**Table S1.** Characterization of selected Peptides based on number of amino acids produced by Gram-positive and Gram-negative organisms in the UniProt Database.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **No.** | **ID** | **Sequence Length** | **Mass (Da)** | **Organism** |
| 1 | A0A0B7MF59 | 50 | 5,557 | *Streptococcus pneumoniae* |
| 2 | A0A0E2P8G1 | 50 | 5,550 | *Streptococcus pseudopneumoniae 22725* |
| 3 | A0A0P0C3P7 | 50 | 5,893 | *Staphylococcus pseudintermedius* |
| 4 | A0A0U0CP47 | 50 | 5,492 | *Streptococcus pseudopneumoniae* |
| 5 | A0A1E9GG72 | 50 | 5,544 | *Streptococcus sp. HMSC073F11* |
| 6 | A0A1S9ZCG0 | 50 | 5,529 | *Streptococcus mitis* |
| 7 | A0A4U1LBM3 | 50 | 5,544 | *Streptococcus mitis* |
| 8 | A0A6G2DJX7 | 50 | 5,588 | *Streptococcus pneumoniae* |
| 9 | A0A6I1TUZ4 | 50 | 5,528 | *Streptococcus mitis* |
| 10 | A0A150NII9 | 50 | 5,192 | *Streptococcus mitis* |
| 11 | B1I839 | 50 | 5,558 | *Streptococcus pneumoniae (strain Hungary19A-6)* |
| 12 | B3A0N4 | 43 | 4,449 | *Weissella paramesenteroides (Leuconostoc paramesenteroides)* |
| 13 | B3EWP5 | 10 | 1,087 | *Lacticaseibacillus paracasei (Lactobacillus paracasei)* |
| 14 | B3EWP7 | 15 | 1,803 | *Bacillus licheniformis* |
| 15 | C0HJC0 | 30 | 3,498 | *Lactiplantibacillus plantarum* |
| 16 | C0HJE5 | 15 | 1,618 | *Bacillus subtilis* |
| 17 | C0HK82 | 47 | 4,448 | *Latilactobacillus curvatus* |
| 18 | C0HL39 | 30 | 2,884 | *Bacillus subtilis* |
| 19 | C0HLT6 | 10 | 1,067 | *Lactiplantibacillus plantarum* |
| 20 | C0HLU4 | 38 | 4,264 | *Weissella confusa (Lactobacillus confusus)* |
| 21 | C1CAF2 | 50 | 5,558 | *Streptococcus pneumoniae (strain 70585)* |
| 22 | C1CBU6 | 50 | 5,557 | *Streptococcus pneumoniae* |
| 23 | C6FX52 | 50 | 5,364 | *Streptomyces actuosus* |
| 24 | D3VML5 | 41 | 4,366 | *Bacillus velezensis* |
| 25 | E0Q3I4 | 47 | 5,235 | *Streptococcus sp. oral taxon 071 str. 73H25AP* |
| 26 | E8K1W4 | 41 | 4,986 | *Streptococcus infantis ATCC 700779* |
| 27 | E9FQC5 | 50 | 5,205 | *Streptococcus sp. M334* |
| 28 | J1P1V9 | 50 | 5,558 | *Streptococcus pneumoniae 2070335* |
| 29 | O07623 | 43 | 4,325 | *Bacillus subtilis (strain 168)* |
| 30 | P0C8P6 | 14 | 1,416 | *Bacillus cereus* |
| 31 | P0C8P7 | 14 | 1,416 | *Bacillus badius* |
| 32 | P0C912 | 14 | 1,365 | *Amycolatopsis sp. (strain MI481-42F4 / FERM P-12739)* |
| 33 | P0DQM5 | 47 | 5,013 | *Burkholderia thailandensis* |
| 34 | P01547 | 45 | 4,680 | *Myxococcus fulvus* |
| 35 | P02987 | 49 | 5,157 | *Escherichia coli* |
| 36 | P36499 | 50 | 5,677 | *Lactococcus lactis subsp. lactis (Streptococcus lactis)* |
| 37 | P36502 | 19 | 2,063 | *Streptomyces sp. (strain R2075) (Streptoverticillium sp. (strain R2075))* |
| 38 | P36503 | 19 | 2,007 | *Streptomyces griseoluteus* |
| 39 | P36504 | 19 | 2,069 | *Streptomyces griseoverticillatus* |
| 40 | P36961 | 39 | 4,346 | *Lactococcus lactis subsp. lactis (Streptococcus lactis)* |
| 41 | P36962 | 35 | 4,110 | *Lactococcus lactis subsp. lactis (Streptococcus lactis)* |
| 42 | P42723 | 42 | 4,406 | *Rhizobium leguminosarum bv. trifolii* |
| 43 | P80214 | 48 | 5,458 | *Lactiplantibacillus plantarum* |
| 44 | P80323 | 38 | 3,431 | *Latilactobacillus curvatus* |
| 45 | P80493 | 42 | 4,402 | *Latilactobacillus sakei* |
| 46 | P80666 | 22 | 2,425 | *Streptococcus mutans* |
| 47 | P80925 | 43 | 4,290 | *Enterococcus mundtii* |
| 48 | P80953 | 41 | 4,061 | *Latilactobacillus sakei* |
| 49 | P80959 | 31 | 3,358 | *Lacticaseibacillus paracasei* |
| 50 | P81052 | 31 | 3,466 | *Leuconostoc mesenteroides* |
| 51 | P81053 | 43 | 4,598 | *Leuconostoc mesenteroides* |
| 52 | P83002 | 37 | 4,145 | *Lactococcus lactis subsp. lactis (Streptococcus lactis)* |
| 53 | P83375 | 9 | 1,095 | *Serratia plymuthica* |
| 54 | P83378 | 22 | 2,545 | *Serratia plymuthica* |
| 55 | P83674 | 47 | 5,360 | *Ruminococcus gnavus* |
| 56 | P84763 | 18 | 2,027 | *Bacillus thuringiensis subsp. entomocidus* |
| 57 | P84886 | 30 | 3,024 | *Latilactobacillus curvatus* |
| 58 | P84962 | 43 | 4,525 | *Carnobacterium divergens* |
| 59 | P85065 | 24 | 2,355 | *Microbispora sp. (strain 107891)* |
| 60 | P85148 | 39 | 4,126 | *Enterococcus faecium* |
| 61 | P85833 | 29 | 3,353 | *Lactococcus lactis subsp. lactis (Streptococcus lactis)* |
| 62 | P85876 | 38 | 3,932 | *Enterococcus faecium* |
| 63 | P86291 | 41 | 4,602 | *Lactococcus sp.* |
| 64 | P86386 | 25 | 2,728 | *Streptococcus mutans* |
| 65 | P86393 | 39 | 4,630 | *Paenibacillus polymyxa* |
