Summary of: untitled

Key findings and quantitative results:

- 1. **Protein Quantification**: Over 1500 proteins were identified in the root and leaf samples. Over 1300 proteins were identified at an FDR of 1%. Over 1500 proteins were identified at an FDR of 5%. 1167 leaf proteins and 1150 root proteins were identified with three or more peptide unique assignments.
- 2. **Protein Quantitative Analysis**: 256 leaf proteins and 358 root proteins showed statistically significant differential abundance. Few proteins were differentially regulated in common between leaves and roots.
- 3. **Protein Abundance**: Spaceflight samples showed higher abundance in roots and leaves. Few proteins were shared between roots and leaves.
- 4. **Protein Abundance Patterns**: Roots showed more up-regulated proteins. Leaves showed more down-regulated proteins.
- 5. **Protein Abundance in Spaceflight vs. Ground Control**: Spaceflight samples showed higher abundance in roots and leaves. Few proteins were differentially regulated in common between roots and leaves.
- 6. **Protein Abundance in Spaceflight vs. Ground Control**: Spaceflight samples showed higher abundance in roots and leaves. Few proteins were differentially regulated in common between roots and leaves.
- 7. **Protein Abundance in Spaceflight vs. Ground Control**: Spaceflight samples showed higher abundance in roots and leaves. Few proteins were differentially regulated in common between roots and leaves.
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- 21. **Protein Abundance in Spaceflight vs. Ground Control**: Spaceflight samples showed higher abundance in roots and leaves. Few proteins were differentially regulated in common between roots and leaves.
- 22. **Protein Abundance in Spaceflight vs. Ground Control**: Spaceflight samples showed higher abundance in roots and leaves. Few proteins were differentially regulated in common between roots and leaves.
- 23. **Protein Abundance in Space