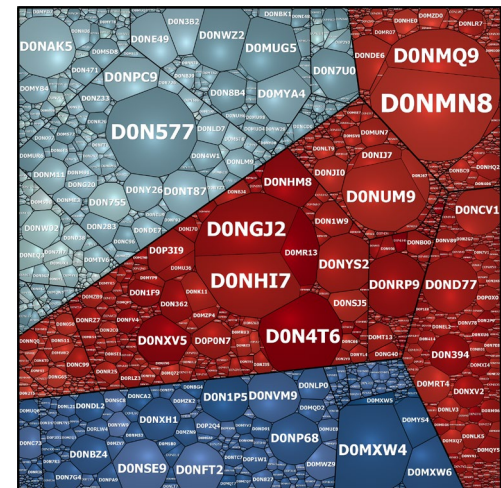
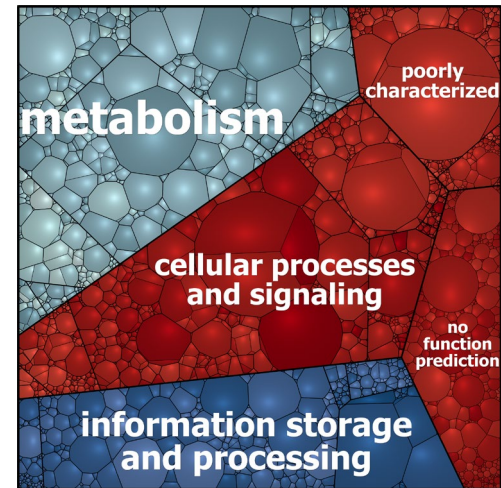
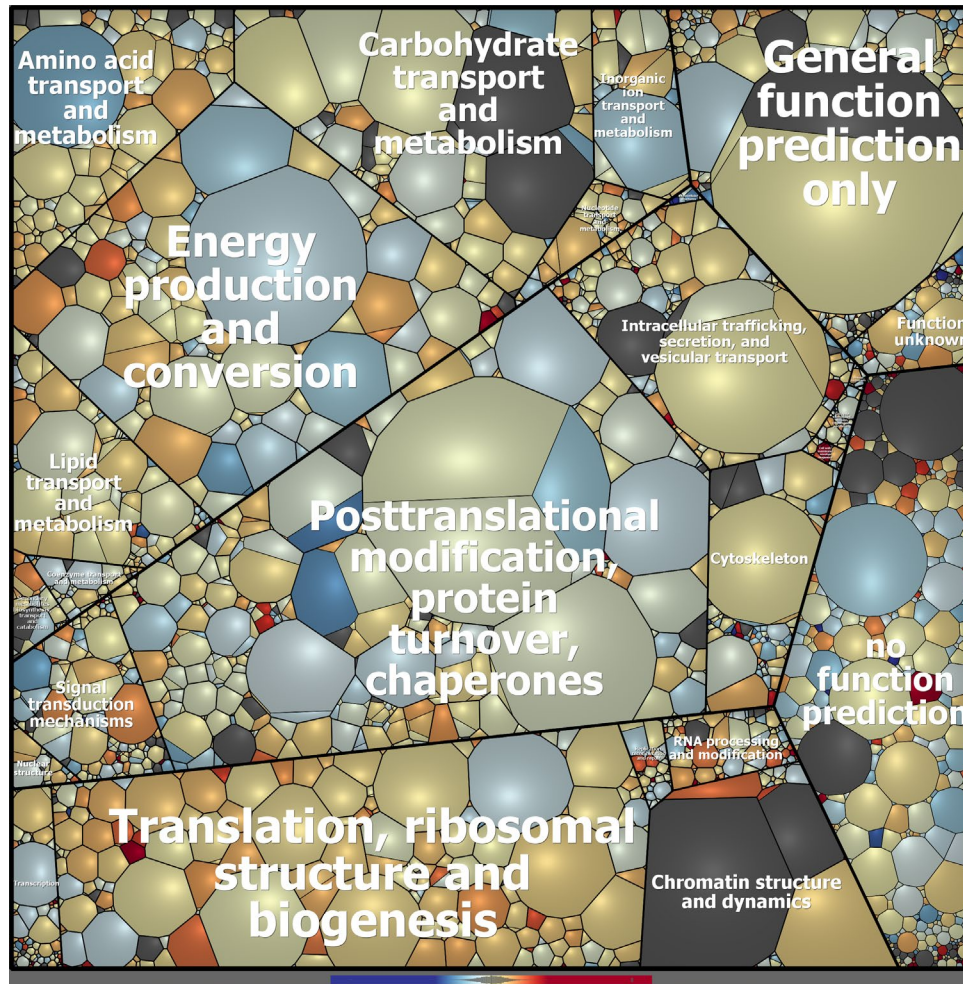