| 66 | P86394 | 35 | 3,629 | *Niallia circulans (Bacillus circulans)* |
| 67 | P86395 | 30 | 3,466 | *Paenibacillus polymyxa* |
| 68 | P86526 | 13 | 1,414 | *Lacticaseibacillus rhamnosus* |
| 69 | Q7M0J8 | 14 | 1,413 | *Planobispora rosea* |
| 70 | Q48501 | 46 | 5,331 | *Lactobacillus acidophilus* |
| 71 | S7Z987 | 50 | 5,511 | *Streptococcus mitis 29/42* |
| 72 | V8IIJ8 | 50 | 5,492 | *Streptococcus pseudopneumoniae 5247* |

**Table 3:** Antimicrobial bacteriocins with their sequence and 3D structures

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **No.** | **UniProt ID** | **AMP Score** | **Sequence length** | **Sequence** | **3D Structure** |
| 1 |  A0A0B7MF59 |  0.556 | 50 | MMKNLNNYREISNKELQEIKGGFGVGVGIALFMAGYTIGKDLRKKFGKSC | Shape  Description automatically generated |
| 2 | A0A0B7MF59m | 0.507 | 50 | MMKNLNNYHEISNKELQEIKGGFGVGVGIALFMAGYTIGKDLRKKFGKSC | Shape  Description automatically generated |
| 3 | A0A1S9ZCG0 | 0.509 | 50 | MMKDLNNYREISNKELQEIKGGFGAGVGIALFMAGYTIGKDLRKKFGKSC | A picture containing tool, opener, weapon  Description automatically generated |
| 4 | A0A1S9ZCG0m | 0.579 | 50 | MVKDLNNYREISNKELQEIKGGFGAGVGIALFMAGYTIGKDLRKKFGKSC | A picture containing tool, opener  Description automatically generated |
| 5 | A0A6I1TUZ4m | 0.581 | 50 | MMKGLNNYREITNKELQEIKGGFGVGVGIALFMIAYTIGKDLRKKFGKSC |  |
| 6 | B3A0N4 | 1.084 | 43 | KNYGNGVYCNKHKCSVDWATFSANIANNSVAMAGLTGGNAGNK | A picture containing logo  Description automatically generated |
| 7 | B3A0N4m | 1.409 | 43 | KNYGNGVYCNKHKCSVDWATFSANIANFSVAMAGLTGGNAGNK | A picture containing shape  Description automatically generated |
| 8 | C0HJC0 | 0.506 | 30 | GRADYNFGYGLGRGTRKFFNGHGRWVRKTF | Shape, arrow  Description automatically generated |
| 9 | C0HL39m | 0.597 | 30 | LGFQLNKGCATCSIGAACLVDGPIPDEIAG | A picture containing shape  Description automatically generated |
| 10 | C1CBU6 | 0.556 | 50 | MMKNLNNYREISNKELQEIKGGFGVGVGIALFMAGYTIGKDLRKKFGKSC | A picture containing opener  Description automatically generated |
| 11 | J1P1V9m | 0.542 | 50 | MMKHLNNYREISNKELQEIKGGFGVGVGIALFMAGYTIGKDLRKKFGKSC | Logo, company name  Description automatically generated |
| 12 | O07623m | 0.832 | 43 | MKKAVIVENKGCATCSIGAACLVDGPIKDFEIAGATGLFGLWG | A picture containing hanger, opener  Description automatically generated |
| 13 | P01547 | 0.961 | 45 | ANCSCSTASDYCPILTFCTTGTACSYTPTGCGTGWVYCACNGNFY | Icon  Description automatically generated |
| 14 | P01547m | 1.048 | 45 | ANCSRSTASDYCPILTFCTTGTACSYTPTGCGTGWVYCACNGNFY | Icon  Description automatically generated with medium confidence |
| 15 | P36502 | 1.648 | 19 | CRQSCSFGPLTFVCDGNTK | Shape  Description automatically generated with low confidence |
| 16 | P36502m | 1.560 | 19 | CRQSCSFGYLTFVCDGNTK | Icon  Description automatically generated with low confidence |
| 17 | P36503 | 1.298 | 19 | CANSCSYGPLTWSCDGNTK | Icon  Description automatically generated |
| 18 | P36503m | 0.837 | 19 | CALSCSYGPLTWSCDGNTK | **Shape, arrow  Description automatically generated** |
| 19 | P36504 | 1.002 | 19 | CKQSCSFGPFTFVCDGNTK | **Icon  Description automatically generated** |
| 20 | P36504m | 0.743 | 19 | CKQSCSFGPFLFVCDGNTK | **Icon  Description automatically generated** |
| 21 | P36961 | 0.923 | 39 | GTWDDIGQGIGRVAYWVGKAMGNMSDVNQASRINRKKKH | **Shape, arrow  Description automatically generated** |
| 22 | P36961m | 0.733 | 39 | GTWDDIGQGILRVAYWVGKAMGNMSDVNQASRINRKKKH | **Shape, arrow  Description automatically generated** |
| 23 | P42723 | 0.661 | 42 | MDNKVAKNVEVKKGSIKATFKAAVLKSKTKVDIGGSRQGCVA | **A blue octopus on a white background  Description automatically generated with low confidence** |
| 24 | P80214 | 1.190 | 48 | MKIQIKGMKQLSNKEMQKIVGGKSSAYSLQMGATAIKQVKKLFKKWGW | **A picture containing opener  Description automatically generated** |
| 25 | P80214m | 1.312 | 48 | MKIQIKGMKQLSNKEMQKIVGGKSSAYSLQMGATAIFQVKKLFKKWGW | **Icon  Description automatically generated with medium confidence** |
| 26 | P80666 | 1.000 | 22 | FKSWSFCTPGCAKTGSFNSYCC | **A picture containing tool  Description automatically generated** |
| 27 | P80666m | 0.833 | 22 | FKSWSFCTPICAKTGSFNSYCC | **Icon  Description automatically generated** |
| 28 | P80925 | 1.576 | 43 | KYYGNGVSCNKKGCSVDWGKAIGIIGNNSAANLATGGAAGWSK | **Logo  Description automatically generated** |
| 29 | P80925m | 1.697 | 43 | KLYGNGVSCNKKGCSVDWGKAIGIIGNNSAANLATGGAAGWSK |  |
