

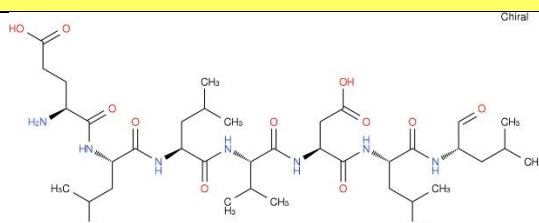
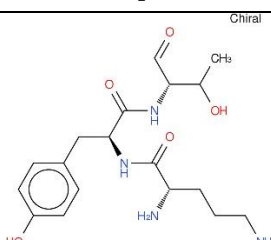
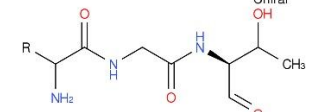
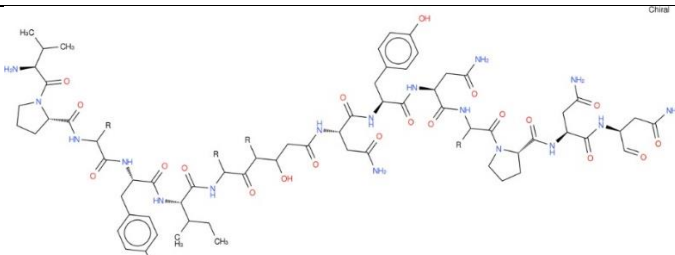
Supplementary Table S1. Details of *Bacillus halotolerans* isolates used in this study.

| Species | Strain | Genome size (Mb) | Plasmid | Description | GenBank Accession |
|------------------------------|--------------|------------------|---------|--|-------------------|
| <i>Bacillus halotolerans</i> | FJAT-2398 | 4.09077 | - | Isolated from agricultural soils (Germany) | LUUT00000000.1 |
| | ATCC 25096 | 4.07273 | - | Isolated from desert soils (Morocco) | LPVF00000000.1 |
| | NRRL B-41617 | 4.09146 | - | Isolated from river (Spain) | LPVD00000000.1 |
| | III-1 | 4.13992 | - | Isolated from roots of <i>Phoenix dactylifera</i> (Tunisia) | MBQV00000000.1 |
| | BFOA2 | 4.44148 | - | Isolated from roots of <i>Limoniastrum monopetalum</i> (Tunisia) | PVXB00000000.1 |
| | BFOA4 | 4.13265 | - | Isolated from rhizosphere of wheat plant (Algeria) | PVXD00000000.1 |
| | BFOA1 | 4.09488 | - | Isolated from roots of <i>Limoniastrum monopetalum</i> (Tunisia) | PVXA00000000.1 |
| | BFOA3 | 4.15264 | - | Isolated from roots of <i>Limoniastrum monopetalum</i> (Tunisia) | PVXC00000000.1 |
| | NRRL B-41618 | 4.20528 | - | Isolated from river (Spain) | LPVE00000000.1 |
| | RSCu-8D | 4.15979 | - | Isolated from <i>Cuminum cyminum</i> (India) | NERK00000000.1 |
| | LNXM37 | 4.08166 | - | Isolated from soil (environmental sample, China) | PVQP00000000.1 |
| | LNXM78 | 4.17021 | - | Isolated from soil (environmental sample, China) | PVQQ00000000.1 |
| | DGL6 | 3.73909 | - | Isolated from a salt lake (China) | PVWC00000000.1 |

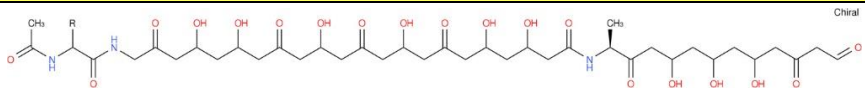
Supplementary Table S2. Details of fungal species used in this study.

| | Species | Strain | Source | Country | GenBank Accession |
|----|--|--------|----------------------------|---------|----------------------|
| | | | | | ITS |
| 1 | <i>Fusarium solani</i> | Fso3 | <i>Olea europaea</i> | Tunisia | KU528851 |
| 2 | <i>Fusarium solani</i> | Fso12 | <i>Olea europaea</i> | Tunisia | KU528862 |
| 3 | <i>Fusarium solani</i> | Fso11 | <i>Olea europaea</i> | Tunisia | KU528861 |
| 4 | <i>Fusarium solani</i> | Fso7 | <i>Olea europaea</i> | Tunisia | KU528857 |
| 5 | <i>Fusarium solani</i> | Fso5 | <i>Olea europaea</i> | Tunisia | KU528854 |
| 6 | <i>Fusarium solani</i> | Fso6 | <i>Olea europaea</i> | Tunisia | KU528855 |
| 7 | <i>Fusarium solani</i> | Fso8 | <i>Olea europaea</i> | Tunisia | KU528858 |
| 8 | <i>Fusarium solani</i> | Fso9 | <i>Olea europaea</i> | Tunisia | KU528859 |
| 9 | <i>Fusarium solani</i> | Fso1 | <i>Olea europaea</i> | Tunisia | KU528848 |
| 10 | <i>Fusarium solani</i> | Fso10 | <i>Olea europaea</i> | Tunisia | KU528860 |
| 11 | <i>Fusarium solani</i> | Fso13 | <i>Olea europaea</i> | Tunisia | KU528863 |
| 12 | <i>Fusarium solani</i> | Fso2 | <i>Olea europaea</i> | Tunisia | KU528850 |
| 13 | <i>Fusarium oxysporum</i> | Fox1 | <i>Olea europaea</i> | Tunisia | KU528844 |
| 14 | <i>Fusarium oxysporum</i> f.sp. <i>radices lycopersicum</i> | FORL | Tomato | Tunisia | - |
| 15 | <i>Fusarium acuminatum</i> | Fac | <i>Olea europaea</i> | Tunisia | KU528866 |
| 16 | <i>Fusarium chlamydosporum</i> | Fch1 | <i>Olea europaea</i> | Tunisia | KU528845 |
| 17 | <i>Alternaria</i> sp. | KT1 | <i>Pistacia vera</i> | Tunisia | - |
| 18 | <i>Rhizoctonia bataticola</i> | MAT1 | <i>Olea europaea</i> | Tunisia | - |
| 19 | <i>Phytophthora infestans</i> | LMA1 | Potato | Algeria | - |
| 20 | <i>Alternaria alternata</i> | LMA2 | Tomato | Algeria | - |
| 21 | <i>Botrytis cinerea</i> | LMA3 | Strawberry | Algeria | - |
| 22 | <i>Fusarium oxysporum</i> f.sp. <i>albedinis</i> | LMA1 | <i>Phoenix dactylifera</i> | Algeria | - |
| 23 | <i>Fusarium oxysporum</i> f.sp. <i>albedinis</i> | LMA2 | <i>Phoenix dactylifera</i> | Algeria | - |
| 24 | <i>Fusarium oxysporum</i> f.sp. <i>albedinis</i> | LMA3 | <i>Phoenix dactylifera</i> | Algeria | - |
| 25 | <i>Fusarium oxysporum</i> f.sp. <i>albedinis</i> | LMA4 | <i>Phoenix dactylifera</i> | Algeria | - |
| | <i>Fusarium oxysporum</i> f.sp. <i>albedinis</i> | LMA5 | <i>Phoenix dactylifera</i> | Algeria | - |
| 23 | <i>Fusarium solani</i> var. <i>coeruleum</i> | IPP1 | Potato | France | - |

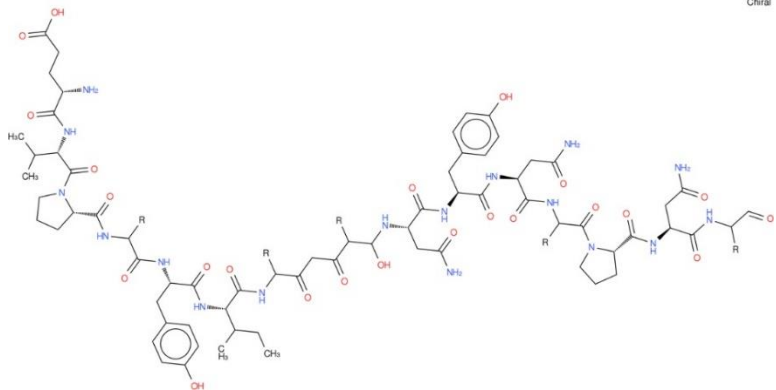
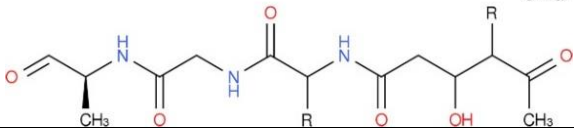
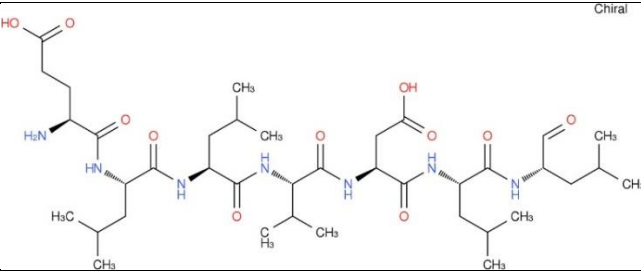
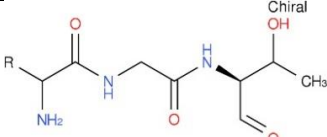
Supplementary Table S3. Secondary metabolites of *Bacillus halotolerans* isolates identified with antiSMASH program.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-------------------|---------------------------|-------------|---|--|
| ATCC 25096 | | | | |
| Cluster 1 | Nrps | 65255 | Surfactin biosynthetic gene cluster (86% of genes show similarity) |  Chiral |
| Cluster 2 | Nrps | 2695 | - | - |
| Cluster 3 | Nrps | 12795 | Plipastatin biosynthetic gene cluster (38% of genes show similarity) |  Chiral |
| Cluster 4 | Nrps | 49738 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  Chiral |
| Cluster 5 | Sactipeptide-Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 6 | Other | 41422 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 7 | Nrps | 90306 | Fengycin biosynthetic gene cluster (80% of genes show similarity) |  Chiral |

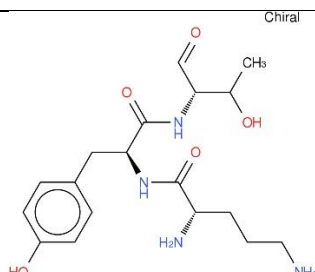
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-------------------|-----------------------------|-------------|--|---|
| ATCC 25096 | | | | |
| Cluster 8 | Transatpks- Otherks-Nrps | 110097 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 9 | Terpene | 20806 | - | - |
| Cluster 10 | Terpene | 21898 | - | - |
| Cluster 11 | T3pks | 41097 | - | - |
| Cluster 12 | Lantipeptide | 18699 | - | - |

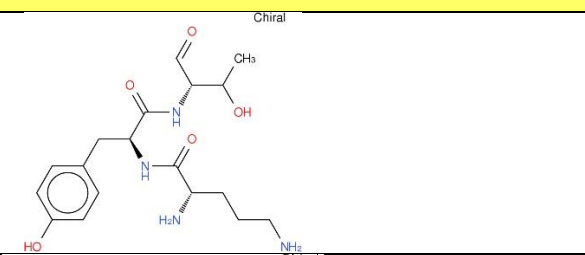
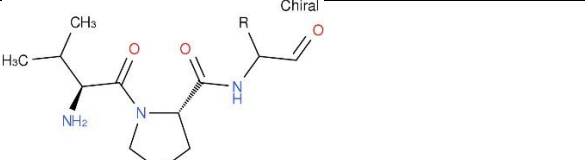