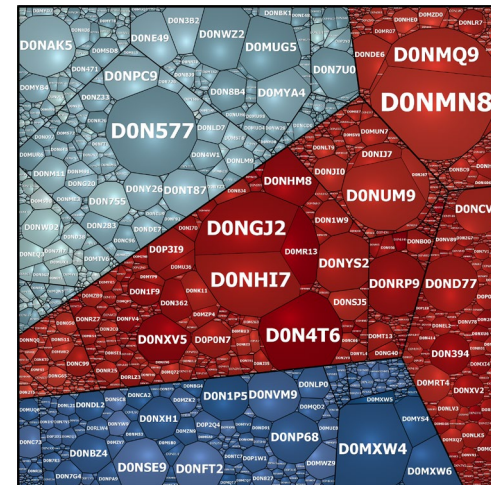
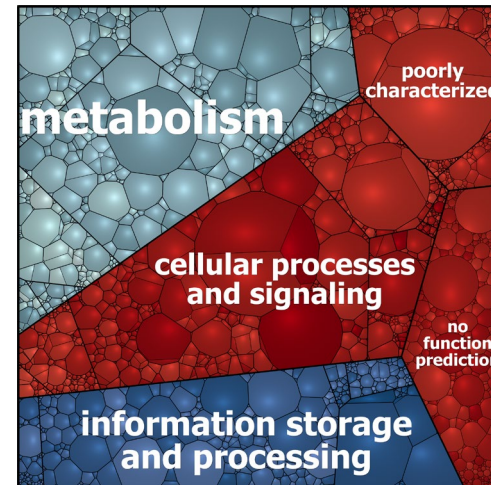
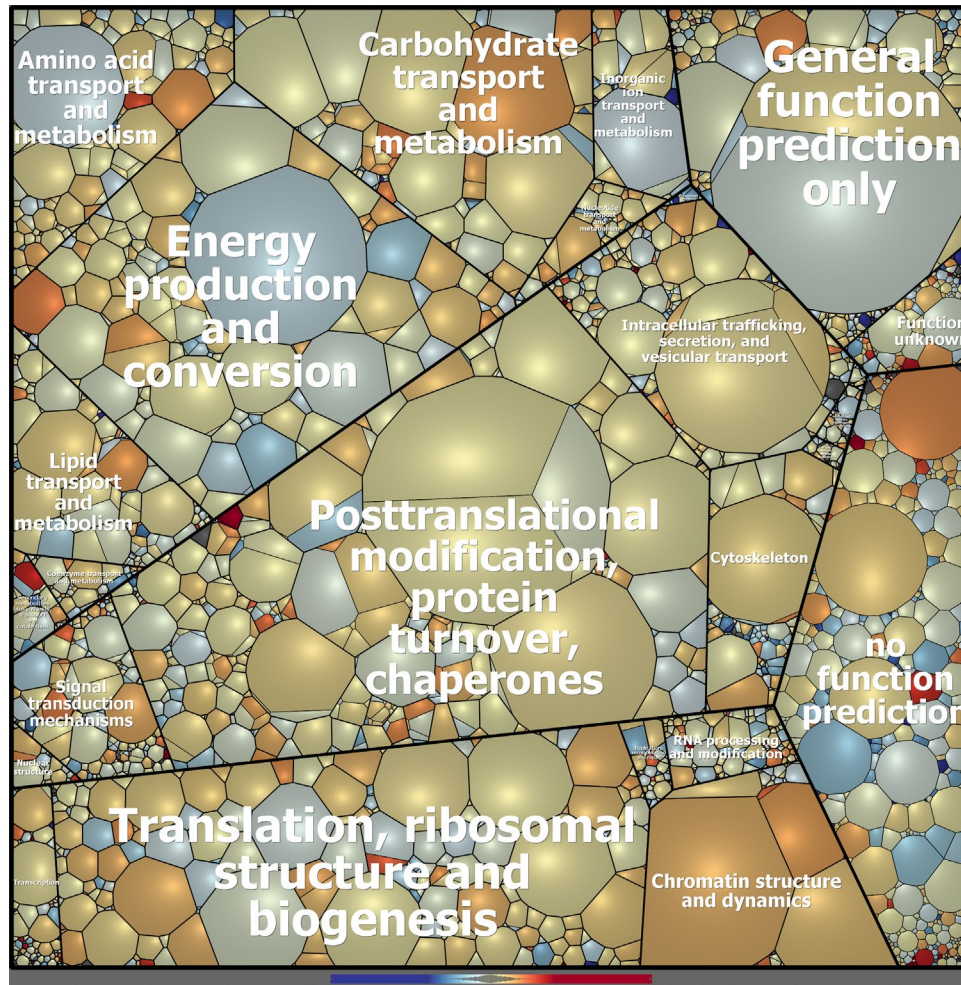
Effect of DMDS exposure on protein pattern of *Phytophthora infestans*