| 30 | P80959 | 1.193 | 31 | GMSGYIQGIPDFLKGYLHGISAANKHKKGRL | **A blue dragonfly on a white background  Description automatically generated with low confidence** |
| 31 | P80959m | 0.949 | 31 | GMSGYIQGIPDFLKGYLHGISAANKHKQGRL | **Logo  Description automatically generated** |
| 32 | P81053 | 0.986 | 43 | KNYGNGVHCTKKGCSVDWGYAWTNIANNSVMNGLTGGNAGWHN | **A picture containing shape  Description automatically generated** |
| 33 | P81053m | 0.925 | 43 | KNLGNGVHCTKKGCSVDWGYAWTNIANNSVMNGLTGGNAGWHN | **A picture containing logo  Description automatically generated** |
| 34 | P83002 | 0.959 | 37 | TSYGNGVHCNKSKCWIDVSELETYKAGTVSNPKDILW | **A picture containing icon  Description automatically generated** |
| 35 | P83002m | 0.546 | 37 | TSYGNGVHCNKSKCWIDVSELETYKAGTVSFPKDILW | **A picture containing icon  Description automatically generated** |
| 36 | P83378 | 1.001 | 22 | ALPKKLKYLNLFNDGFNYMGVV | **A picture containing shape  Description automatically generated** |
| 37 | P83378m | 0.522 | 22 | ALPKSLKYLNLFNDGFNYMGVV | **Icon  Description automatically generated** |
| 38 | P84886 | 1.164 | 30 | AYPGNGVHCGKYSCTVDKQTAIGNIGNNAA | **A picture containing logo  Description automatically generated** |
| 39 | P84886m | 1.340 | 30 | ALPGNGVHCGKYSCTVDKQTAIGNIGNNAA | **A picture containing logo  Description automatically generated** |
| 40 | P84962 | 1.000 | 43 | TKYYGNGVYCNSKKCWVDWGTAQGCIDVVIGQLGGGIPGKGKC | **Logo  Description automatically generated with medium confidence** |
| 41 | P84962m | 1.061 | 43 | TKYAGNGVYCNSKKCWVDWGTAQGCIDVVIGQLGGGIPGKGKC | **A picture containing icon  Description automatically generated** |
| 42 | P85065m | 0.515 | 24 | VTSWSLCTPGCTSFGGGSNCSFCC | **Icon  Description automatically generated with low confidence** |
| 43 | P85148 | 1.039) | 39 | TTKNYGNGVCNSVNWCQCGNVWASCNLATGCAAWLCKLA | A picture containing logo  Description automatically generated |
| 44 | P85148m | 0.920 | 39 | TTKNYGNGVCNSVNWCQCGNCWASCNLATGCAAWLCKLA | A picture containing opener, tool, cable, hydrozoan  Description automatically generated |
| 45 | P85876 | 1.270 | 38 | ATRSYGNGVYCNNSKCWNVGEAKENIAGIVISGKASGL | A pair of blue headphones  Description automatically generated with medium confidence |
| 46 | P85876m | 1.315 | 38 | ATRSYGNGVYCNNSKCWNVGEAKENIAGFVFSGKASGL | Icon  Description automatically generated |
| 47 | P86291 | 0.551 | 41 | TSYGNGVHCNKSKCWIDVSELETYKAGTVSNPKDILWSLKE |  |
| 48 | P86291m | 0.587 | 41 | TSYGNGVHCNKSKCWIDVSELETYKAGTVSNPKFILWSLKE |  |
| 49 | P86393 | 1.005 | 39 | ATYYGNGLYCNKQKHYTWVDWNKASREIGKITVNGWVQH | A pair of blue headphones  Description automatically generated with medium confidence |
| 50 | P86393m | 0.927 | 39 | ATYLGNGLYCNKQKHYTWVDWNKASREIGKITVNGWVQH | Logo  Description automatically generated |
| 51 | P86394 | 1.057 | 35 | VNYGNGVSCSKTKCSVNWGIITHQAFRVTSGVASG | Shape  Description automatically generated |
| 52 | P86394m | 1.010 | 35 | VNYGNGVSCSKTKCSVNWGISTHQAFRVTSGVASG | A picture containing shape  Description automatically generated |
| 53 | S7Z987 | 0.781 | 50 | MIKDLNNYREISNKELQEIKGGFGAGVGIALFMAGYTIGKDLRKKFGKSC | A picture containing opener, tool  Description automatically generated |
| 54 | S7Z987m | 0.521 | 50 | MIKDLNNYREISNKELQEIKGGFGAGVGIALFMAGYTIGKDLRYKFGKSC | A picture containing logo  Description automatically generated |
| 55 | V8IIJ8m | 0.519 | 50 | MLKDLNNYCEISTKELQEIKGGFGVGVGIALFMAGYTIGKDLRKKFGKSC | Shape  Description automatically generated |

**Table S3:** Virulent proteins present in *Serratia marcescens* and their physiochemical properties

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **No.** | **ID** | **Molecular Weight (Da)** | **Theoretical PI** | **Instability index** | **Extinction coefficients** | **GRAVY** |
| 1 | UPI0002AF2D88 | 32230.53 | 4.99 | 46.73 | 0.509 | 0.303 |
| 2 | UPI0002AF3E70 | 43195.42 | 5.13 | 28.76 | 0.441 | -0.153 |
| 3 | UPI0002AF25B6 | 36786.57 | 5.36 | 23.48 | 1.165 | -0.174 |
| 4 | UPI0002AF42AC | 27628.2 | 5.27 | 51.19 | 1.083 | -0.497 |
| 5 | UPI0002AF278A | 27862.93 | 4.62 | 27.77 | 0.518 | -0.34 |
| 6 | UPI0002AF352E | 38210.07 | 4.81 | 53.78 | 0.705 | -0.539 |
| 7 | UPI0002AF3971 | 37957.56 | 9 | 31.11 | 0.23 | 0.174 |
| 8 | UPI0002AF4256 | 59164.94 | 5.9 | 37.52 | 1.686 | -0.402 |
| 9 | UPI0002AF4326 | 23738.69 | 6.75 | 32.63 | 0.546 | 0.002 |
| 10 | UPI0002B86BA6 | 48972.25 | 6.29 | 47.66 | 0.571 | -0.051 |
| 11 | UPI0002B88CF4 | 33642.28 | 7.82 | 43.07 | 0.814 | -0.273 |
| 12 | UPI0002B87870 | 42206.51 | 8.41 | 36.7 | 0.742 | 0.089 |