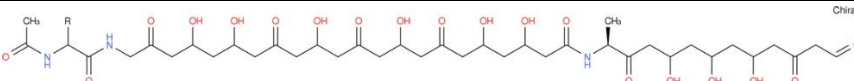
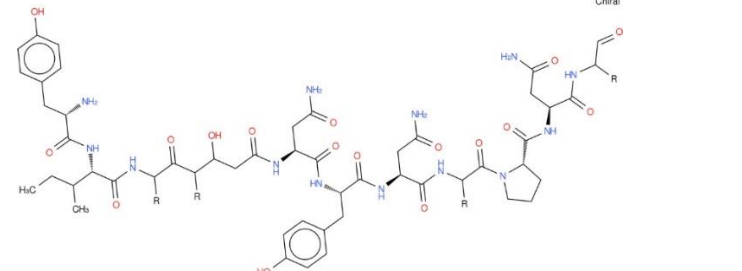
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|------------------|-------------------------|-------------|---|---|
| FJAT-2398 | | | | |
| Cluster 1 | T3pks | 41097 | - | - |
| Cluster 2 | Terpene | 21898 | - | - |
| Cluster 3 | Transatpks-Nrps | 114372 | Fengycin biosynthetic gene cluster (93% of genes show similarity) |  |
| Cluster 4 | Transatpks-Otherks-Nrps | 110096 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 5 | Terpene | 20806 | - | - |
| Cluster 6 | Nrps | 65255 | Surfactin biosynthetic gene cluster (86% of genes show similarity) |  |
| Cluster 7 | Lantipeptide | 18464 | - | - |
| Cluster 8 | Nrps | 49738 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  |

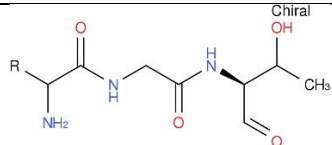
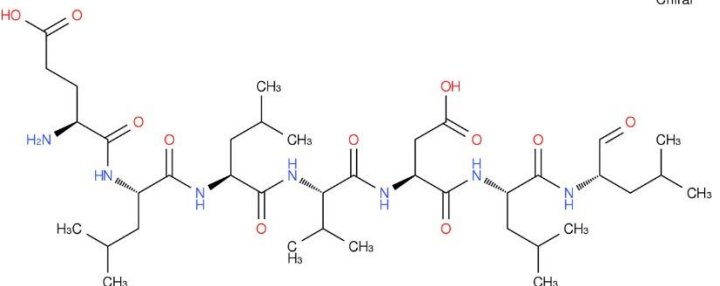
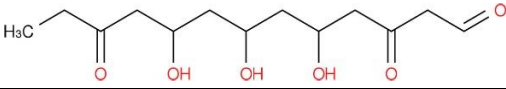
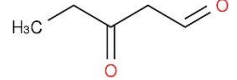
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|------------------|-------------------------------|-------------|--|---|
| FJAT-2398 | | | | |
| Cluster 9 | Sactipeptide- Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 10 | Other | 41422 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 11 | Nrps | 12929 | Plipastatin biosynthetic gene cluster (38% of genes show similarity) |  Chiral |

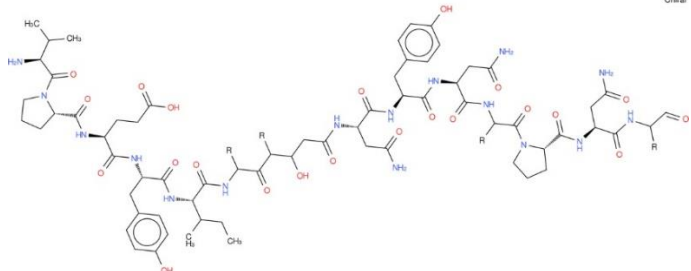
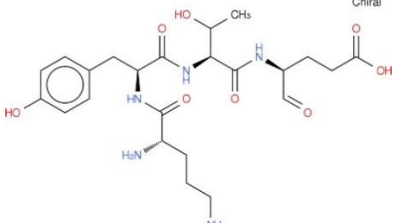
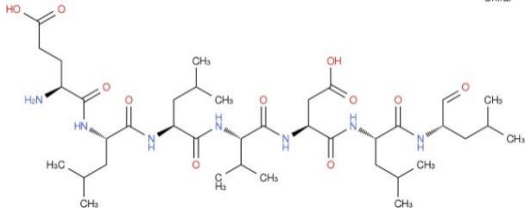
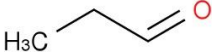
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-----------------|---------------------------------|-------------|--|---|
| III-1 | | | | |
| Cluster 1 | Nrps | 14107 | Fengycin biosynthetic gene cluster (26% of genes show similarity) |  |
| Cluster 2 | Nrps | 10652 | Plipastatin biosynthetic gene cluster (23% of genes show similarity) |  |
| Cluster 3 | Transatpks | 8404 | Elansolid biosynthetic gene cluster (10% of genes show similarity) |  |
| Cluster 4 | Transatpks | 5353 | - | - |
| Cluster 5 | Transatpks | 2123 | - | - |
| Cluster 6 | Terpene | 20806 | - | - |
| Cluster 7 | Nrps- Transatpks- Otherks | 110082 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 8 | Transatpks- Nrps | 80362 | Fengycin biosynthetic gene cluster (86% of genes show similarity) |  |
| Cluster 9 | Other | 41418 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |

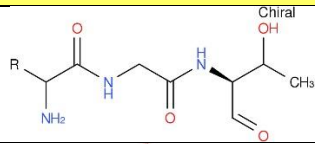
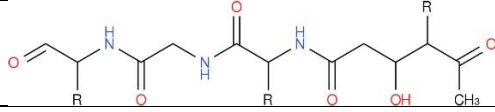
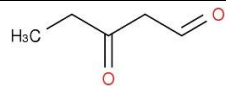
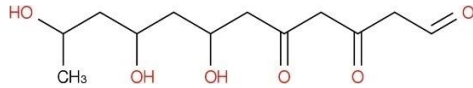
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-----------------|----------------------------------|-------------|---|---|
| III-1 | | | | |
| Cluster 10 | Sactipeptide- Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 11 | Nrps | 49738 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 12 | T3pks | 41097 | - | - |
| Cluster 13 | Terpene | 21898 | - | - |
| Cluster 14 | Nrps | 23006 | Plipastatin biosynthetic gene cluster (38% of genes show similarity) | - |
| Cluster 15 | Nrps | 65394 | Surfactin biosynthetic gene cluster (86% of genes show similarity) |  |
| Cluster 16 | Transatpks | 48051 | Phormidolide biosynthetic gene cluster (17% of genes show similarity) |  |
| Cluster 17 | Transatpks- T1pks- Otherks | 32386 | Phormidolide biosynthetic gene cluster (21% of genes show similarity) |  |

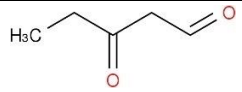
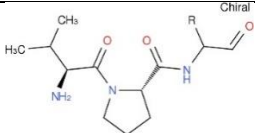
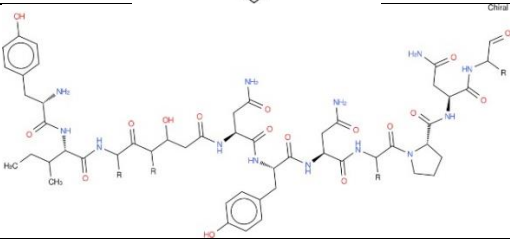
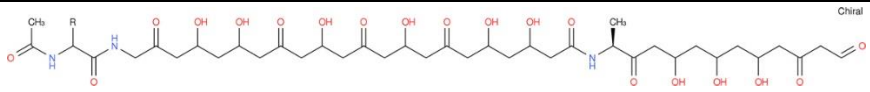
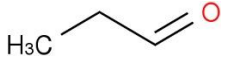
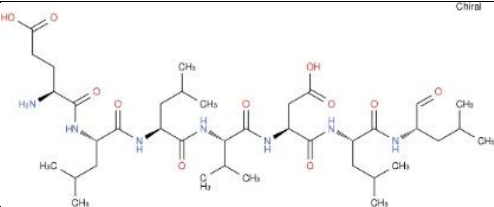
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|---------------------|---------------------------|-------------|--|---|
| NRRL B-41618 | | | | |
| Cluster 1 | Transatpks-Nrps | 90713 | Fengycin biosynthetic gene cluster (80% of genes show similarity) |  |
| Cluster 2 | Transatpks | 5608 | - | - |
| Cluster 3 | Nrps | 14364 | Plipastatin biosynthetic gene cluster (30% of genes show similarity) |  |
| Cluster 4 | Nrps | 65395 | Surfactin biosynthetic gene cluster (86% of genes show similarity) |  |
| Cluster 5 | Transatpks | 8190 | Elansolid biosynthetic gene cluster (10% of genes show similarity) |  |
| Cluster 6 | Nrps | 1122 | - | - |
| Cluster 7 | Other | 41418 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 8 | Sactipeptide-Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |

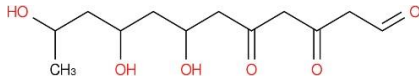
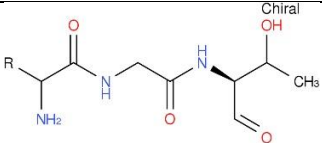
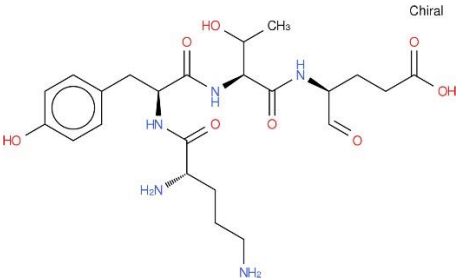
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|---------------------|----------------------------------|-------------|---|---|
| NRRL B-41618 | | | | |
| Cluster 9 | Nrps | 49738 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 10 | Transatpks- Otherks-Nrps | 110097 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 11 | Terpene | 20806 | - | - |
| Cluster 12 | Nrps | 22596 | Plipastatin biosynthetic gene cluster (38% of genes show similarity) | - |
| Cluster 13 | Terpene | 21898 | - | - |
| Cluster 14 | Transatpks- T1pks- Otherks | 32280 | Thiomarinol biosynthetic gene cluster (8% of genes show similarity) |  |
| Cluster 15 | T3pks | 41097 | - | - |
| Cluster 16 | Transatpks | 46509 | Myxovirescin biosynthetic gene cluster (13% of genes show similarity) |  |