Voronoi treemap visualization of protein expression pattern of *P. infestans* exposed to DMDS.

Relative changes compared to untreated control are color-coded: red – higher amount in treated, blue – higher amount in untreated cultures. Cells sizes were determined by the average protein occurrence level within all samples. Proteins are represented by cells, cells are clustered according their functional classification. Classification was done using Prophan 2.0. Grey cells indicate proteins that were not identified.

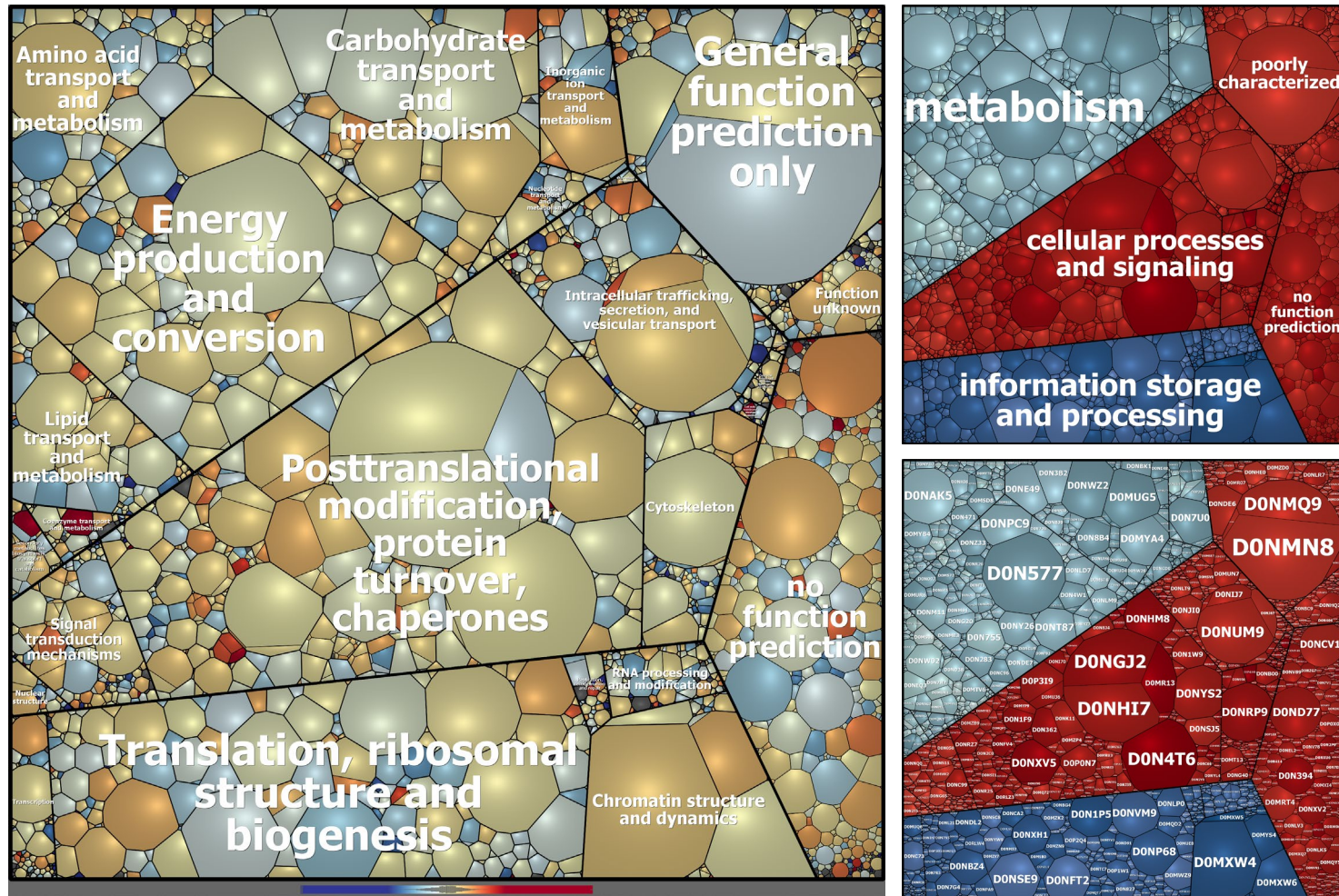
Effect of **DMTS** exposure on protein pattern of *Phytophthora infestans*