| 13 | UPI0002C18763 | 47271.17 | 5.27 | 30.16 | 1.506 | -0.196 |
| 14 | UPI0002DC8D87 | 25635.4 | 5.95 | 20.21 | 0.447 | 0.2 |
| 15 | UPI0002E08E29 | 54743.61 | 5.38 | 40.39 | 1.114 | -0.154 |
| 16 | UPI0002E65B3D | 50931.39 | 5.25 | 25.39 | 0.502 | 0.032 |
| 17 | UPI0002F3BD56 | 42785.47 | 9.66 | 45.41 | 1.015 | 0.011 |
| 18 | UPI0002F98715 | 56658.38 | 5.05 | 47.43 | 1.311 | -0.532 |
| 19 | UPI0003C9AF1D | 52615.25 | 6.41 | 53.93 | 3.039 | -0.445 |
| 20 | UPI0003C9BA85 | 73818.73 | 5.68 | 23.04 | 1.534 | -0.512 |
| 21 | UPI0003C9C1DC | 92534.12 | 5.55 | 32.34 | 1.608 | -0.405 |
| 22 | UPI0003C9C802 | 32747.57 | 5.6 | 37.95 | 1.176 | -0.169 |
| 23 | UPI0003C9C858 | 99789.17 | 6.11 | 36.97 | 1.058 | 0.172 |
| 24 | UPI0003C9CE2E | 48794.87 | 5.97 | 27.16 | 0.588 | -0.048 |
| 25 | UPI0003C9D4FC | 92283.32 | 6.59 | 26.12 | 1.363 | -0.388 |
| 26 | UPI0003C9D160 | 25445.28 | 6.08 | 35.19 | 0.784 | -0.252 |
| 27 | UPI0003C9E72D | 165116.71 | 8.82 | 24.08 | 0.444 | -0.554 |
| 28 | UPI0003C9E649 | 54852.47 | 6.15 | 33.88 | 1.166 | -0.629 |
| 29 | UPI0003C9EDE5 | 49575.12 | 6.4 | 25.4 | 0.842 | -0.045 |
| 30 | UPI0003C95C92 | 33949.88 | 5.53 | 39.43 | 0.551 | -0.052 |
| 31 | UPI0003C95D10 | 52721.57 | 6.13 | 44.49 | 0.633 | -0.065 |
| 32 | UPI0003C95E0A | 39726.64 | 5.45 | 42.78 | 1.722 | -0.285 |
| 33 | UPI0003C97B07 | 47688.62 | 9.58 | 39.06 | 1.017 | -0.044 |
| 34 | UPI0003C97E1F | 111609.1 | 5.65 | 34.99 | 0.773 | 0.251 |
| 35 | UPI0003C99A74 | 30686.31 | 5.41 | 29.6 | 0.198 | -0.011 |
| 36 | UPI0003C953F8 | 313314.86 | 5.41 | 46.08 | 1.031 | -0.16 |
| 37 | UPI0003C967AA | 49222.6 | 9.11 | 35.81 | 1.308 | 0.263 |
| 38 | UPI0003C982D4 | 71335.45 | 5.4 | 31.67 | 1.582 | -0.509 |
| 39 | UPI0003C992A0 | 84289.03 | 5.87 | 35.21 | 0.409 | -0.303 |
| 40 | UPI0003C9590B | 33340.73 | 5.85 | 35.18 | 0.695 | -0.102 |
| 41 | UPI0003C9723E | 80166.9 | 6.28 | 39.18 | 0.778 | -0.182 |
| 42 | UPI0003C9725E | 32808.23 | 6.27 | 29.71 | 0.783 | -0.044 |
| 43 | UPI0003C9839B | 52187.73 | 4.64 | 24.32 | 1.386 | -0.392 |
| 44 | UPI0003C94769 | 50540.6 | 5.22 | 29.12 | 0.833 | -0.196 |
| 45 | UPI0003C95066 | 57890.07 | 6.11 | 44.83 | 1.559 | -0.406 |
| 46 | UPI0003C98210 | 87545.47 | 5.67 | 26.49 | 1.437 | -0.55 |
| 47 | UPI0003C99600 | 60141.85 | 4.92 | 40.54 | 0.547 | -0.128 |
| 48 | UPI000012A963 | 21809.64 | 8.87 | 45.84 | 1.115 | -0.002 |
| 49 | UPI000320AA72 | 55709.48 | 6.53 | 35.67 | 1.932 | 0.624 |
| 50 | UPI000320ADAB | 93267.75 | 5.69 | 34.75 | 1.571 | -0.431 |
| 51 | UPI000320CDF9 | 32400.77 | 4.68 | 52.69 | 1.564 | -0.275 |
| 52 | UPI000320FFD4 | 111696.07 | 5.41 | 32.15 | 0.617 | 0.392 |
| 53 | UPI000321B1CB | 101694.21 | 5.51 | 43.58 | 0.416 | -0.094 |
| 54 | UPI000321CC59 | 143884.12 | 5.34 | 36.37 | 1.09 | -0.05 |
| 55 | UPI000322E099 | 95038.34 | 8.23 | 32.8 | 0.96 | -0.092 |
| 56 | UPI000322F750 | 58166.99 | 4.77 | 20.81 | 0.479 | -0.376 |
| 57 | UPI000323D213 | 53516.47 | 4.66 | 38.49 | 0.72 | -0.364 |
| 58 | UPI000323EE92 | 39365.14 | 5.2 | 35.5 | 1.663 | -0.238 |
| 59 | UPI000324AAB7 | 110572.43 | 8.98 | 37.35 | 0.596 | 0.296 |
| 60 | UPI000324CA26 | 135143.64 | 8.58 | 34.25 | 1.324 | -0.27 |
| 61 | UPI000324DAE7 | 41918.13 | 5.76 | 40.56 | 0.659 | -0.103 |
| 62 | UPI000325E0FA | 92142.77 | 5.69 | 31.87 | 1.479 | -0.4 |
| 63 | UPI000326AD5D | 41726.71 | 6.33 | 34.26 | 1.255 | -0.21 |
| 64 | UPI000326D3C8 | 113221.97 | 6.34 | 46.15 | 1.084 | -0.159 |
| 65 | UPI0003211BB5 | 45918.15 | 5.61 | 42.88 | 0.672 | -0.258 |
| 66 | UPI0003213C38 | 82299.04 | 5.17 | 32 | 1.55 | -0.519 |
| 67 | UPI0003224D5E | 53029.24 | 5.52 | 36.96 | 0.984 | -0.184 |
| 68 | UPI0003227B2E | 57356.99 | 4.89 | 29.16 | 0.21 | 0.018 |
| 69 | UPI0003229BFB | 50493 | 4.55 | 21.85 | 1.264 | -0.371 |
| 70 | UPI0003235D33 | 36028.5 | 4.93 | 26 | 1.445 | -0.321 |
| 71 | UPI0003236E46 | 37704.23 | 5.23 | 39.56 | 0.568 | -0.19 |
| 72 | UPI0003236F10 | 84272.26 | 6.15 | 38.95 | 1.079 | -0.36 |
| 73 | UPI0003245BEC | 36613.8 | 4.65 | 46.77 | 0.163 | -0.163 |
| 74 | UPI0003246B37 | 49786.64 | 5.92 | 48.59 | 1.619 | -0.261 |
| 75 | UPI0003251A93 | 112931.17 | 5.78 | 32.07 | 0.721 | 0.28 |
| 76 | UPI0003259EC6 | 32073.91 | 5.81 | 39.95 | 1.076 | -0.053 |
| 77 | UPI0003261DB2 | 42571.03 | 6.87 | 40.36 | 1.056 | -0.183 |
| 78 | UPI0003262E92 | 91538.01 | 8.33 | 35.48 | 1.256 | -0.356 |
| 79 | UPI00032087BB | 72878.04 | 9.4 | 37.26 | 2.301 | 0.539 |
| 80 | UPI00032128B6 | 61170.15 | 5.89 | 31.48 | 1.267 | -0.268 |
