

Viral Ecology of Onion Bulbs with Bacterial Rot

Pedro H Lebre¹, Christopher Liakos², Verushka Ibanez¹, Don A Cowan¹, Teresa Coutinho¹

Centre for Microbial Ecology and Genomics,¹ Department of Biochemistry, Genetics and Microbiology,² Department of Plant and Soil Sciences, University of Pretoria, South Africa

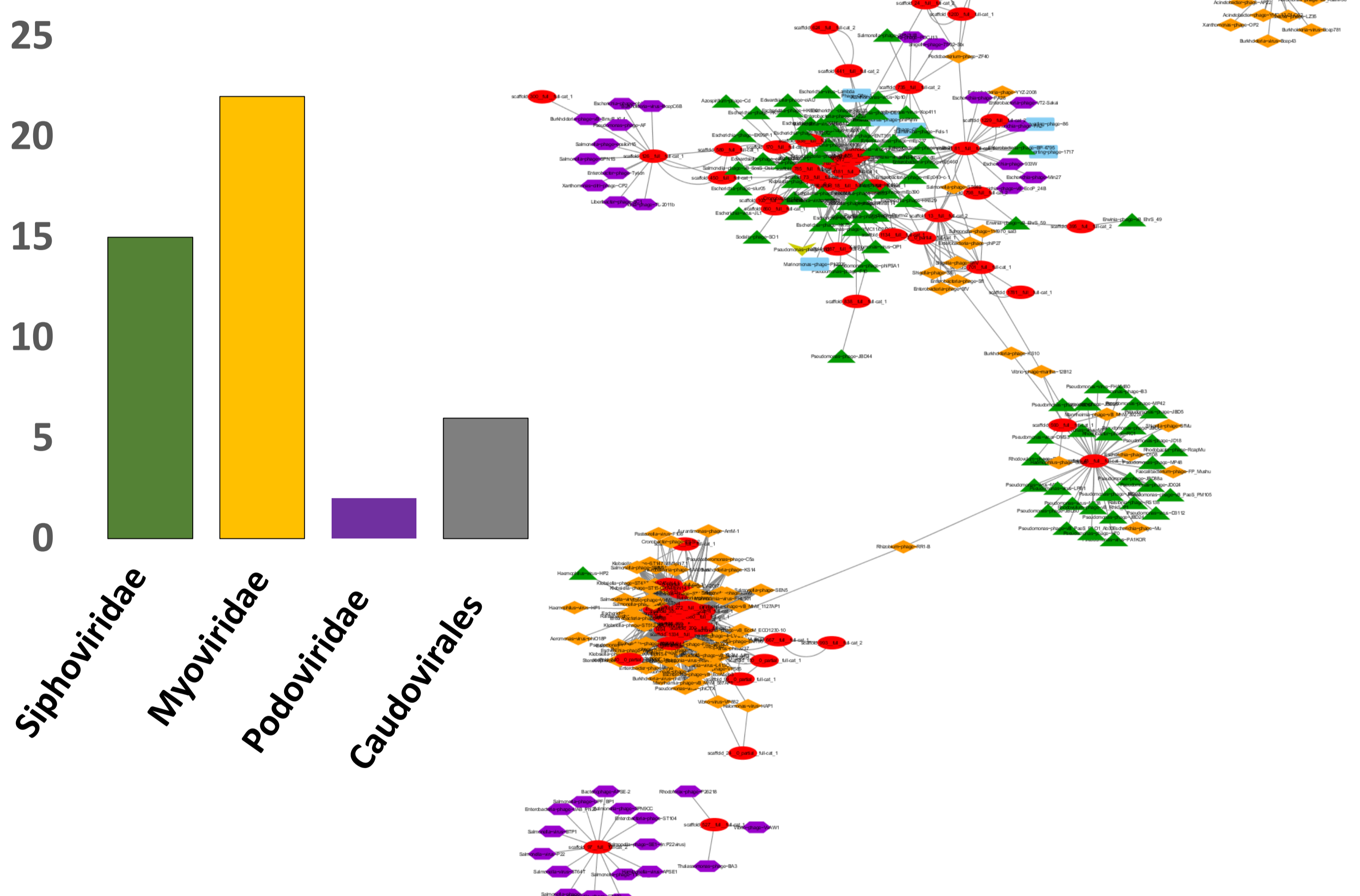


1. Introduction

- Microbes are key players in both plant health and plant disease (Ortíz-Castro et al, 2009).
- Examples include *Rhizobium*, which is essential for nitrogen fixing in leguminous plants (Gourion et al. 2014); and *Pantoea ananatis*, one of the pathogens responsible for bulb rot in onions (Asselin et al, 2018).
- Viruses have been shown to also play important roles in shaping the ecosystem around them (Rohwer et al, 2009).
- However, there is little information regarding viral ecology during crop disease development.

Aim: To characterize the endophytic viral populations of diseased onion bulbs using a metagenomics approach.