Voronoi treemap visualization of protein expression pattern of *P. infestans* exposed to DMTS.

Relative changes compared to untreated control are color-coded: red – higher amount in treated, blue – higher amount in untreated cultures. Cells sizes were determined by the average protein occurrence level within all samples. Proteins are represented by cells, cells are clustered according their functional classification. Classification was done using Prophan 2.0. Grey cells indicate proteins that were not identified.

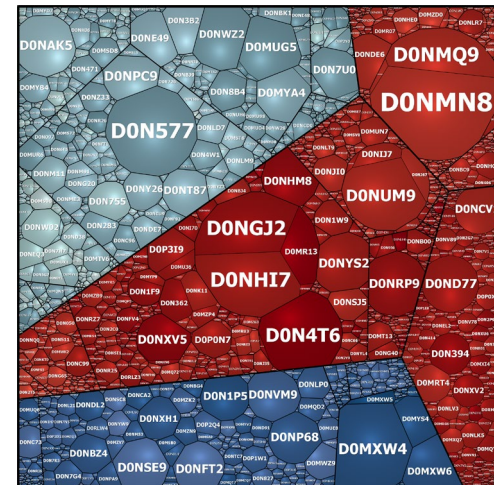
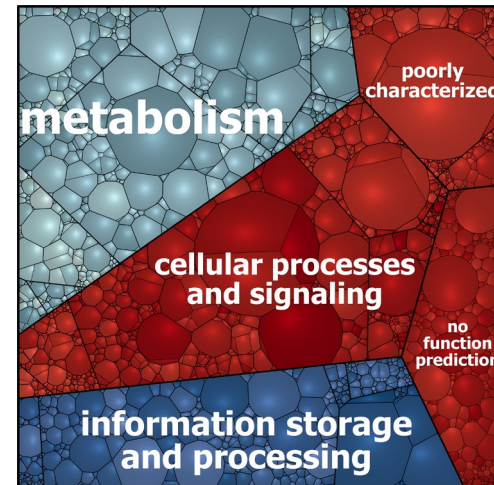
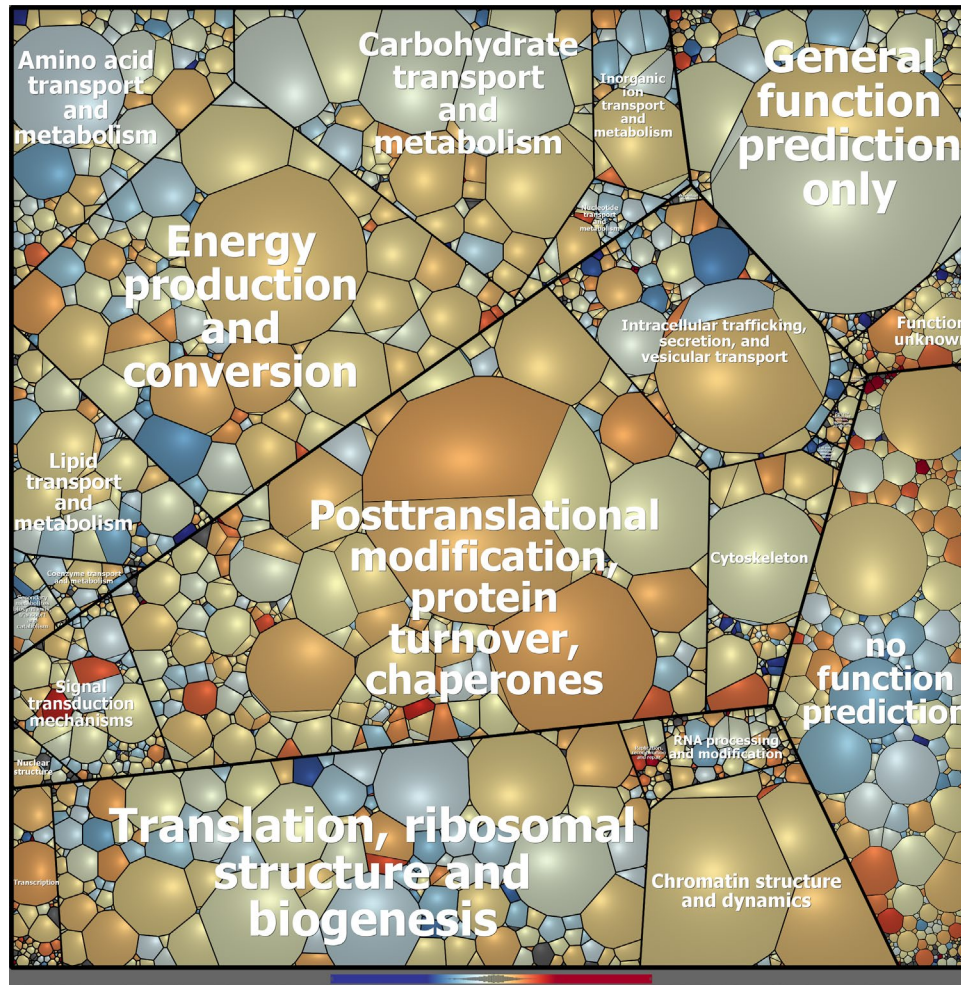
Effect of MMTS exposure on protein pattern of *Phytophthora infestans*