| 81 | UPI00032189DE | 112422.67 | 7.01 | 34.27 | 0.75 | 0.332 |
| 82 | UPI00032237FC | 26520.96 | 4.96 | 35.12 | 0.169 | -0.234 |
| 83 | UPI00032291F5 | 52656.09 | 5.57 | 41.52 | 1.109 | -0.198 |
| 84 | UPI00032325DB | 107539.05 | 4.87 | 41.5 | 0.824 | -0.006 |
| 85 | UPI00032386A1 | 83329.46 | 6.86 | 39 | 1.024 | 0.439 |
| 86 | UPI00032427A7 | 33087.79 | 9.44 | 30.1 | 1.119 | -0.303 |
| 87 | UPI00032500A3 | 82587.34 | 6.5 | 32.25 | 1.378 | -0.574 |
| 88 | UPI00032555AB | 110446.43 | 6.04 | 37.2 | 0.678 | 0.36 |
| 89 | UPI00032589FC | 42638.09 | 5.02 | 26.76 | 0.677 | -0.336 |
| 90 | UPI00032621DF | 48874.27 | 6.15 | 36.04 | 1.302 | -0.331 |
| 91 | UPI00032632F6 | 49390.58 | 4.49 | 19.84 | 1.594 | -0.456 |
| 92 | UPI00032637BE | 27676.29 | 9.05 | 48.83 | 0.865 | 0.777 |
| 93 | UPI00032649DA | 32665 | 5.6 | 34.96 | 0.618 | -0.026 |
| 94 | UPI000321385E | 91236.31 | 6.17 | 33.03 | 1.446 | -0.372 |
| 95 | UPI000322301E | 42488.15 | 10.02 | 39.91 | 1.5 | 0.749 |
| 96 | UPI000322408E | 59726.6 | 5.16 | 43.51 | 1.008 | -0.193 |
| 97 | UPI000324881A | 37695.86 | 8.43 | 42.58 | 0.201 | 0.001 |
| 98 | UPI000326241C | 49944.92 | 6.87 | 16.12 | 0.408 | -0.512 |
| 99 | UPI000326561B | 36506.65 | 5.61 | 56.94 | 1.055 | -0.32 |
| 100 | UPI0003211044 | 38959.03 | 5.95 | 55.59 | 0.615 | -0.485 |
| 101 | UPI0003218851 | 38576.49 | 8.27 | 51.76 | 1.221 | -0.129 |
| 102 | UPI0003222101 | 39711.76 | 8.9 | 22.43 | 1.547 | -0.333 |
| 103 | UPI0003226237 | 113297.76 | 6.2 | 32.49 | 0.843 | 0.192 |
| 104 | UPI0003227771 | 57867.08 | 4.85 | 39.42 | 0.416 | -0.077 |
| 105 | UPI0003235528 | 91673.41 | 6.08 | 29.33 | 1.547 | -0.402 |
| 106 | UPI0003235719 | 79438.9 | 6.37 | 46.1 | 1.713 | -0.513 |
| 107 | UPI0003257892 | 41176.94 | 5.99 | 40.25 | 1.102 | -0.067 |
| 108 | UPI0003265092 | 80149.13 | 6.19 | 29.28 | 0.678 | -0.133 |
| 109 | UPI0003265136 | 28203.26 | 6.89 | 34.79 | 0.837 | 1.06 |
| 110 | UPI0003267316 | 64767.18 | 7.08 | 37.82 | 0.703 | 0.103 |
| 111 | UPI000313986E | 34472.34 | 4.73 | 34.19 | 0.346 | -0.492 |

**Table S4:** Likelihood of Subcellular localization for virulent proteins in *Serratia marscens*

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  | **UniProt ID** | **Extracellular** | **Periplasmic** | **Outer Membrane** | **Cytoplasmic Membrane** | **Cytoplasmic** | **Final prediction** |
| 1 | UPI0002AF3E70 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 2 | UPI0002AF25B6 | 0.01 | 0.48 | 0.01 | 0.24 | 9.26 | Periplasmic |
| 3 | UPI0002AF278A | 10.00 | 0.00 | 0.00 | 0.00 | 0.00 | Extracellular |
| 4 | UPI0002AF3971 | 0.11 | 9.76 | 0.06 | 0.00 | 0.00 | Periplasmic |
| 5 | UPI0002AF4256 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 6 | UPI0002AF4326 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 7 | UPI0002B87870 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 8 | UPI0002C18763 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 9 | UPI0002DC8D87 | 0.01 | 0.48 | 0.01 | 0.24 | 9.26 | Periplasmic |
| 10 | UPI0002E65B3D | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 11 | UPI0003C9BA85 | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| 12 | UPI0003C9C1DC | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| 13 | UPI0003C9C802 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 14 | UPI0003C9C858 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 15 | UPI0003C9CE2E | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 16 | UPI0003C9D4FC | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| 17 | UPI0003C9D160 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 18 | UPI0003C9E72D | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| 19 | UPI0003C9E649 | 0.05 | 4.48 | 0.00 | 0.06 | 5.41 | Unknown(may have multiple site |
| 20 | UPI0003C9EDE5 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 21 | UPI0003C95C92 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 22 | UPI0003C97B07 | 0.26 | 0.26 | 0.01 | 0.51 | 8.96 | Periplasmic |
| 23 | UPI0003C97E1F | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 24 | UPI0003C99A74 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 25 | UPI0003C967AA | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 2627 | UPI0003C982D4 | 0.01 | 0.48 | 0.01 | 0.24 | 9.26 | Periplasmic |
| UPI0003C992A0 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 28 | UPI0003C9590B | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 29 | UPI0003C9723E | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 30 | UPI0003C9725E | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 31 | UPI0003C9839B | 10.00 | 0.00 | 0.00 | 0.00 | 0.00 | Extracellular |
