitate proin localized toxin inactivation.	
CarOa and CarOb classified on the basis of non-variable N-terminal	Catel-Ferreira et al., 2011
domain (1-131) and two variable domains (132–162 and 200–238)	
CarOa and CarOb are both specific for imipenem uptake. L-orinthin	Mussi et al., 2007; Catel-
competes with imipenem.	Ferreira et al., 2011
CarOa is less specific to imipenem than CarOb	Catel-Ferreira et al., 2011
Insertion disruption of CarO by ISAba 10, ISAba 15, ISAba36,	Mussi et al., 2005; Khorsi
ISAba825, and ISAba125 in Carbapenem resistant strains	et al., 2018; Kim et al.,
	2015; Lee <i>et al.</i> , 2010.

Supplementary Table 1: Genetic basis of CarO function and its role in antibiotic resistance in *A. baumannii*.