Voronoi treemap visualization of protein expression pattern of *P. infestans* exposed to MMTS.

Relative changes compared to untreated control are color-coded: red – higher amount in treated, blue – higher amount in untreated cultures. Cells sizes were determined by the average protein occurrence level within all samples. Proteins are represented by cells, cells are clustered according their functional classification. Classification was done using Prophan 2.0. Grey cells indicate proteins that were not identified.

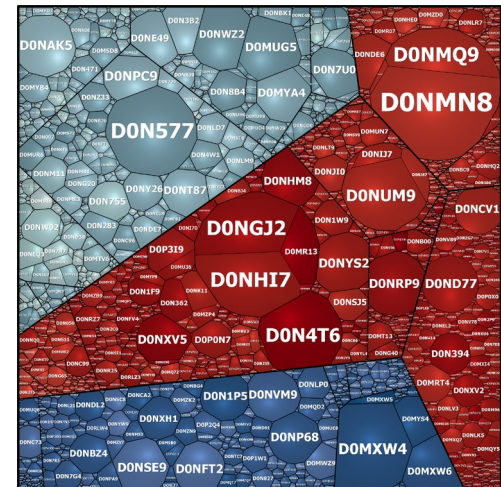
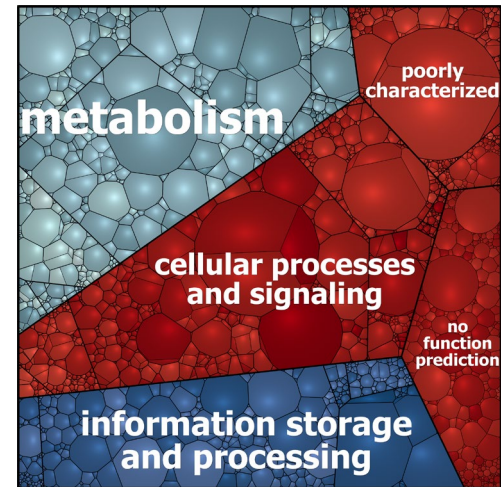
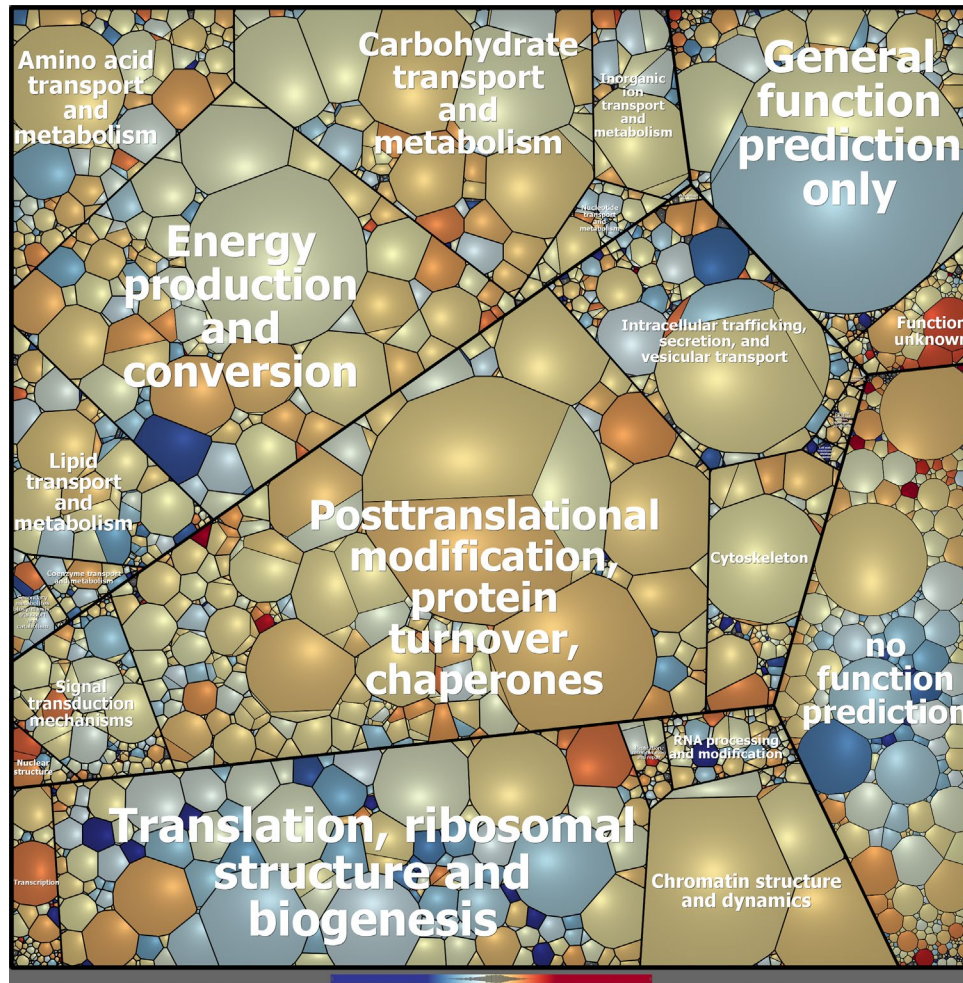
Effect of SM exposure on protein pattern of *Phytophthora infestans*



Voronoi treemap visualization of protein expression pattern of *P. infestans* exposed to SM.

Relative changes compared to untreated control are color-coded: red – higher amount in treated, blue – higher amount in untreated cultures. Cells sizes were determined by the average protein occurrence level within all samples. Proteins are represented by cells, cells are clustered according their functional classification. Classification was done using Prophan 2.0. Grey cells indicate proteins that were not identified.

Effect of **BMTMS** exposure on protein pattern of *Phytophthora infestans*



Voronoi treemap visualization of protein expression pattern of *P. infestans* exposed to BMTMS.

Relative changes compared to untreated control are color-coded: red – higher amount in treated, blue – higher amount in untreated cultures. Cells sizes were determined by the average protein occurrence level within all samples. Proteins are represented by cells, cells are clustered according their functional classification. Classification was done using Prophan 2.0. Grey cells indicate proteins that were not identified.