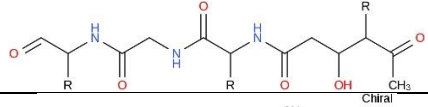
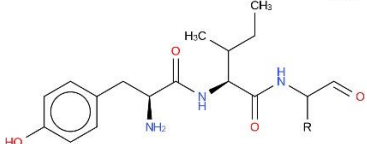
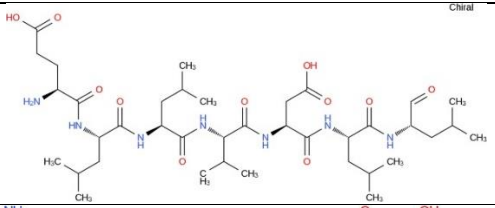
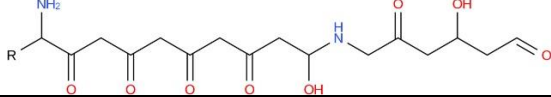
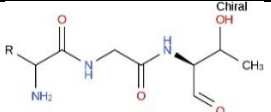
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|---------------------|--------------------------|-------------|---|---|
| NRRL B-41617 | | | | |
| Cluster 1 | T3pks | 41097 | - | - |
| Cluster 2 | Terpene | 21898 | - | - |
| Cluster 3 | Nrps | 22905 | Plipastatin biosynthetic gene cluster (38% of genes show similarity) | - |
| Cluster 4 | Transatpks-T1pks-Otherks | 32259 | Phormidolide biosynthetic gene cluster (21% of genes show similarity) |  |
| Cluster 5 | Nrps | 10181 | Plipastatin biosynthetic gene cluster (23% of genes show similarity) |  |
| Cluster 6 | Transatpks-Nrps | 63211 | Fengycin biosynthetic gene cluster (80% of genes show similarity) |  |
| Cluster 7 | Terpene | 20806 | - | - |
| Cluster 8 | Nrps-Transatpks-Otherks | 110093 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 9 | Transatpks | 8405 | Elansolid biosynthetic gene cluster (10% of genes show similarity) |  |
| Cluster 10 | Transatpks | 2083 | - | - |
| Cluster 11 | Nrps | 65394 | Surfactin biosynthetic gene cluster (86% of genes show similarity) |  |

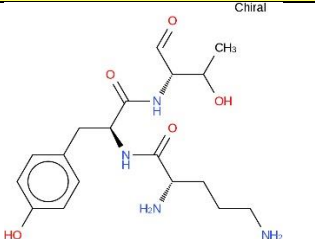

Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|---------------------|---------------------------|-------------|---|--|
| NRRL B-41617 | | | | |
| Cluster 12 | Other | 41418 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 13 | Transatpks | 46517 | Myxovirescin biosynthetic gene cluster (13% of genes show similarity) |  |
| Cluster 14 | Nrps | 49738 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 15 | Sactipeptide-Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 16 | Nrps | 15038 | Plipastatin biosynthetic gene cluster (30% of genes show similarity) |  |

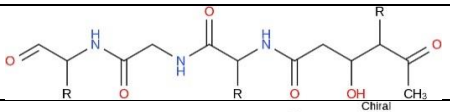
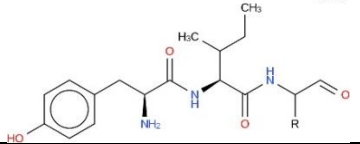
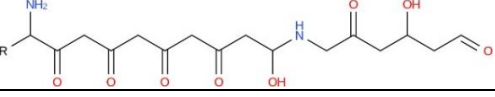
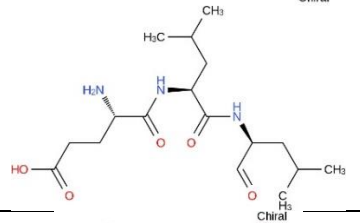
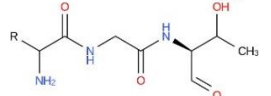
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-------------------|---------------------------------|-------------|---|---|
| BFOA2 (72) | | | | |
| Cluster 1 | Nrps- Transatpks- Otherks | 110091 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 2 | Nrps | 33415 | Fengycin biosynthetic gene cluster (80% of genes show similarity) |  |
| Cluster 3 | Sactipeptide- Head_to_tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 4 | Other | 41418 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 5 | Bacteriocin | 12183 | - | - |
| Cluster 6 | T3pks | 41097 | - | - |
| Cluster 7 | Terpene | 21898 | - | - |
| Cluster 8 | Nrps | 65395 | Surfactin biosynthetic gene cluster (86% of genes show similarity) |  |
| Cluster 9 | Transatpks- Nrps | 61278 | Kijanamicin biosynthetic gene cluster (4% of genes show similarity) |  |
| Cluster 10 | Terpene | 20806 | - | - |
| Cluster 11 | Nrps | 43656 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 12 | Nrps | 23268 | Plipastatin biosynthetic gene cluster (46% of genes show similarity) | - |

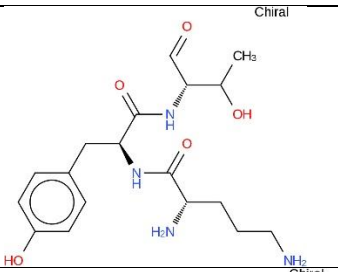
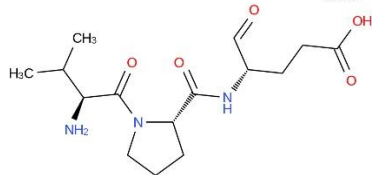

Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-------------------|------|-------------|--|---|
| BFOA2 (72) | | | | |
| Cluster 13 | Nrps | 13200 | Plipastatin biosynthetic gene cluster (38% of genes show similarity) |  |
| Cluster 14 | Nrps | 10296 | Plipastatin biosynthetic gene cluster (15% of genes show similarity) |  |

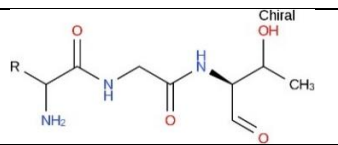
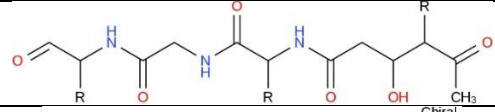
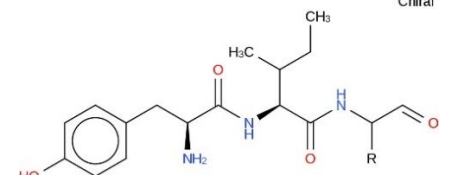
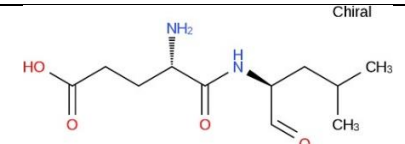
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-------------------|---------------------------------|-------------|---|---|
| BFOA1 (86) | | | | |
| Cluster 1 | Terpene | 20806 | - | - |
| Cluster 2 | Nrps- Transatpks- Otherks | 110091 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 3 | Nrps | 33415 | Fengycin biosynthetic gene cluster (80% of genes show similarity) |  |
| Cluster 4 | Sactipeptide- Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 5 | Other | 41418 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 6 | Bacteriocin | 12183 | - | - |
| Cluster 7 | Transatpks- Nrps | 61278 | Kijanamicin biosynthetic gene cluster (4% of genes show similarity) |  |
| Cluster 8 | T3pks | 41097 | - | - |
| Cluster 9 | Terpene | 21898 | - | - |
| Cluster 10 | Nrps | 25324 | Surfactin biosynthetic gene cluster (43% of genes show similarity) | - |
| Cluster 11 | Nrps | 28547 | Surfactin biosynthetic gene cluster (47% of genes show similarity) |  |
| Cluster 12 | Nrps | 43656 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  |

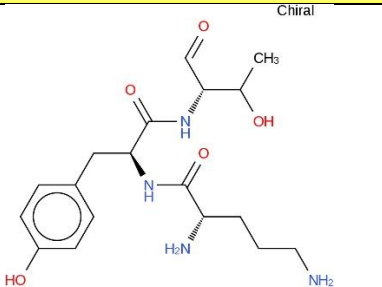
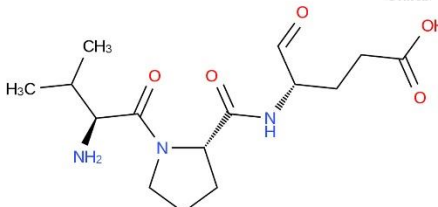
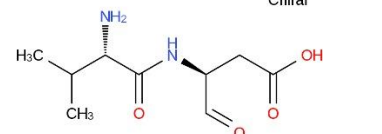
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-------------------|-------|-------------|--|---|
| BFOA1 (86) | | | | |
| Cluster 13 | Nrps | 23268 | Plipastatin biosynthetic gene cluster (46% of genes show similarity) | - |
| Cluster 14 | Nrps | 13615 | Fengycin biosynthetic gene cluster (26% of genes show similarity) |  |
| Cluster 15 | Nrps | 10296 | Plipastatin biosynthetic gene cluster (15% of genes show similarity) |  |
| Cluster 16 | Nrps | 9326 | Surfactin biosynthetic gene cluster (8% of genes show similarity) |  |
| Cluster 17 | Other | 1009 | - | - |

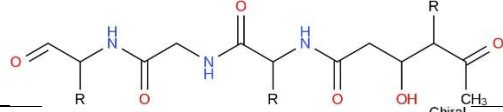
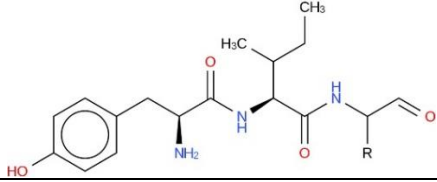
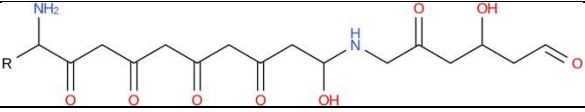
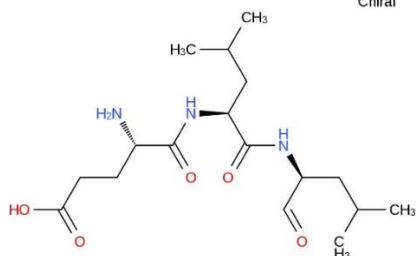
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-----------------|---------------------------|-------------|---|---|
| LNXM37 | | | | |
| Cluster 1 | Other | 41418 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 2 | Sactipeptide-Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 3 | Nrps | 49738 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 4 | Nrps-Transatpks-Otherks | 110098 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 5 | Nrps | 33354 | Fengycin biosynthetic gene cluster (80% of genes show similarity) |  |
| Cluster 6 | T3pks | 41097 | - | - |
| Cluster 7 | Terpene | 21898 | - | - |
| Cluster 8 | Nrps | 26072 | Surfactin biosynthetic gene cluster (43% of genes show similarity) | - |
| Cluster 9 | Terpene | 20806 | - | - |
| Cluster 10 | Lantipeptide | 22597 | - | - |
| Cluster 11 | Nrps | 21754 | Plipastatin biosynthetic gene cluster (30% of genes show similarity) | - |
| Cluster 12 | Nrps | 28299 | Surfactin biosynthetic gene cluster (47% of genes show similarity) |  |

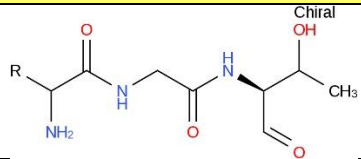
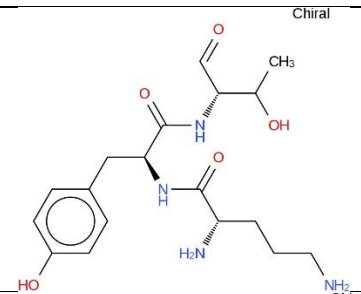
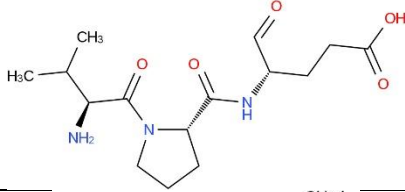
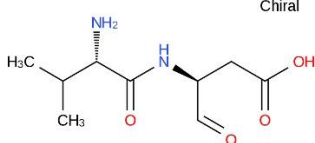
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-----------------|------|-------------|--|---|
| LNXM37 | | | | |
| Cluster 13 | Nrps | 13179 | Plipastatin biosynthetic gene cluster (23% of genes show similarity) |  |
| Cluster 14 | Nrps | 9879 | Plipastatin biosynthetic gene cluster (15% of genes show similarity) |  |
| Cluster 15 | Nrps | 9291 | Surfactin biosynthetic gene cluster (8% of genes show similarity) |  |

