SUPPLEMENTARY INFORMATION

Spatial characteristics of the efflux pump MexB determine inhibitor binding

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Supplementary Materials

Supplementary Figure 1. Alignment of E. coli AcrB and P. aeruginosa MexB.

Supplementary Figure 2. Growth of *Escherichia coli* cells in the presence of various concentrations of ABI-PP without antibiotics

Supplementary Figure 3. Growth of *Escherichia coli* cells in the presence of 0.016 μ g/mL of levofloxacin and various concentrations of ABI-PP

Supplementary Figure 4. Growth of *Escherichia coli* cells in the presence of 0.125 μ g/mL of aztreonam and various concentrations of ABI-PP

MexB	1	MSKFFIDRPIFAWVIALVIMLAGGLSILSLPVNQYPAIAPPAIAVQVSYPGASAETVQDT	60
AcrB	1	MPNFFIDRPIFAWVIAIIIMLAGGLAILKLPVAQYPTIAPPAVTISASYPGADAKTVQDT	60
MexB	61	VVQVIEQQMNGIDNLRYISSESNSDGSMTITVTFEQGTDPDIAQVQVQNKLQLATPLLPQ V QVIEQ MNGIDNL Y+SS S+S G++ IT+TFE GTD DIAQVQVQNKLQLA PLLPQ	120
AcrB	61	VTQVIEQNMNGIDNLMYMSSNSDSTGTVQITLTFESGTDADIAQVQVQNKLQLAMPLLPQ	120
MexB	121	EVQRQGIRVTKAVKNFLMVVGVVSTDGSMTKEDLSNYIVSNIQDPLSRTKGVGDFQVFGS EVQ+QG+ V K+ +FLMVVGV++TDG+MT+ED+S+Y+ +N++D +SRT GVGD Q+FGS	180
AcrB	121	EVQQQGVSVEKSSSSFLMVVGVINTDGTMTQEDISDYVAANMKDAISRTSGVGDVQLFGS	180
MexB	181	QYSMRIWLDPAKLNSYQLTPGDVSSAIQAQNVQISSGQLGGLPAVKGQQLNATIIGKTRL OY+MRIW++P +LN +0LTP DV +AI+A0N 0+++G0LGG P VKG00LNA+II +TRL	240
AcrB	181	QYAMRIWMNPNELNKFQLTPVDVITAIKAQNAQVAAGQLGGTPPVKGQQLNASIIAQTRL	240
MexB	241	QTAEQFENILLKVNPDGSQVRLKDVADVGLGGQDYSINAQFNGSPASGIAIKLATGANAL + E+F ILLKVN DGS+V L+DVA + LGG++Y I A+FNG PASG+ IKLATGANAL	300
AcrB	241	TSTEEFGKILLKVNQDGSRVLLRDVAKIELGGENYDIIAEFNGQPASGLGIKLATGANAL	300
MexB	301	DTAKAIRQTIANLEPFMPQGMKVVYPYDTTPVVSASIHEVVKTLGEAILLVFLVMYLFLQ DTA ATR +A +EPF P G+K+VVPYDTTP V STHEVVKTL EAT+LVFLVMYLFLQ	360
AcrB	301	DTAAAIRAELAKMEPFFPSGLKIVYPYDTTPFVKISIHEVVKTLVEAIILVFLVMYLFLQ	360
MexB	361	NFRATLIPTIAVPVVLLGTFGVLAAFGFSINTLTMFGMVLAIGLLVDDAIVVVENVERVM NFRATLIPTIAVPVVLLGTF VLAAFGFSINTLTMFGMVLAIGLLVDDAIVVVENVFRVM	420
AcrB	361	NFRATLIPTIAVPVVLLGTFAVLAAFGFSINTLTMFGMVLAIGLLVDDAIVVVENVERVM	420
MexB	421	AEEGLSPREAARKSMGQIQGALVGIAMVLSAVFLPMAFFGGSTGVIYRQFSITIVSAMAL	480
AcrB	421	AEEGLPPKEATRKSMGQIQGALVGIAMVLSAVFVPMAFFGGSTGAIYRQFSITIVSAMAL	480
MexB	481	SVIVALILTPALCATMLKPIEKGDHGEHKAGFFGWFNRMFLSTTHGYERGVASILKHRAP	540
AcrB	481	SVLVALILTPALCATMLKPIAKGDHGEGKKGFFGWFNRMFEKSTHHYTDSVGGILRSTGR	540
MexB	541	YLLIYVVIVAGMIWMFTRIPTAFLPDEDQGVLFAQVQTPPGSSAERTQVVVDSMREYLLE	600
AcrB	541	YLVLYLIIVVGMAYLFVRLPSSFLPDEDQGVFMTMVQLPAGATQERTQKVLNEVTNYYLT	600
MexB	601	KESSSVSSVFTVTGFNFAGRGQSSGMAFIMLKPWEERPGGENSVFELAKRAQMHFFSFKD KE ++V SVF V GF FAGRG0++G+AF+ LK W +RPG EN V + RA F KD	660
AcrB	601	KEKNNVESVFAVNGFGFAGRGQNTGIAFVSLKDWADRPGEENKVEAITMRATRAFSQIKD	660
MexB	661	AMVFAFAPPSVLELGNATGFDLFLQDQAGVGHEVLLQARNKFLMLAAQNP-ALQRVRPNG AMVFAF P+++ELG ATGFD L D0AG+GHE L 0ARN+ L AA++P L VRPNG	719
AcrB	661	AMVFAFNLPAIVELGTATGFDFELIDQAGLGHEKLTQARNQLLAEAAKHPDMLTSVRPNG	720
MexB	720	MSDEPQYKLEIDDEKASALGVSLADINSTVSIAWGSSYVNDFIDRGRVKRVYLQGRPDAR + D_P0+K++TD_EKA_ALGVS+_DIN+T+_AWG_SYVNDFIDRGRVK+VY+R	779
AcrB	721	LEDTPQFKIDIDQEKAQALGVSINDINTTLGAAWGGSYVNDFIDRGRVKKVYVMSEAKYR	780
MexB	780	MNPDDLSKWVVRNDKGEMVPFNAFATGKWEYGSPKLERYNGVPAMEILGEPAPGLSSGDA M PDD+ WVVR G+MVPF+AF++ +WFYGSP+ FRYNG+P+MEILG+ APG S+G+A	839
AcrB	781	MLPDDIGDWYVRAADGQMVPFSAFSSSRWEYGSPRLERYNGLPSMEILGQAAPGKSTGEA	840
MexB	840	MAAVEEIVKQLPKGVGYSWTGLSYEERLSGSQAPALYALSLLVVFLCLAALYESWSIPFS	899
AcrB	841	MELMEQLASKLPTGVGYDWTGMSYQERLSGNQAPSLYAISLIVVFLCLAALYESWSIPFS	900
MexB	900	VMLVVPLGVIGALLATSMRGLSNDVFFQVGLLTTIGLSAKNAILIVEFAKEL-HEQGKGI	958
AcrB	901	VMLVVPLGVIGALLAATFRGLTNDVYFQVGLLTTIGLSANNAILIVEFAKHL ++6KG+	960
MexB	959	VEAAIEACRMRLRPIVMTSLAFILGVVPLAISTGAGSGSQHAIGTGVIGGMVTATVLAI	1017
AcrB	961	IEATTA WINLIN ITTI JEATTE ISIGAGSGAUNAVGTGWINIAIVEAI IEATLDAVRMRLRPILMTSLAFILGVMPLVISTGAGSGAQNAVGTGVMGGMVTATVLAI	1019