| 32 | UPI0003C94769 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 33 | UPI0003C98210 | 0.01 | 0.48 | 0.01 | 0.24 | 9.26 | Periplasmic |
| 34 | UPI000320AA72 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 35 | UPI000320ADAB | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| 36 | UPI000320FFD4 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 37 | UPI000321CC59 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 38 | UPI000322E099 | 0.38 | 0.09 | 9.49 | 0.01 | 0.03 | OuterMembrane |
| 39 | UPI000322F750 | 9.96 | 0.00 | 0.04 | 0.00 | 0.00 | Extracellular |
| 40 | UPI000323D213 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 41 | UPI000323EE92 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 42 | UPI000324AAB7 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 43 | UPI000324CA26 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 44 | UPI000325E0FA | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| 45 | UPI000326AD5D | 0.03 | 0.03 | 9.92 | 0.01 | 0.01 | OuterMembrane |
| 4647 | UPI0003213C38 | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| UPI0003224D5E | 0.26 | 0.26 | 0.01 | 0.51 | 8.96 | Periplasmic |
| 48 | UPI0003227B2E | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 49 | UPI0003229BFB | 10.00 | 0.00 | 0.00 | 0.00 | 0.00 | Extracellular |
| 50 | UPI0003235D33 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 51 | UPI0003236E46 | 0.00 | 0.00 | 0.00 | 7.88 | 2.11 | CytoplasmicMembrane |
| 52 | UPI0003236F10 | 0.26 | 0.26 | 0.01 | 0.51 | 8.96 | Extracellular |
| 53 | UPI0003251A93 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 54 | UPI0003259EC6 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 55 | UPI0003262E92 | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| 56 | UPI00032087BB | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 57 | UPI00032128B6 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 58 | UPI00032189DE | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 59 | UPI00032237FC | 0.11 | 9.44 | 0.06 | 0.06 | 0.33 | Periplasmic |
| 60 | UPI00032386A1 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 61 | UPI00032427A7 | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 62 | UPI00032500A3 | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| 63 | UPI00032555AB | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 64 | UPI00032589FC | 9.84 | 0.16 | 0.00 | 0.00 | 0.00 | Extracellular |
| 65 | UPI00032621DF | 0.01 | 0.48 | 0.01 | 0.24 | 9.26 | Periplasmic |
| 66 | UPI00032632F6 | 10.00 | 0.00 | 0.00 | 0.00 | 0.00 | Extracellular |
| 67 | UPI00032649DA | 0.00 | 0.01 | 0.00 | 0.01 | 9.97 | Periplasmic |
| 68 | UPI000321385E | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| 69 | UPI000322301E | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 70 | UPI000326241C | 9.71 | 0.20 | 0.07 | 0.01 | 0.01 | Extracellular |
| 71 | UPI0003222101 | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| 72 | UPI0003226237 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 73 | UPI0003227771 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 74 | UPI0003235528 | 0.00 | 0.00 | 10.00 | 0.00 | 0.00 | OuterMembrane |
| 75 | UPI0003265092 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 76 | UPI0003265136 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 77 | UPI0003267316 | 0.00 | 0.00 | 0.00 | 10.00 | 0.00 | CytoplasmicMembrane |
| 78 | UPI000313986E | 10.00 | 0.00 | 0.00 | 0.00 | 0.00 | Extracellular |

**Table S5:** Docking results between proteinUPI0002AF278A (*Serratia marcescens*)and 55 bacteriocins peptides.

|  |  |  |
| --- | --- | --- |
| **Sr #** | **Peptide ID** | **Peptide-protein interaction** |
| **Peptide residue** | **Protein residue** | **Bond Length(A)** | **Score** |
| 1 | A0A0B7MF59m | No interaction |
| 2 | A0A0B7MF59w | Gly(22) | Phe(41) | 2.21 | -9.47 |
| Phe(23) | Gln(15) | 3.16 |
| 3 | A0A1S9ZCG0 | Leu(16) | Gln(235) | 2.79 | -2.37 |
| 4 | A0A1S9ZCG0m | Leu(16) | Gln(235) | 2.79 | -4.64 |
| 5 | A0A6I1TUZ4m | Gly(21) | Gln(51) | 2.65 | -6.91 |
| Leu(31) | Gln(67) | 3.11 |
| Gly(24) | Gln(67) | 2.54 |
| Val(27) | Gln(67) | 2.84 |
| 6 | B3A0N4 | Trp(18) | Gln(16) | 2.65 | -5.89 |
| Ser(15) | Asp(20) | 3.23 |
| 7 | B3A0N4m | Trp(18) | Gln(16) | 3.20 | -7.93 |