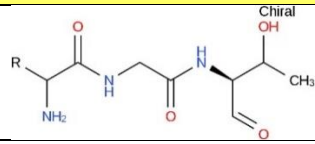
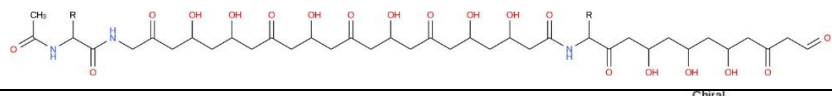
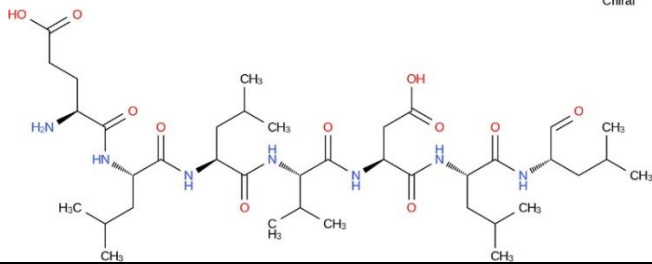
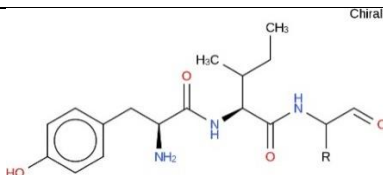
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-------------------|---------------------------------|-------------|--|---|
| BFOA3 (36) | | | | |
| Cluster 1 | Nrps- Transatpks- Otherks | 110091 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 2 | Nrps | 33415 | Fengycin biosynthetic gene cluster (80% of genes show similarity) |  |
| Cluster 3 | Sactipeptide- Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 4 | Other | 41418 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 5 | Bacteriocin | 12183 | - | - |
| Cluster 6 | Transatpks- Nrps | 61278 | Kijanamicin biosynthetic gene cluster (4% of genes show similarity) |  |
| Cluster 7 | T3pks | 41097 | - | - |
| Cluster 8 | Terpene | 21898 | - | - |
| Cluster 9 | Terpene | 20806 | - | - |
| Cluster 10 | Nrps | 25324 | Surfactin biosynthetic gene cluster (43% of genes show similarity) | - |
| Cluster 11 | Nrps | 28547 | Surfactin biosynthetic gene cluster (47% of genes show similarity) |  |

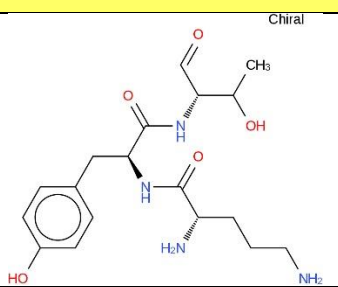
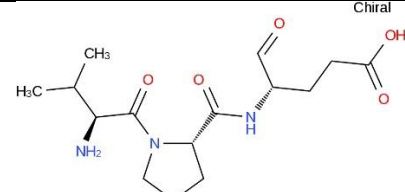
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-------------------|-------|-------------|---|---|
| BFOA3 (36) | | | | |
| Cluster 12 | Nrps | 43656 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 13 | Nrps | 23268 | Plipastatin biosynthetic gene cluster (46% of genes show similarity) | - |
| Cluster 14 | Nrps | 13200 | Plipastatin biosynthetic gene cluster (38% of genes show similarity) |  |
| Cluster 15 | Nrps | 10296 | Plipastatin biosynthetic gene cluster (15% of genes show similarity) |  |
| Cluster 16 | Nrps | 9326 | Surfactin biosynthetic gene cluster (8% of genes show similarity) |  |
| Cluster 17 | Other | 1009 | - | - |

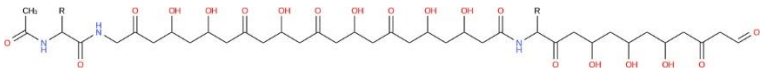
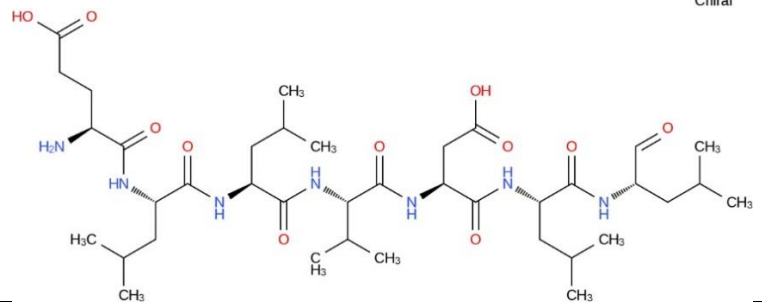
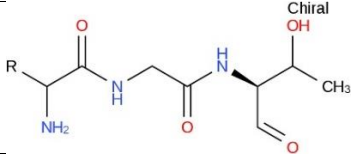
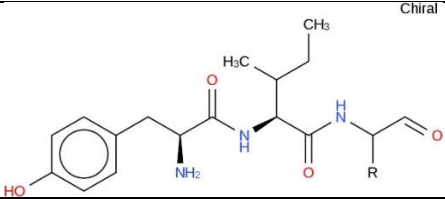
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-----------------|---------------------------|-------------|---|---|
| LNXM78 | | | | |
| Cluster 1 | Nrps | 49738 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 2 | Sactipeptide-Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 3 | Other | 41418 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 4 | T3pks | 41097 | - | - |
| Cluster 5 | Terpene | 20806 | - | - |
| Cluster 6 | Nrps-Transatpks-Otherks | 110098 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 7 | Nrps | 65395 | Surfactin biosynthetic gene cluster (86% of genes show similarity) |  |
| Cluster 8 | Terpene | 21898 | - | - |
| Cluster 9 | Lantipeptide | 21755 | - | - |
| Cluster 10 | Nrps | 33542 | Fengycin biosynthetic gene cluster (80% of genes show similarity) |  |
| Cluster 11 | Nrps | 21938 | Plipastatin biosynthetic gene cluster (30% of genes show similarity) | - |

Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-----------------|-------|-------------|--|---|
| LNXM78 | | | | |
| Cluster 12 | Nrps | 13531 | Plipastatin biosynthetic gene cluster (23% of genes show similarity) |  |
| Cluster 13 | Nrps | 10317 | Plipastatin biosynthetic gene cluster (23% of genes show similarity) |  |
| Cluster 14 | Other | 1052 | - | - |
| Cluster 15 | Other | 1052 | - | - |

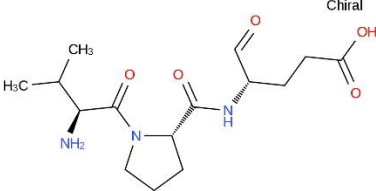
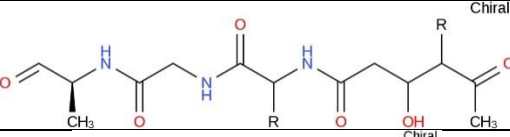
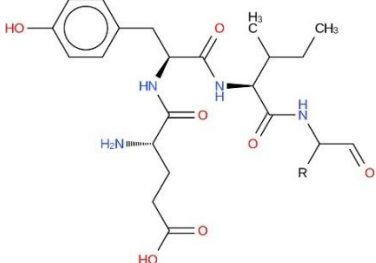
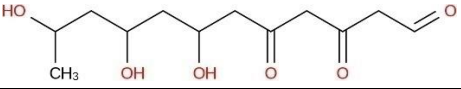
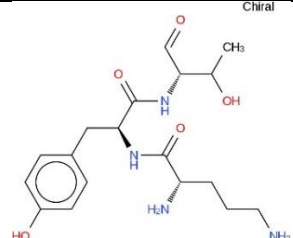
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|--------------------|---------------------------|-------------|---|--|
| BFOA4 (B19) | | | | |
| Cluster 1 | Terpene | 20806 | - | - |
| Cluster 2 | Other | 41418 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 3 | Sactipeptide-Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 4 | Transatpks-Otherks-Nrps | 105327 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 5 | Lantipeptide | 22597 | - | - |
| Cluster 6 | Nrps | 65395 | Surfactin biosynthetic gene cluster (86% of genes show similarity) |  Chiral |
| Cluster 7 | Nrps | 49738 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  Chiral |
| Cluster 8 | T3pks | 41097 | - | - |
| Cluster 9 | Nrps | 33485 | Fengycin biosynthetic gene cluster (80% of genes show similarity) |  Chiral |
| Cluster 10 | Terpene | 21898 | - | - |

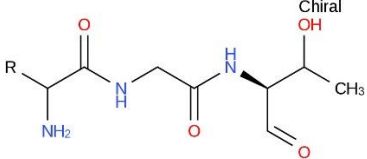
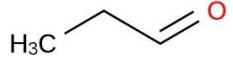
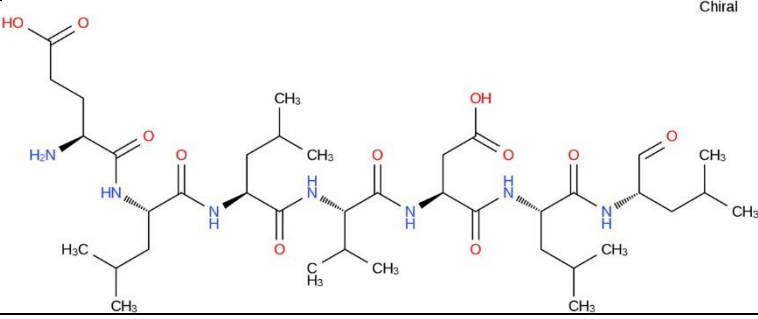
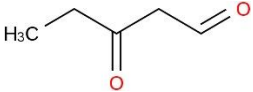
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|--------------------|------|-------------|--|-------------------------|
| BFOA4 (B19) | | | | |
| Cluster 11 | Nrps | 13174 | Plipastatin biosynthetic gene cluster (30% of genes show similarity) | |
| Cluster 12 | Nrps | 9618 | Plipastatin biosynthetic gene cluster (23% of genes show similarity) | |
| Cluster 13 | Nrps | 5050 | Plipastatin biosynthetic gene cluster (23% of genes show similarity) | - |

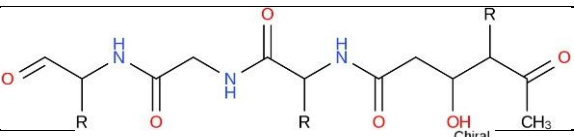
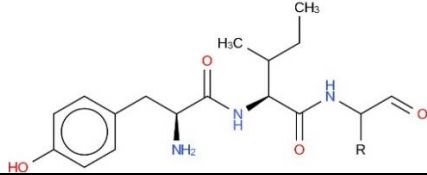
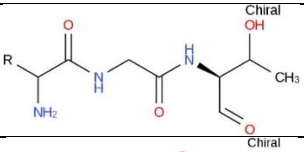
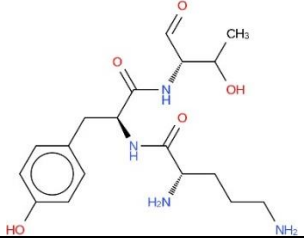
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-----------------|---------------------------------|-------------|---|---|
| RSCu-8D | | | | |
| Cluster 1 | Nrps | 9233 | Plipastatin biosynthetic gene cluster (15% of genes show similarity) |  |
| Cluster 2 | Terpene | 20806 | - | - |
| Cluster 3 | Nrps- Transatpks- Otherks | 110097 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 4 | Nrps | 57204 | Fengycin biosynthetic gene cluster (93% of genes show similarity) |  |
| Cluster 5 | Terpene | 21898 | - | - |
| Cluster 6 | T3pks | 41097 | - | - |
| Cluster 7 | Transatpks | 46339 | Myxovirescin biosynthetic gene cluster (13% of genes show similarity) |  |
| Cluster 8 | Transatpks | 5388 | - | - |
| Cluster 9 | Nrps | 12848 | Plipastatin biosynthetic gene cluster (38% of genes show similarity) |  |