Supplementary Figure 1. Alignment of *E. coli* AcrB and *P. aeruginosa* MexB.

Alignment of Escherichia coli AcrB (Ref. NP_414995.1) and Pseudomonas aeruginosa

MexB (Ref. NP_249117.1) by BLAST. Identity 71%, positives 84%, Gaps 0%.





a The growth of *E. coli* MG1655 $\Delta acrB\Delta tolC$ cell harbouring the plasmid pMMB67HE in the presence of various concentrations of ABI-PP without antibiotics. **b-g** The growth of *E. coli* MG1655 $\Delta acrB\Delta tolC$ cells expressing wild-type MexB (b), MexB(I277W) (c), MexB(A279W) (d), MexB(V612W) (e), MexB(V139W) (f) and MexB(V571W) (g) in the presence of various concentrations of ABI-PP without antibiotics. These tests were performed in triplicates. Abbreviations: PP, ABI-PP inhibitor.



Supplementary Figure 3. Growth of *Escherichia coli* cells in the presence of 0.016 µg/mL of levofloxacin and various concentrations of ABI-PP

a The growth of *E. coli* MG1655 $\Delta acrB\Delta tolC$ cell harbouring the plasmid pMMB67HE in the presence of 0.016 µg/mL of levofloxacin and various concentrations of ABI-PP. **b-g** The growth of *E. coli* MG1655 $\Delta acrB\Delta tolC$ cells expressing wild-type MexB (b), MexB(I277W) (c), MexB(A279W) (d), MexB(V612W) (e), MexB(V139W) (f) and MexB(V571W) (g) in the presence of 0.016 µg/mL of levofloxacin and various concentrations of ABI-PP. These tests were performed in triplicates. Abbreviations: PP, ABI-PP inhibitor.



Supplementary Figure 4. Growth of *Escherichia coli* cells in the presence of $0.125 \mu g/mL$ of aztreonam and various concentrations of ABI-PP

a The growth of *E. coli* MG1655 $\Delta acrB\Delta tolC$ cell harbouring the plasmid pMMB67HE in the presence of 0.125 µg/mL of aztreonam and various concentrations of ABI-PP. **b-g** The growth of *E. coli* MG1655 $\Delta acrB\Delta tolC$ cells expressing wild-type MexB (b), MexB(I277W) (c), MexB(A279W) (d), MexB(V612W) (e), MexB(V139W) (f) and MexB(V571W) (g) in the presence of 0.125 µg/mL of aztreonam and various concentrations of ABI-PP. These tests were performed in triplicates. Abbreviations: PP, ABI-PP inhibitor.