| 8 | C0HJC0 | Gly(1) | Asn(232) | 2.93 | -9.92 |
| 9 | C0HL39m | Cys(18) | Thr(70) | 2.54 | -2.68 |
| Thr(11) | Gly(53) | 2.68 |
| 10 | C1CBU6 | Gly(22) | Phe(41) | 2.21 | -12.94 |
| Phe(23) | Gln(15) | 3.16 |
| 11 | J1P1V9m | Gly(22) | Phe(41) | 2.21 | -9.34 |
| Phe(23) | Gln(15) | 3.16 |
| 12 | O07623m | Asn(9) | Ser(186) | 2.51 | -1.03 |
| Leu(38) | Glu(189) | 2.81 |
| 13 | P01547 | Cys(18) | Ser(132) | 2.57 | -10.56 |
| Thr(16) | Ser(131) | 2.58 |
| Leu(15) | Glu(189) | 2.69 |
| Tyr(11) | Ser(119) | 2.16 |
| Val(36) | Ser(30) | 2.45 |
| 14 | P01547m | Phe(17) | Ala(39) | 3.22 | -13.40 |
| Trp(35) | Glu(228) | 3.03 |
| Thr(16) | Asn(18) | 1.99 |
| 15 | P36502 | Cyx(14) | Arg(238) | 3.11 | -15.67 |
| Ser(6) | Glu(241) | 2.17 |
| Cyx(5) | Ser(244) | 3.14 |
| Thr(18) | Glu(241) | 2.57 |
| Phe(7) | Ser(244 | 2.59 |
| 16 | P36502m | No interaction |
| 17 | P36503 | Asn(3) | Gln(15) | 2.60 | -11.84 |
| Asn(3) | Asn(232) | 3.24 |
| Cyx(1) | Phe(41) | 2.73 |
| Tyr(7) | Val(72) | 3.06 |
| Trp(12) | Gly(69) | 3.09 |
| 18 | P36503m | Cys(14) | Leu(45) | 2.65 | -9.92 |
| 19 | P36504 | No interaction |
| 20 | P36504m | No interaction |
| 21 | P36961 | Arg(32) | Glu(189) | 2.84 | -4.03 |
| Asn(34) | Val(187) | 3.11 |
| 22 | P36961m | Asn(34) | Asp(101) | 3.03 | -6.87 |
| Gln(29) | Glu(185) | 2.35 |
| Ser(25) | Val(187) | 2.56 |
| 23 | P42723 | Val(31) | Glu(189) | 2.43 | -15.57 |
| 24 | P80214 | Lys(44) | Gln(193) | 2.67 | -4.60 |
| Lys(40) | Glu(194) | 2.69 |
| 25 | P80214m | Trp(46) | Asn(28) | 1.67 | -7.01 |
| Leu(11) | Glu(189) | 2.16 |
| Lys(9) | Ser(119) | 3.19 |
| 26 | P80666 | Trp(4) | Thr(77) | 2.88 | -10.66 |
| 27 | P80666m | Phe(17) | Pro(108) | 2.70 | -10.21 |
| 28 | P80925 | Gly(36) | Arg(238) | 3.33 | -4.95 |
| Asn(32) | Gln(235) | 2.71 |
| Ser(29) | Val(231) | 2.26 |
| 29 | P80925m | Asn(5) | Gln(235) | 2.77 | -5.43 |
| Asn(5) | Asn(232) | 2.12 |
| Trp(41) | Phe(41) | 3.15 |
| 30 | P80959 | Arg(30) | Ser(186) | 2.77 | -3.70 |
| His(26) | Ala(76) | 2.85 |
| 31 | P80959m | Lys(14) | Gln(237) | 2.67 | -1.87 |
| Gly(15) | Gln(237) | 1.64 |
| 32 | P81053 | Gly(37) | Gln(15) | 2.30 | -14.59 |
| 33 | P81053m | Asn(43) | Asp(20) | 3.25 | -7.90 |
| Asn(32) | Glu(185) | 1.70 |
| 34 | P83002 | Asp(17) | Asn(28) | 2.09 | -14.55 |
| Asp(17) | Asn(24) | 2.77 |
| Glu(20) | Ser(186) | 2.98 |
| 35 | P83002m | Thr(28) | Val(187) | 2.01 | -13.60 |
| Trp(15) | Glu(189) | 2.83 |
| Glu(20) | Arg(79) | 2.51 |
| Asp(17) | Asn(24 | 2.65 |
| 36 | P83378 | Asn(13) | Thr(70) | 2.65 | -5.90 |
| 37 | P83378m | Gly(15) | Ala(39) | 2.79 | -8.58 |
|  Asn(17) |  Ala(39) | 3.10 |
| Met(19) | Gln(235) | 2.68 |
| 38 | P84886 | Tyr(12) | Ser(250) | 2.06 | -4.90 |
| Tyr(12) | Ala(246) | 2.65 |
| Asn(5) |  Met(49) | 2.74 |
| 39 | P84886m | Tyr(12) | Glu(241) | 3.23 | -5.69 |
| 40 | P84962 | Gly(24) | Ser(30) | 3.26 | -8.37 |
| Asp(27) | Thr(221) | 2.44 |
| Leu(33) | Ser(87) | 2.59 |
| Leu(33) | Asn(85) | 3.22 |
| 41 | P84962m | Thr(21) | Gln(37) | 3.26 | -5.49 |
| Cyx(25) | Asn(24) | 2.97 |
| Tyr(9) | Glu(185) | 2.04 |
| Tyr(9) | Arg(79) | 2.91 |
| 42 | P85065m | Cys(23) | Tyr(46) | 2.35 | -10.48 |
| Gly(15) | Gly(71) | 1.79 |
| Ser(13) | Asn(243) | 2.33 |
| 43 | P85148 | Tyr(5) | Glu(185) | 2.69 | -5.99 |
| Leu(27) | Glu(185) | 3.16 |
| Thr(1) | Ser(197) | 2.52 |
| Gly(30) | Val(75) | 1.74 |
| 44 | P85148m | Lys(37) | Val(187) | 2.54 | -13.54 |
| Cys(31) | Phe(134) | 3.12 |
| Cys(31) | Ser(131) | 3.12 |
| Ser(24) | Ser(132) | 3.35 |
| Asn(20) | Asp(94) | 3.17 |
| 45 | P85876 | Gly(28) | Thr(77) | 2.72 | -4.46 |
| Tyr(5) | Leu(45) | 2.81 |
| 46 | P85876m | Ser(36) | Glu(42) | 2.84 | -7.68 |
| Ser(36) | Arg(73) | 2.60 |
| Ser(36) | Asp(43) | 3.26 |
| 47 | P86291 | Asp(17) | Asn(24) | 2.77 | -14.04 |
| Asp(17) | Asn(28) | 2.09 |
| Glu(20) | Ser(186) | 2.98 |
| 48 | P86291m | Asp(17) | Asn(24) | 2.77 | -10.53 |
| Asp(17) | Asn(28) | 2.09 |
| Glu(20) | Ser(186) | 2.98 |
| 49 | P86393 | Asp(20) | Gln(59) | 3.08 | -8.40 |
| Val(19) | Thr(60) | 2.80 |
| 50 | P86393m | Asp(20) | Gln(59) | 2.96 | -9.47 |
| Val(19) | Gln(51) | 2.88 |
| Trp(21) | Leu(66) | 2.24 |
| 51 | P86394 | Ala(33) | His(81) | 2.94 | -16.79 |
| Ala(25) | Ser(182) | 1.96 |
| Thr(22) | Ser(197) | 3.29 |
| Tyr(3) | Asp(43) | 2.66 |
| 52 | P86394m | Phe(26) | Val(187) | 2.99 | -12.47 |
| 53 | S7Z987 | Leu(16) | Gln(235) | 2.79 | -10.58 |
| 54 | S7Z987m | Lys(20) | Val(75) | 2.42 | -10.34 |
| 55 | V8IIJ8m | No interaction |