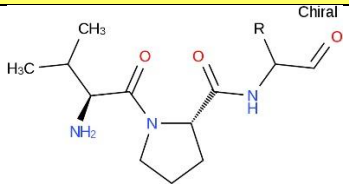
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-----------------|---------------------------|-------------|---|---|
| RSCu-8D | | | | |
| Cluster 10 | Nrps | 49738 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 11 | Transatpks | 8471 | Elansolid biosynthetic gene cluster (10% of genes show similarity) |  |
| Cluster 12 | Nrps | 65394 | Surfactin biosynthetic gene cluster (86% of genes show similarity) |  |
| Cluster 13 | Sactipeptide-Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 14 | Other | 41418 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 15 | Transatpks-T1pks-Otherks | 32875 | Thiomarinol biosynthetic gene cluster (8% of genes show similarity) |  |

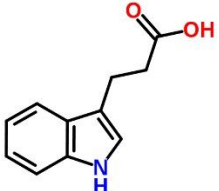
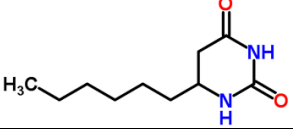
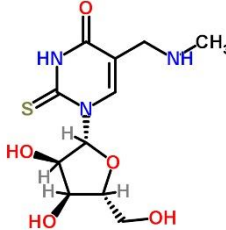
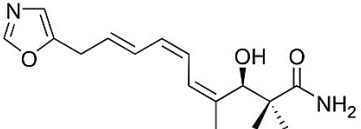

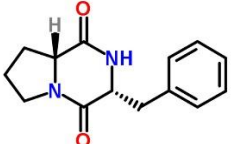
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-----------------|---------------------------------|-------------|---|---|
| DGL6 | | | | |
| Cluster 1 | T3pks | 41097 | - | - |
| Cluster 2 | Terpene | 21898 | - | - |
| Cluster 3 | Nrps | 21680 | Plipastatin biosynthetic gene cluster (30% of genes show similarity) | - |
| Cluster 4 | Terpene | 20806 | - | - |
| Cluster 5 | Nrps- Transatpks- Otherks | 110103 | Bacillaene biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 6 | Nrps | 33899 | Fengycin biosynthetic gene cluster (86% of genes show similarity) |  |
| Cluster 7 | Sactipeptide- Head to tail | 21612 | Subtilosin A biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 8 | Other | 41418 | Bacilysin biosynthetic gene cluster (100% of genes show similarity) | - |
| Cluster 9 | Nrps | 49738 | Bacillibactin biosynthetic gene cluster (100% of genes show similarity) |  |
| Cluster 10 | Nrps | 14513 | Plipastatin biosynthetic gene cluster (30% of genes show similarity) |  |

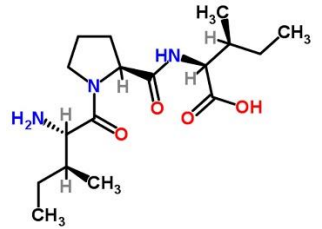
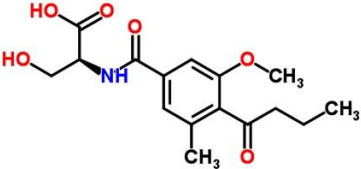
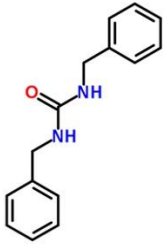
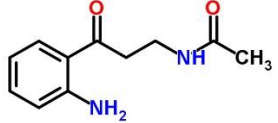
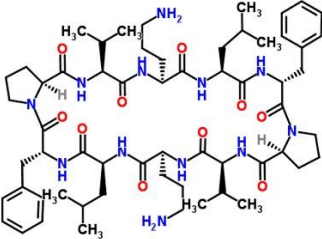
Supplementary Table S3. Continued.

| Strain clusters | Type | Length (bp) | Most similar known clusters | Predicted core clusters |
|-----------------|------|-------------|--|--|
| DGL6 | | | | |
| Cluster 11 | Nrps | 11005 | Plipastatin biosynthetic gene cluster (23% of genes show similarity) |  The chemical structure shows a pyrrolidine ring with an amide group attached to its nitrogen. The amide nitrogen is also bonded to a chiral center (labeled 'Chiral') which is further substituted with an aldehyde group and an 'R' group. Another amide group is attached to the chiral center, which is also substituted with a methyl group (H3C) and an amino group (NH2). |
| Cluster 12 | Nrps | 1354 | - | - |

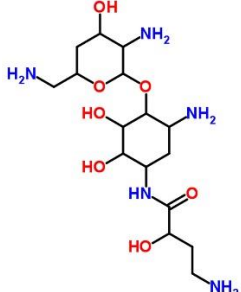
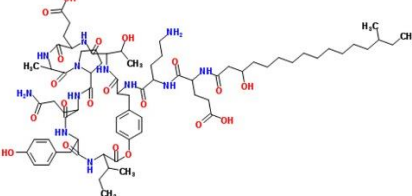
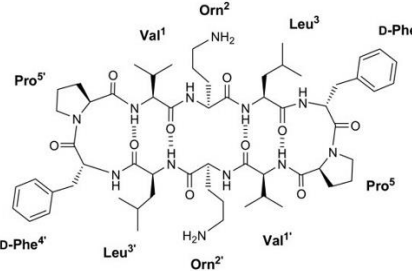
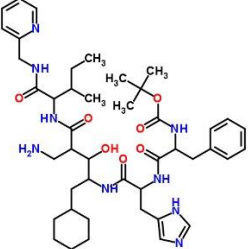
Supplementary Table S4. Compounds identified in *Bacillus halotolerans* isolates secreted fraction by LC-MS analysis.

| Rt (min) | HRESIMS | Molecular formula | Tentative identification | Structure | BFOA 4 | BFOA 3 | BFOA 2 | BFOA 1 | PGP activity | Biocontrol / antimicrobial activity | Herbicidal activity |
|----------|----------|---|--------------------------------------|--|--------|--------|--------|--------|-------------------------|-------------------------------------|---------------------|
| 1.39 | 188.0699 | C ₁₁ H ₁₁ NO ₂ | Indole propionic acid |  | + | + | + | + | Zhang and Davies (2016) | - | - |
| 1.71 | 197.1277 | C ₁₀ H ₁₈ N ₂ O ₂ | Cyclo(L-Leu-N-methyl-L-Ala) |  | - | + | + | + | - | - | - |
| 2.20 | 302.0819 | C ₁₁ H ₁₇ N ₃ O ₅ S | 2-thio-5(N-methylaminomethyl)uridine |  | - | - | + | + | - | - | - |
| 2.62 | 289.1550 | C ₁₆ H ₂₂ N ₂ O ₃ | Inthomycin-A |  | + | + | + | + | - | Webb et al. (2008) | Webb et al. (2008) |
| 2.90 | 211.1433 | C ₁₁ H ₂₀ N ₂ O ₂ | Cyclo(L-Val-L-Leu) |  | + | + | + | + | - | - | - |
| 3.49 | 245.1270 | C ₁₄ H ₁₈ N ₂ O ₂ | Cyclo(L-Val-L-Phe) |  | - | + | - | + | - | Guo et al. (2011) | - |

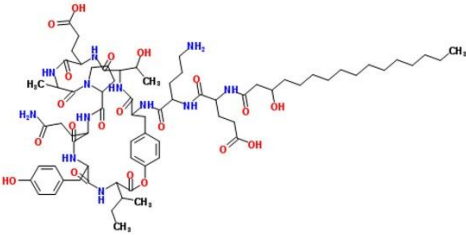
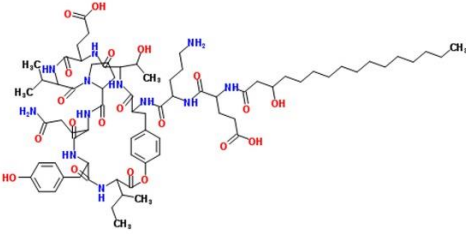
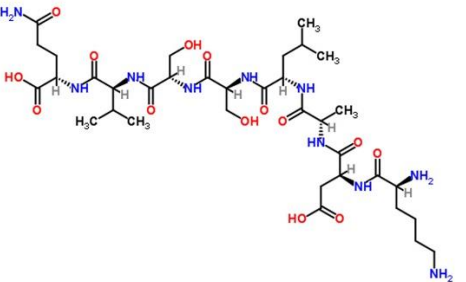
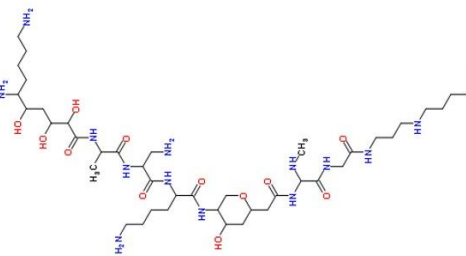
Supplementary Table 4. Continued.

| Rt (min) | HRESIMS | Molecular formula | Tentative identification | Structure | BFOA 4 | BFOA 3 | BFOA 2 | BFOA 1 | PGP activity | Biocontrol / antimicrobial activity | Herbicidal activity |
|----------|-----------|---|--|--|--------|--------|--------|--------|--------------|-------------------------------------|---------------------|
| 4.46 | 340.2242 | C ₁₇ H ₃₁ N ₃ O ₄ | Diprotin A (Isoleucyl Proyl Isoleucine) |  | + | + | + | + | - | - | - |
| 5.07 | 322.1295 | C ₁₆ H ₂₁ NO ₆ | Methyl 3-acetamido-4,6-O-benzylidene-3-deoxy-β-D-glucopyranoside |  | + | - | - | - | - | - | - |
| 5.17 | 239.1187 | C ₁₅ H ₁₆ N ₂ O | 3-(2-Butyl-5-oxazolyl)-1H-indole |  | + | - | - | - | - | - | - |
| 5.97 | 205.0966 | C ₁₁ H ₁₄ N ₂ O ₂ | N-Acetylkynuramine |  | + | - | - | - | - | - | - |
| 6.90 | 1139.6985 | C ₆₀ H ₉₂ N ₁₂ O ₁₀ | Gramicidin S |  | + | + | + | + | - | Gause and Brazhnikova (1944) | - |

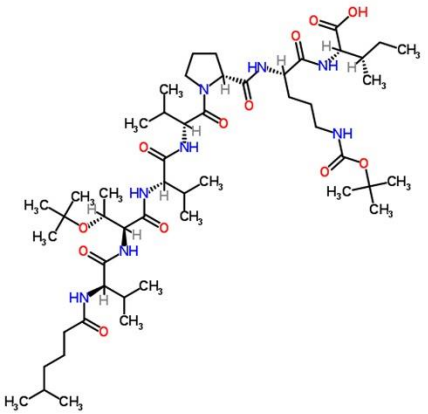
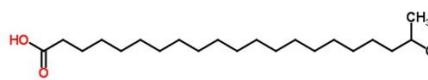
Supplementary Table 4. Continued.

