Table S3: Confusion matrix for Viridiplantae representing the number of proteins for each targeting peptides predicted by TargetP 2.0 (rows) versus observed in the test set (columns).

|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
|

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|   | Class | SP | mTP | cTP | luTP | noTP |
|   | SP | 272 | 0 | 2 | 0 | 3 |
|   | mTP | 0 | 117 | 2 | 0 | 11 |
|   | cTP | 0 | 2 | 197 | 9 | 9 |
|   | luTP | 0 | 0 | 11 | 34 | 0 |
|   | noTP | 10 | 6 | 15 | 2 | 1779 |

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