**Table S6:** Docking results between UPI0003C9BA85 (*Serratia* *marcescens*)and 55 bacteriocins peptides.

|  |  |  |
| --- | --- | --- |
| **Sr #** | **Peptide ID** | **Peptide-protein interaction** |
| **Peptide residue** | **Protein residue** | **Bond Length(A)** | **Score** |
| 1 | A0A0B7MF59m | Arg(43) | Gln(500) | 2.36 | -5.91 |
| Arg(43) | Thr(501) | 3.16 |
| 2 | A0A0B7MF59w | Glu(18) | Gly(515) | 2.17 | -8.34 |
| 3 | A0A1S9ZCG0 | Tyr(36) | Asp(454) | 2.59 | -13.33 |
| Gly(21) | Trp(493) | 2.78 |
| Ile(19) | Arg(512) | 2.59 |
| 4 | A0A1S9ZCG0m | Tyr(36) | Asp(454) | 2.59 | -13.01 |
| Gly(21) | Trp(493) | 2.78 |
| Ile(19) | Arg(512) | 2.59 |
| 5 | A0A6I1TUZ4m | No interaction |
| 6 | B3A0N4 | Gly(38) | Thr(501) | 1.94 | -8.88 |
| Asn(42) | Asp(454) | 2.85 |
| 7 | B3A0N4m | Cyx(9) | Ser(542) | 2.32 | -13.67 |
| Cyx(14) | Ser(542) | 2.85 |
| 8 | C0HJC0 | Thr(15) | Tyr(510) | 1.73 | -9.79 |
| Leu(11) | Tyr(510) | 2.74 |
| Phe(4) | Trp(493) | 3.01 |
| Lys(28) | Tyr(499) | 2.96 |
| Asn(6) | Tyr(462) | 2.48 |
| 9 | C0HL39m | Glu(27) | Arg(512) | 2.90 | -2.66 |
| 10 | C1CBU6 | Glu(18) | Gly(515) | 2.17 | -3.20 |
| 11 | J1P1V9m | Glu(18) | Gly(515) | 2.17 | -6.79 |
| 12 | O07623m | Leu(41) | Leu(451) | 2.59 | -13.55 |
| 13 | P01547 | Thr(27) | Thr(434) | 3.32 | -1.56 |
| Ser(9) | Trp(491) | 2.53 |
| 14 | P01547m | No interaction |
| 15 | P36502 | Ser(4) | Asp(454) | 2.30 | -9.42 |
| Lys(19) | Thr(391) | 3.17 |
| 16 | P36502m | Thr(11) | Trp(506) | 2.40 | -8.61 |
| 17 | P36503 | Thr(18) | Asp(454) | 3.08 | -1.04 |
| 18 | P36503m | Ser(6) | Tyr(510) | 2.32 | -3.79 |
| 19 | P36504 | Gln(3) | Lys(588) | 5.16 | -13.55 |
| Ser(6) | Ser(28) | 2.23 |
| Thr(11) | Ile(24) | 2.38 |
| Phe(7) | Ser(543) | 3.20 |
| Phe(7) | Pro(29) | 3.08 |
| 20 | P36504m | Leu(11) | Trp(506) | 2.33 | -15.79 |
| Cyx(5) | Tyr(510) | 3.04 |
| Asp(15) | Ser(461) | 3.12 |
| 21 | P36961 | Thr(2) | Trp(557) | 3.05 | -9.09 |
| Val(17) | Tyr(499) | 2.74 |
| 22 | P36961m | Asp(26) | Ser(146) | 3.08 | -9.56 |
| 23 | P42723 | Ser (27) | Leu (444) | 3.18 | -4.91 |
| 24 | P80214 | No interaction |
| 25 | P80214m | Lys(44) | Ala(440) | 3.17 | -7.99 |
| Lys(40) | Met(395) | 2.39 |
| Lys(40) | Ser(389) | 2.52 |
| 26 | P80666 | Phe(6) | Glu(550) | 2.38 | -9.34 |
| Lys(13) | Ile(24) | 2.07 |
| 27 | P80666m | No interaction |
| 28 | P80925 | No interaction |
| 29 | P80925m | Asn(10) | Tyr(499) | 2.31 | -11.76 |
| Asn(10) | Thr(501) | 3.24 |
| Lys(11) | Phe(504) | 2.78 |
| Lys(1) | Asn(505) | 3.00 |
| 30 | P80959 | No interaction |
| 31 | P80959m | No interaction |
| 32 | P81053 | Asp(17) | Ser(542) | 2.82 | -15.70 |
| 33 | P81053m | His(42) | Tyr(388) | 2.12 | -15.45 |
| Asn(43) | Tyr(388) | 2.65 |
| Lys(12) | His(309) | 3.32 |
| 34 | P83002 | Tyr(3) | Asn(511) | 2.52 | -6.03 |
| Tyr(3) | Asp(494) | 2.83 |
| 35 | P83002m | Glu(22) | Arg(512) | 2.74 | -7.90 |
| 36 | P83378 | No interaction |
| 37 | P83378m | Lys(7) | Asp(454) | 2.55 | -10.70 |
| Ser(5) | Asp(454) | 2.94 |
| 38 | P84886 | Tyr(2) | Trp(493) | 2.35 | -6.80 |
| 39 | P84886m | No interaction |
| 40 | P84962 | Asp(18) | Lys(502) | 2.75 | -9.36 |
| Asp(18) | Tyr(499) | 2.11 |
| 41 | P84962m | Trp(19) | Tyr(510) | 2.56 | -8.49 |
| 42 | P85065m | Ser(13) | Asn(27) | 3.24 | -14.03 |
| 43 | P85148 | Lys(37) | Thr(480) | 2.55 | -10.69 |
| Lys(37) | Met(473) | 3.19 |
| Val(21) | Tyr(474) | 2.79 |
| Cys(36) | Tyr(474) | 2.56 |
| 44 | P85148m | Asn(26) | Arg(512) | 3.00 | -8.09 |
| Cys(31) | Trp(493) | 2.57 |
| 45 | P85876 | Lys(23) | Tyr(510) | 2.14 | -14.99 |
| Ala(27) | Tyr(510) | 3.23 |
| Glu(24) | Ser(496) | 2.54 |
| Thr(2) | Asp(454) | 3.09 |
| 46 | P85876m | No interaction |
| 47 | P86291 | Tyr(3) | Asp(494) | 2.83 | -7.88 |
| Tyr(3) | Asn(511) | 2.52 |
| 48 | P86291m | Tyr(3) | Asp(494) | 2.83 | -2.49 |
| Tyr(3) | Asn(511) | 2.52 |
| 49 | P86393 | Tyr(16) | Leu(451) | 2.77 | -9.01 |
| Thr(17) | Thr(501) | 2.77 |
| 50 | P86393m | No interaction |
| 51 | P86394 | Gly(35) | Leu(451) | 2.86 | -7.12 |
| 52 | P86394m | Thr(29) | Glu(25) | 2.46 | -5.01 |
| Thr(29) | Ile(24) | 2.78 |
| Ala(33) | Thr(22) | 3.21 |
| Asn(5) | Arg(590) | 3.13 |
| 53 | S7Z987 | Tyr(36) | Asp(454) | 2.59 | -9.30 |
| Gly(21) | Trp(493) | 2.73 |
| Ile(19) | Arg(519) | 2.59 |
| 54 | S7Z987m | Lys(20) | Ser(455) | 2.11 | -4.60 |
| 55 | V8IIJ8m | Arg(43) | Thr(501) | 3.16 | -14.89 |
| Arg(43) | Gln(500) | 2.36 |

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