| Rt (min) | HRESIMS | Molecular formula | Tentative identification | Structure | BFOA 4 | BFOA 3 | BFOA 2 | BFOA 1 | PGP activity | Biocontrol / antimicrobial activity | Herbicidal activity |
|----------|-----------|--|--------------------------|--|--------|--------|--------|--------|--------------|-------------------------------------|---------------------|
| 7.28 | 406.2300 | C ₁₆ H ₃₃ N ₅ O ₇ | 5-Deoxybutirosamine |  | + | - | - | - | - | Taylor and Schmitz (1976) | - |
| 7.33 | 1475.8043 | C ₇₃ H ₁₁₂ N ₁₂ O ₂₀ | Plipastatin A2 |  | + | + | + | + | - | Umezawa et al. (1986) | - |
| 7.44 | 1125.6832 | C ₅₉ H ₉₀ N ₁₂ O ₁₀ | Gramicidin S2 |  | + | + | + | + | - | Solanas et al. (2009) | - |
| 7.51 | 830.4935 | C ₄₄ H ₆₅ N ₉ O ₇ | Gramicidin-J |  | + | + | + | + | - | Hunter and Schwartz (1967) | - |

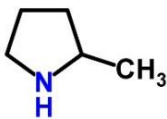
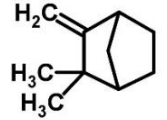
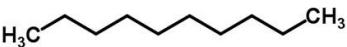
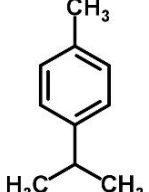
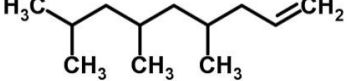
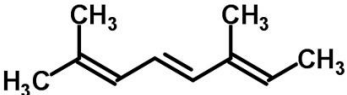
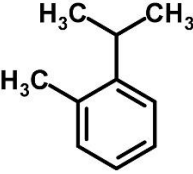
Supplementary Table 4. Continued.

| Rt (min) | HRESIMS | Molecular formula | Tentative identification | Structure | BFOA 4 | BFOA 3 | BFOA 2 | BFOA 1 | PGP activity | Biocontrol / antimicrobial activity | Herbicidal activity |
|----------|-----------|--|--------------------------|--|--------|--------|--------|--------|---------------------|-------------------------------------|---------------------|
| 7.61 | 1461.7884 | C ₇₂ H ₁₁₀ N ₁₂ O ₂₀ | Plipastatin A1 |  | + | - | + | + | - | Ma and Hu (2018) | - |
| 7.66 | 1489.8196 | C ₇₄ H ₁₁₄ N ₁₂ O ₂₀ | Plipastatin B1 |  | + | - | + | + | - | Tsuge et al. (1996) | - |
| 7.97 | 845.4376 | C ₃₅ H ₆₂ N ₁₀ O ₁₄ | Rotihibin-B |  | + | - | - | - | Fukuchi et al. 1995 | - | - |
| 8.39 | 961.6161 | C ₄₁ H ₈₂ N ₁₄ O ₁₂ | Galantin-I |  | + | - | - | - | - | Sakai and Ohfuné (1992) | - |

Supplementary Table 4. Continued.

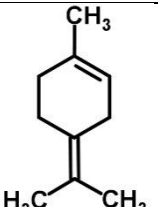
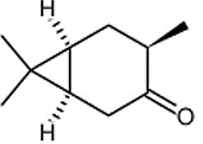

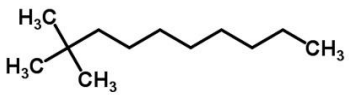
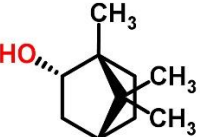
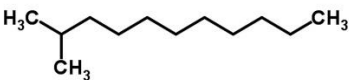
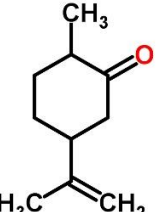
| Rt (min) | HRESIMS | Molecular formula | Tentative identification | Structure | BFOA 4 | BFOA 3 | BFOA 2 | BFOA 1 | PGP activity | Biocontrol / antimicrobial activity | Herbicidal activity |
|----------|-----------|---|-----------------------------|--|--------|--------|--------|--------|--------------|--|---------------------|
| 8.46 | 993.6427 | C ₄₂ H ₈₆ N ₁₄ O ₁₃ | Lysinegalantin I | – | + | + | + | + | – | – | – |
| 8.57 | 1007.6763 | C ₅₁ H ₉₂ N ₈ O ₁₂ | Pseudofactin I |  | + | + | + | + | – | Janek et al. (2012); Biniarz et al. (2015) | – |
| 8.62 | 1021.6916 | C ₅₂ H ₉₄ N ₈ O ₁₂ | Pseudofactin II | – | | | | | | | |
| 9.28 | 339.3266 | C ₂₂ H ₄₄ O ₂ | 20-methylheneicosanoic acid |  | – | + | + | + | – | – | – |

Supplementary Table S5. Compounds identified in *Bacillus halotolerans* strains secreted fraction by GC-MS analysis.

| Rt (min) | Kovat's Index | Molecular formula | Tentative identification | Structure | BFOA4 | BFOA3 | BFOA2 | BFOA1 | PGP activity | Biocontrol / antimicrobial activity | Herbicidal/ Insecticide activity |
|----------|---------------|----------------------------------|---------------------------|--|-------|-------|-------|-------|--------------|-------------------------------------|---|
| 4.69 | 873 | C ₅ H ₁₁ N | Ethyl-2-methylpyrrolidine |  | + | + | - | - | - | - | - |
| 5.10 | 942 | C ₁₀ H ₁₆ | Camphene |  | + | + | - | - | - | - | - |
| 5.92 | 1003 | C ₁₀ H ₂₂ | Decane |  | - | + | - | - | - | - | - |
| 6.02 | 1011 | C ₁₀ H ₁₄ | p-Cymene |  | - | + | - | - | - | - | - |
| 6.15 | 1014 | C ₁₂ H ₂₄ | Trimethyl-1-nonene |  | + | + | - | - | - | - | - |
| 6.43 | 1018 | C ₁₀ H ₁₆ | D-Limonene |  | - | - | + | + | - | - | Avenger Material Safety Data Sheet ¹ |
| 6.78 | 1025 | C ₁₀ H ₁₄ | o-Cymene |  | + | + | - | - | - | - | - |

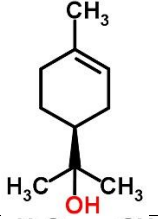
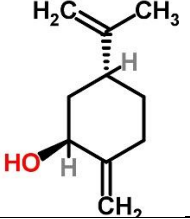
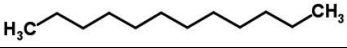
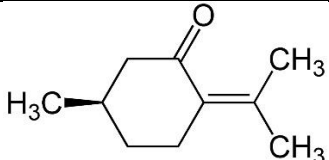
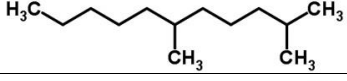
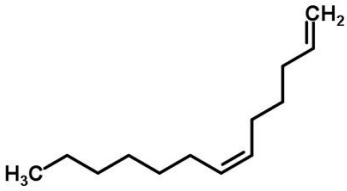
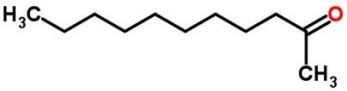
¹ <http://nebula.wsimg.com/07de45c0af774ba73e06362ad1a56f06?AccessKeyId=C67FD801C8FC93742D64&disposition=0&alloworigin=1>

Supplementary Table S5. Continued.


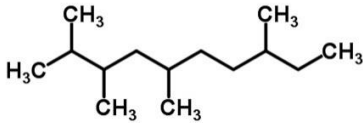
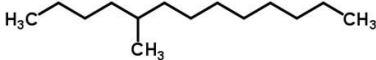
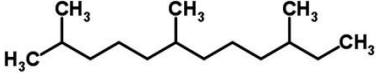
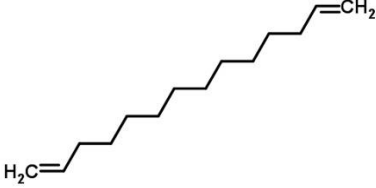
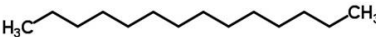
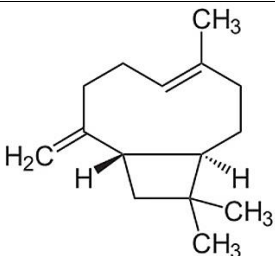
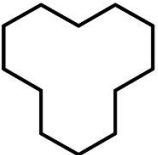
| Rt (min) | Kovat's Index | Molecular formula | Tentative identification | Structure | BFOA4 | BFOA3 | BFOA2 | BFOA1 | PGP activity | Biocontrol / antimicrobial activity | Herbicidal/ Insecticide activity |
|----------|---------------|-----------------------------------|--------------------------|--|-------|-------|-------|-------|--------------|-------------------------------------|-----------------------------------|
| 6.87 | 1083 | C ₁₀ H ₁₆ | α -terpinolene |  | + | + | + | - | - | - | - |
| 7.51 | 1109 | C ₁₀ H ₁₆ O | cis-4-Caranone |  | + | + | - | - | - | - | - |
| 7.66 | 1115 | C ₁₁ H ₂₄ | Undecane |  | + | + | - | - | - | - | - |
| 7.88 | 1151 | C ₁₂ H ₂₆ | Dimethyldecane |  | + | + | - | - | - | - | - |
| 8.15 | 1158 | C ₁₀ H ₁₈ O | Borneol |  | + | + | - | - | - | - | Chemical Information ² |
| 8.28 | 1170 | C ₁₂ H ₂₆ | 2-methyl-undecane |  | + | + | - | - | - | - | - |
| 8.41 | 1179 | C ₁₀ H ₁₆ O | Dihydrocarvone |  | + | + | - | - | - | - | - |

² <https://web.archive.org/web/20041107161247/http://sun.ars-grin.gov:8080/npgspub/xsql/duke/chemdisp.xsql?chemical=BORNEOL>

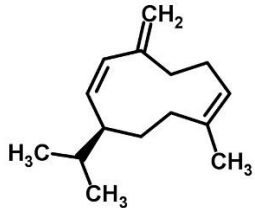
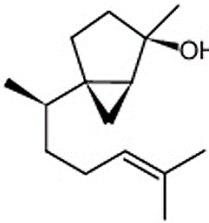
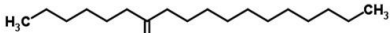
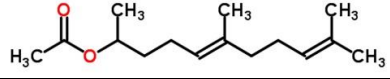
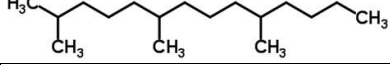
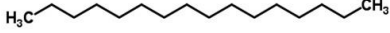
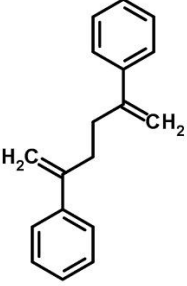
Supplementary Table S5. Continued.

