

```

17978      mnhnikglsmnsvsynrnfktgndqrinhsvflngvasaqmalrlgylgvtvflcasn 60
19606      mnqnikglsmnsvsynrnfktgndqrinhsvflngvasaqmalrlgylgvtvflcasn 60
AC12       mnqnikglsmnsvsynrnfktgndqrinhsvflngvasaqialrlgylgvtvflcasn 60
          *****

17978      tyaavidnsaktleqqtaqtnvaalpaitvkaeqddtyaggqvsskssvflgntkmet 120
19606      tyaavidnstktleqqtaqtnvaalpaitvkaeqddtyaggqvatsnsvflgskkfldt 120
AC12       tyaavidnsaktleqqtaqtnvaalpaitvkaeqddtyaggqvsskssvflgntkmet 120
          *****

17978      pfntiaytekyiadqqakditdviaktgpsifstgaygiaesyaairfrsssvaaagrdt 180
19606      pfntisytdkyiedkqakditeviaatdpsiytngasggwsenyiirgyasst----ndm 176
AC12       pfntiaytdtyiadkqakditsviaktgpsvftnaasggwsenyiirgfessp----sdm 176
          *****

17978      smgglgyiaplyraspemferidvlgkpsamlngmmpgstlgtvnlvpkragddpltrf 240
19606      smnglfgitpfyrtspemfgrvevlgkpsallngmmpagsvggtvnlvtkyaaedpfarl 236
AC12       smnglfgitpyrtpemfekinvlkpsallngmmpptsgigtvdlttkratdepltrl 236
          *****

17978      tttymdsqfghldigrffgenkefgvrfnsvyrdgegpikqekthlflslgldwrgd 300
19606      tttymdaqfghvdvgrffgenkefgvringmyrdgdaavndqskesrlflslgldwqge 296
AC12       tttymdsqfghldvgrffgsdkefgvranavyrdsgpvekqdlkelfslgmdwhgd 296
          *****

17978      narlsadlytsrdhadginrgltilpgvnlpkapnptllnptwtvntnrdtggmirgey 360
19606      narvfvdaydalhdvgtvrgvntavgiqppkadtllspdwgsvetkdkgamirgey 356
AC12       rarvstdlytskdrvdgvtvrginlgkiaipkpppetllnpdwsfvdidqkgamirgey 356
          *****

17978      dlndkwmayatagmskteyntlgaakteiqneagdikfniahlgfkyerksaevglrgkf 420
19606      dlfsqlmayayqsttekyngasagtitstgtlsslqqlafdvdkksadagfkgkf 416
AC12       dltnnmgyatyqgsktekyngamsatvldntgtfqtsglqtfvndkksadavglkgkf 416
          *****

17978      dtgavkhalalnalnathyretdeagirqgfppeg-drvtniyn--pnwgskpnrviapifs 477
19606      etgsvkhqwanatyntqdddygyriipgfsdpvitniydpnpwgpkeftpp-flfh 475
AC12       ntgavhgwvwanatyqkhnqddyvrvnvaeg--wtnlyd--piwgkavpfnap-lisn 471
          *****

17978      nkenltsvgladtmsvlqdvqlltllgryqnipteslngqgvrringrydknaitpaaai 537
19606      stltsstsfgladtlsfaqdkvqlltllgrrhqvkatss-----vntlpenaksattpgval 530
AC12       selqlnsygladtlsfiedrltllgvryqvevsnlfv--ptstltykknattpgaaai 529
          *****

17978      lvkandhvslyanyieglssggtvptgqnyenegqifspfkktkmetgikidsnftntf 597
19606      likatdkisvyanyiegltkgdqapat--asnpeifppqktkqqlgkvdltgftahtl 588
AC12       lfkatdkvisyanyiegltkgdaapl--aanypitifapyktkqtefgvkdqgftahtl 587
          *****

17978      svfqikrsnyvikstdekiksiygadqeqrnrngvewgfygspvndirvmggltyidpki 657
19606      safeitkpslylpsklvnnlptfvsdqeqrnrngiewsfsgpiehvrimggftyldepel 648
AC12       slfeikkpsaytdlvt-----niytsggeqrnrngiewsfsgspievrlimggasyiepev 642
          *****

17978      tktriaseelnvavgvpkvqaklgiegdvsiyigp--lrltanataaskqiyndnstsvpg 716
19606      tktksggndghtavavpknqaklgaewdtqvaqgtllsgninavskqiyinaentlsvpg 708
AC12       ktaiqsnegnmavgvpkqgklgvewdnevaqgvltlssnatavskqiyidqentlhipg 702
          *****

17978      revyigaryntkiaqfpvtirgnidnltnksywsmpqftnlmlgaprtyllsatidf 774
19606      rttldvgarystkvdehpvtfraniynltnkaywaqqltnlalgaprtymlsvsydf 766
AC12       rttldvgaryktsisnhpitfkadinnltnkaywmpklsnlalgaprtymlsvsydf 760
          *****

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% identity	ATCC 17978	ATCC 19606	AC12
ATCC 17978	100.00	59.84	61.92
ATCC 19606	59.84	100.00	70.28
AC12	61.92	70.28	100.00

Fig S8. Sequence comparison among three *A. baumannii* BauA variants.

The sequences of three BauA variants from *A. baumannii* ATCC 17978, ATCC 19606, and AC12 were aligned using Clustal Omega, and the calculated identity matrix was shown as a table.