| Rt (min) | Kovat's Index | Molecular formula | Tentative identification | Structure | BFOA4 | BFOA3 | BFOA2 | BFOA1 | PGP activity | Biocontrol / antimicrobial activity | Herbicidal/ Insecticide activity |
|----------|---------------|-----------------------------------|---------------------------------|--|-------|-------|-------|-------|--------------|-------------------------------------|----------------------------------|
| 8.57 | 1185 | C ₁₀ H ₁₈ O | α -terpineol |  | + | + | - | - | - | - | - |
| 9.06 | 1201 | C ₁₀ H ₁₆ O | trans-p-mentha-1(7),8-dien-2-ol |  | + | + | - | - | - | - | - |
| 9.13 | 1209 | C ₁₂ H ₂₆ | Dodecane |  | + | + | - | - | - | - | - |
| 9.21 | 1216 | C ₁₀ H ₁₆ O | Pulegone |  | + | + | - | - | - | - | Franzios et al. (1997) |
| 9.34 | 1234 | C ₁₃ H ₂₈ | 2,3-dimethylundecane |  | + | + | + | - | - | - | - |
| 9.64 | 1271 | C ₁₃ H ₂₄ | Tridecadiene |  | + | + | + | - | - | - | - |
| 9.99 | 1298 | C ₁₁ H ₂₂ O | 2-undecanone |  | + | + | - | - | - | - | Stephen et al. (2002) |

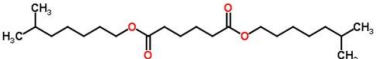
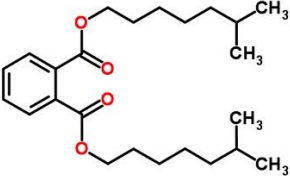
Supplementary Table S5. Continued.

| Rt (min) | Kovat's Index | Molecular formula | Tentative identification | Structure | BFOA4 | BFOA3 | BFOA2 | BFOA1 | PGP activity | Biocontrol / antimicrobial activity | Herbicidal/ Insecticide activity |
|----------|---------------|---------------------------------|-----------------------------|--|-------|-------|-------|-------|--------------|-------------------------------------|----------------------------------|
| 10.51 | 1313 | C ₁₃ H ₂₈ | Tridecane |  | + | + | + | - | - | - | - |
| 10.67 | 1319 | C ₁₄ H ₃₀ | Tetramethyldecane |  | + | + | + | - | - | - | - |
| 10.83 | 1349 | C ₁₄ H ₃₀ | Methyltridecane |  | + | + | + | - | - | - | - |
| 11.14 | 1381 | C ₁₅ H ₃₂ | Farnesane |  | - | + | + | - | - | - | - |
| 11.93 | 1411 | C ₁₄ H ₂₆ | Tetradecadiene |  | + | + | + | - | - | - | - |
| 12.32 | 1417 | C ₁₄ H ₃₀ | Tetradecane |  | - | + | + | - | - | - | - |
| 13.18 | 1423 | C ₁₅ H ₂₄ | <i>trans</i> -Caryophyllene |  | - | + | + | - | - | - | - |
| 13.26 | 1439 | C ₁₂ H ₂₄ | Cyclododecane |  | - | - | + | - | - | - | - |

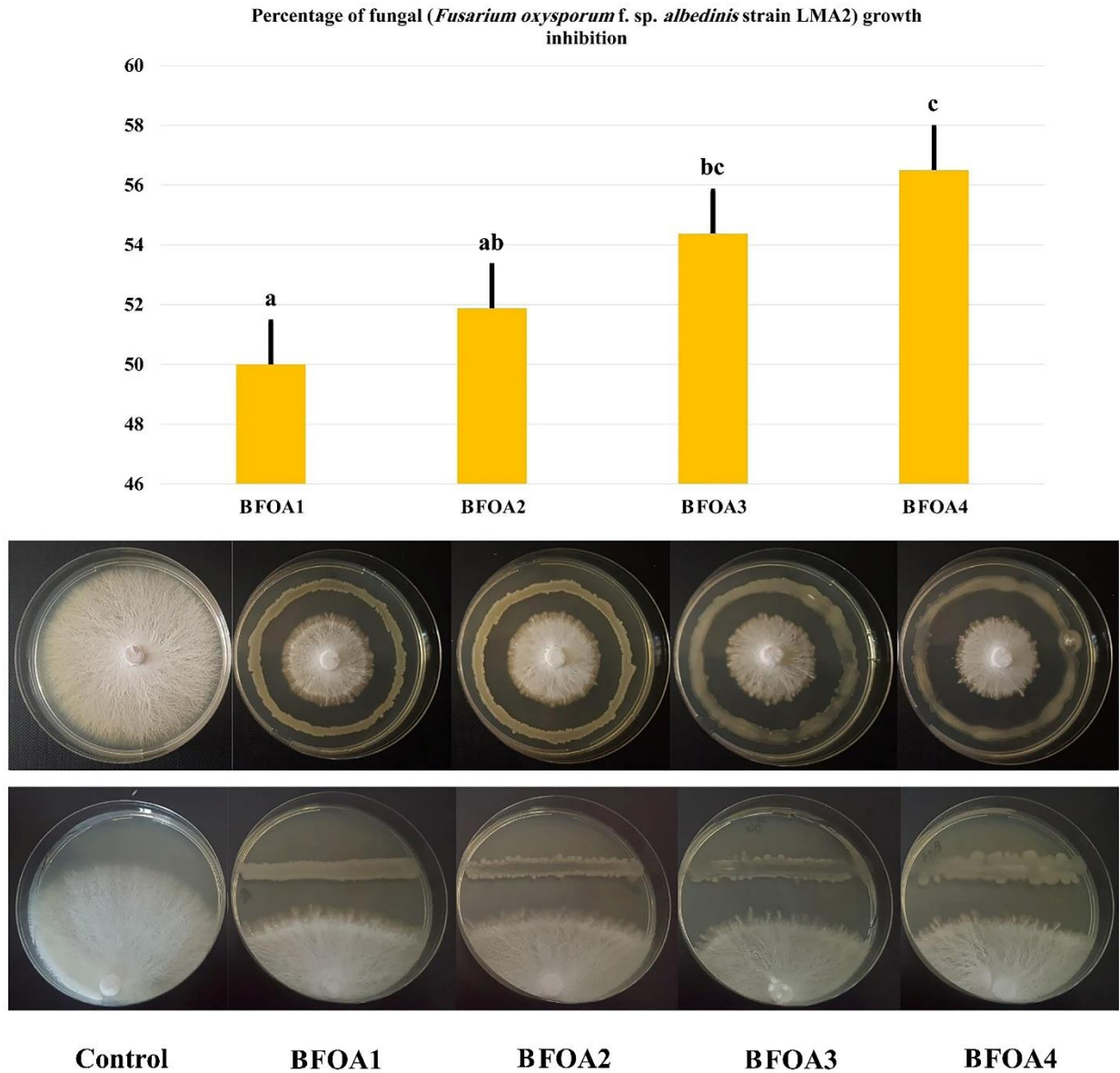
Supplementary Table S5. Continued.

| Rt (min) | Kovat's Index | Molecular formula | Tentative identification | Structure | BFOA4 | BFOA3 | BFOA2 | BFOA1 | PGP activity | Biocontrol / antimicrobial activity | Herbicidal/ Insecticide activity |
|----------|---------------|--|----------------------------|--|-------|-------|-------|-------|--------------|-------------------------------------|----------------------------------|
| 13.38 | 1477 | C ₁₅ H ₂₄ | Germacrene D |  | - | - | + | - | - | Adio (2009) | Adio (2009) |
| 13.57 | 1523 | C ₁₅ H ₂₆ O | 7-epi-cis-sesquisabinene |  | - | - | + | - | - | - | - |
| 13.79 | 1563 | C ₁₈ H ₃₆ O | Hexa-hydro-farnesyl |  | - | + | + | - | - | - | - |
| 14.50 | 1573 | C ₁₅ H ₂₆ O ₂ | Geranyl isovalerate |  | - | + | - | - | - | - | - |
| 15.94 | 1557 | C ₁₇ H ₃₆ | Trimethyltetradecane |  | - | - | + | - | - | - | - |
| 16.96 | 1612 | C ₁₆ H ₃₄ | Hexadecane |  | - | - | + | - | - | - | - |
| 34.71 | 1939 | C ₁₈ H ₁₈ | 1,5-Diphenyl-1,5-hexadiene |  | - | - | + | + | - | - | - |

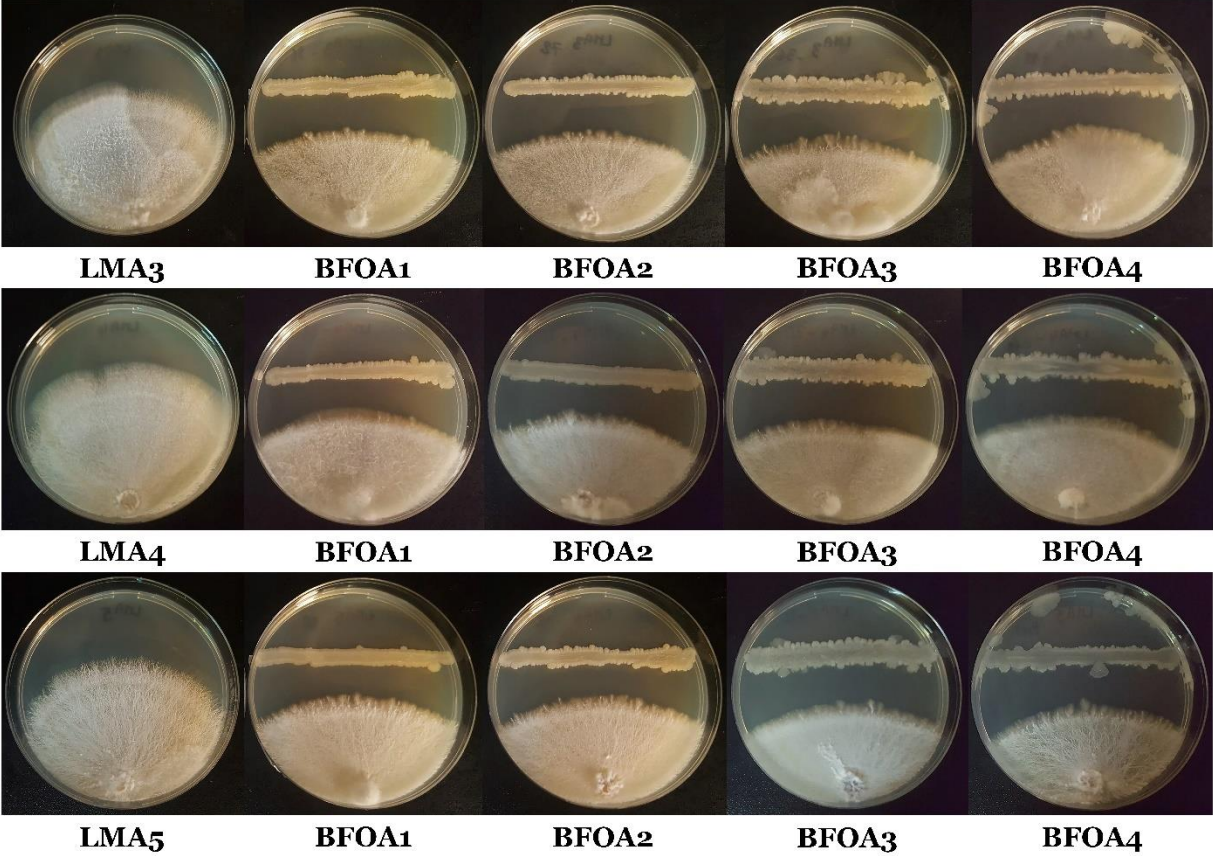
Supplementary Table S5. Continued.

| Rt (min) | Kovat's Index | Molecular formula | Tentative identification | Structure | BFOA4 | BFOA3 | BFOA2 | BFOA1 | PGP activity | Biocontrol / antimicrobial activity | Herbicidal/ Insecticide activity |
|----------|---------------|--|--------------------------|--|-------|-------|-------|-------|--------------|-------------------------------------|----------------------------------|
| 35.82 | 2244 | C ₂₂ H ₄₂ O ₄ | Diisooctyl adipate |  | - | - | + | + | - | - | - |
| 37.00 | 2704 | C ₂₄ H ₃₈ O ₄ | Diisooctyl phthalate |  | - | - | + | - | - | - | - |

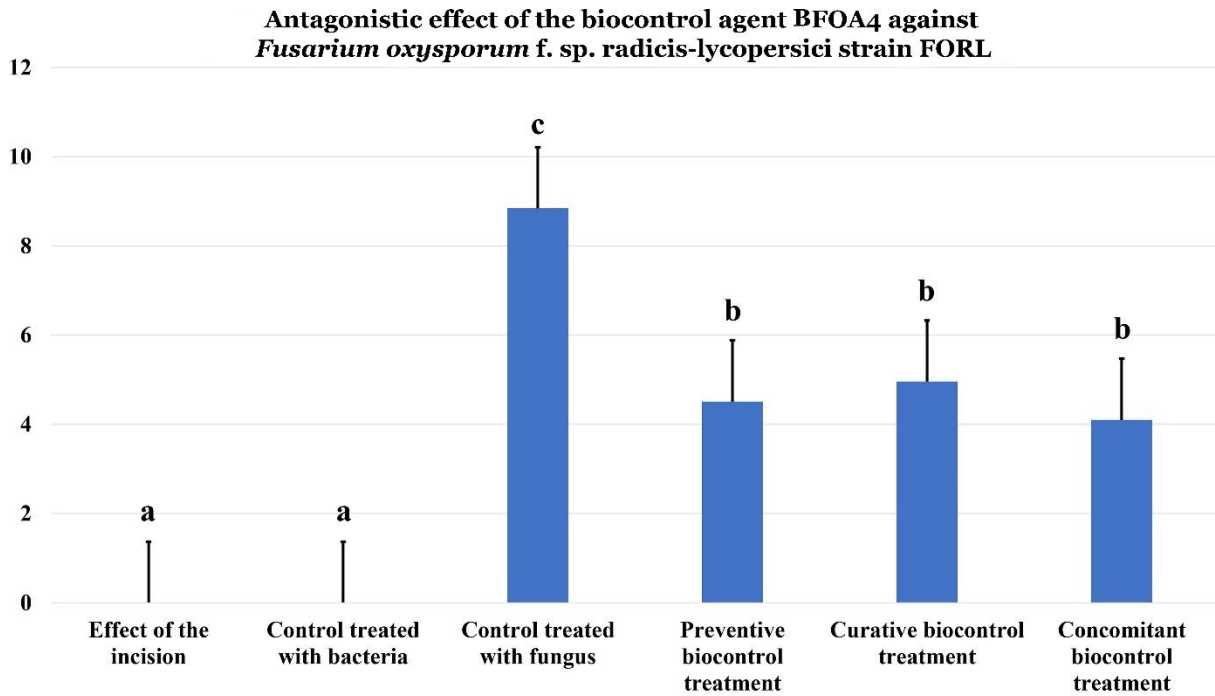
Supplementary Figure S1. Percentage of fungal (*Fusarium oxysporum* f. sp. *albedinis* strain LMA2) growth inhibition against *Bacillus halotolerans* isolates. Data presents mean \pm standard error. Bars labelled with different letters are significantly different among the treatments at $p < 0.05$ using the Tukey's HSD test.



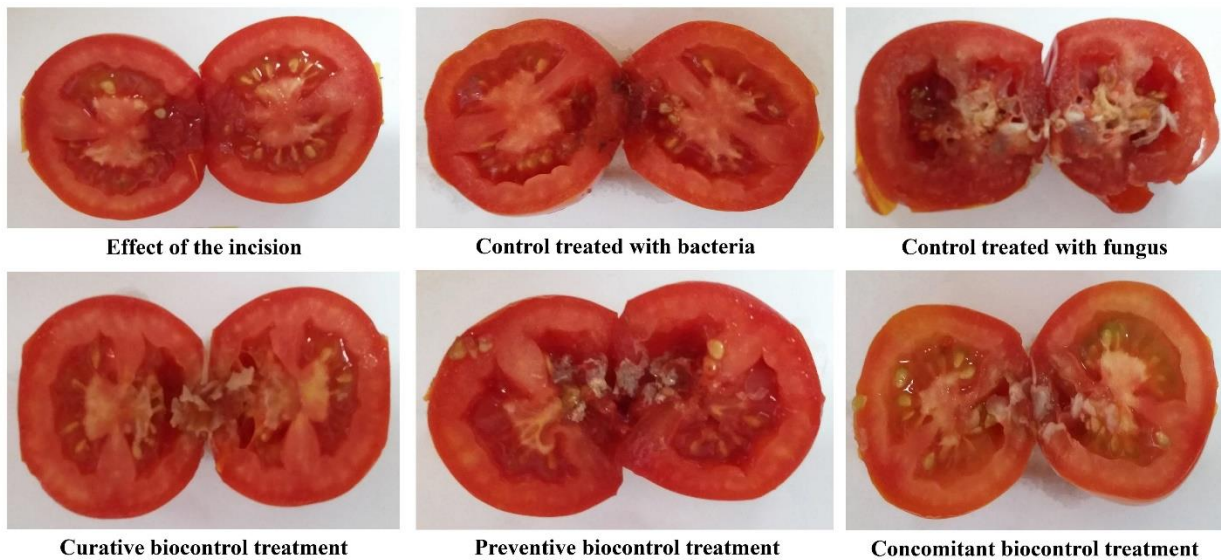
Supplementary Figure S2. *Fusarium oxysporum* f. sp. *albedinis* strains LMA3, LMA4 and LMA5 growth inhibition against *Bacillus halotolerans* isolates.



Supplementary Figure S3. Antagonistic effect of the biocontrol agent FOA4 against *Fusarium oxysporum* f. sp. radicis-lycopersici strain FORL. (A) Effects of FOA1-4 bacterial isolates on *Fusarium oxysporum* f. sp. radicis-lycopersici strain FORL tomato fruit rot severity as compared to tomato fruit controls. Data presents mean \pm standard error. Bars labelled with different letters are significantly different among the treatments at $p < 0.05$ using the Tukey's HSD test. (B) Effect of preventive, concomitant, and curative biocontrol treatment on tomato fruits at 7 days post-inoculation with *Fusarium oxysporum* f. sp. radicis-lycopersici strain FORL.



A



References

- Adio, A. M. (2009). Germacrenes A–E and related compounds: thermal, photochemical and acid induced transannular cyclizations. *Tetrahedron*. 65, 1533–1552 10.1016/j.tet.2008.11.050
- Biniarz, P., Baranowska, G., Feder-Kubis, J. and Krasowska, A. (2015). The lipopeptides pseudofactin II and surfactin effectively decrease *Candida albicans* adhesion and hydrophobicity. *Antonie Leeuwenhoek*. 108, 343–353. doi: 10.1007/s10482-015-0486-3
- Franzios, G., Mirotsoy, M., Hatziapostolou, E., Kral, J., Scouras, Z. G., Mavragani-Tsipidou, P. (1997). Insecticidal and genotoxic activities of mint essential oils. *J. Agri. Food Chem.* 45 (7), 2690–2694. doi:10.1021/jf960685f
- Fukuchi, N., Furihata, K., Nakayama, J., and Isogai, A. (1995). Rotihibins, novel plant growth regulators from *Streptomyces graminofaciens*. *J. Antibiot.* 48(9), 1004–1010. doi: 10.7164/antibiotics.48.1004
- Gause, G. F., Brazhnikova, M. G. (1944). Gramicidin S and its use in the treatment of infected Wounds. *Nature*. 154(3918): 703–703, doi:10.1038/154703a0
- Guo, X., Zheng, L., Zhou, W. et al. (2011). A case study on chemical defense based on quorum sensing: antibacterial activity of sponge-associated bacterium *Pseudoalteromonas* sp. NJ6-3-1 induced by quorum sensing mechanisms. *Ann Microbiol.* 61(2), 247–255. doi: 10.1007/s13213-010-0129-x
- Hunter, F. E. and Schwartz, L. S. (1967). Tyrocidines and Gramicidin S (J1, J2). In: Gottlieb, D., Shaw, P. D. (eds) *Antibiotics*. Springer, Berlin, Heidelberg.
- Janek, T., Lukaszewicz, M. and Krasowska, A. (2012). Antiadhesive activity of the biosurfactant pseudofactin II secreted by the arctic bacterium *Pseudomonas fluorescens* BD5. *BMC Microbiol.* 12:24.
- Ma, Z. and Hu, J. (2018). Plipastatin A1 produced by a marine sediment-derived *Bacillus amyloliquefaciens* SH-B74 contributes to the control of gray mold disease in tomato. *3 Biotech.* 8(2):125. doi: 10.1007/s13205-018-1144-z
- Sakai, N. and Ohfuné, Y. (1992). Total synthesis of Galantin I. Acid-catalyzed cyclization of galantinic acid. *J. Am. Chem. Soc.* 114, 998–1010.
- Solanas, C., de la Torre, B. G., Fernandez-Reyes, M., et al. (2009). Therapeutic Index of Gramicidin S is Strongly Modulated by D-Phenylalanine Analogues at the Beta-Turn. *J. Med. Chem.* 52, 664–674.
- Stephen, J., Toth, Jr. and Buhler, W. G. (2002). North Carolina State University scientist discovers mosquito repellent in tomatoes. *Pesticide Broadcast*. 12: (5).
- Taylor, H. D. and Schmitz, H. (1976). Antibiotics derived from a mutant of *Bacillus circulans*. *J Antibiot.* 29(5), 532–535.
- Tsuge, K., Ano, T. and Shoda, M. (1996). Isolation of a gene essential for biosynthesis of the lipopeptide antibiotics plipastatin B1 and surfactin in *Bacillus subtilis* YB8. *Arch. Microbiol.* 165(4), 243–251.

Umezawa, H., Aoyagi, T., Nishikiori, T., Okuyama, A., Yamagishi, Y., Hamada, M. and Takeuchi, T. (1986). Plipastatins: new inhibitors of phospholipase A2, produced by *Bacillus cereus* BMG302-fF67. I. Taxonomy, production, isolation and preliminary characterization. *J Antibiot.* (Tokyo). 39 (6), 737-44.

Webb, M. R., Addie, M. S., Crawforth, C. M. et al. (2008). The syntheses of rac-inthomycin A, (+)-inthomycin B and (+)-inthomycin C using a unified synthetic approach. *Tetrahedron.* 64 (21), 4778-4791. doi: 10.1016/j.tet.2008.01.116

Zhang, L. S. and Davies, S. S. (2016). Microbial metabolism of dietary components to bioactive metabolites: opportunities for new therapeutic interventions. *Genome Med.* 8:46. doi: 10.1186/s13073-016-0296-x