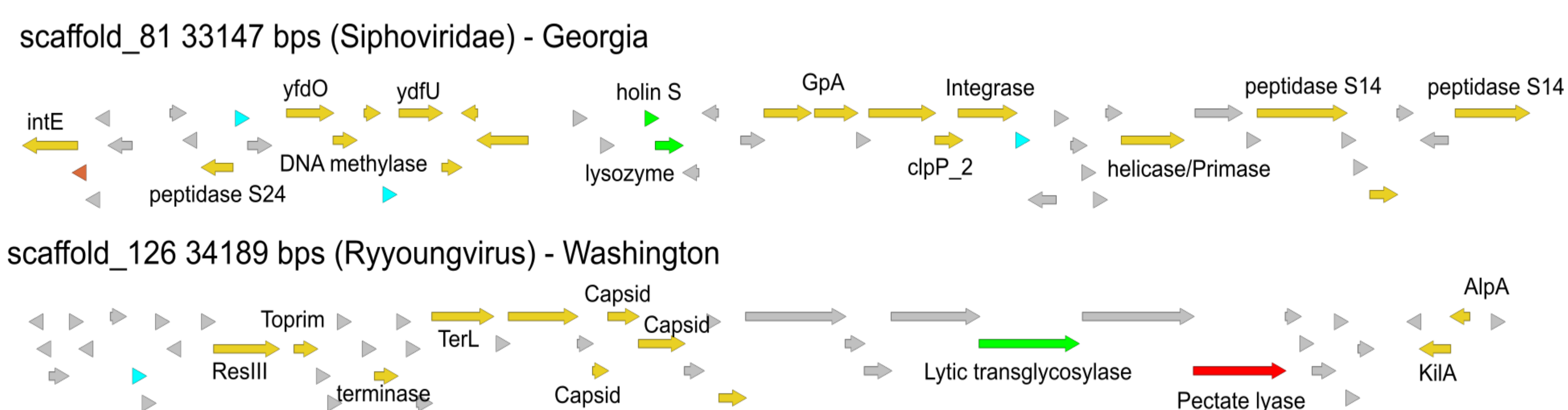
3. Results – Taxonomy of viral community:



- A total of 89 bacterial viral contigs were extracted from the diseased bulbs (no signal from healthy).
- 67 viral contigs (in red) were linked to known viruses

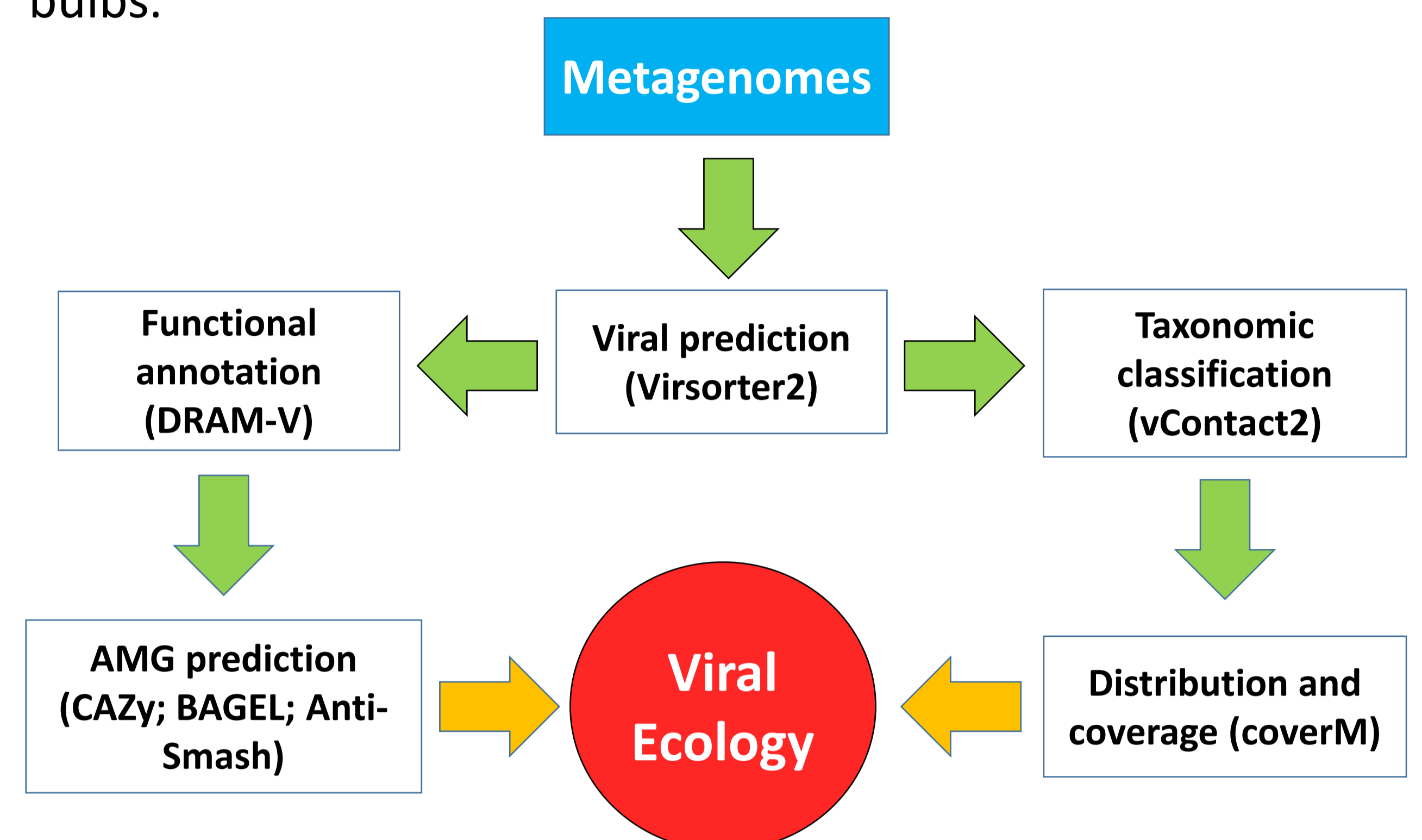
5. Results – Functional Annotations

- Several of the viral contigs contained gene markers for replication and evasion.
- These also include genes for lysis of the bacterial host (green) and that might help bacterial host pathogenicity (red).



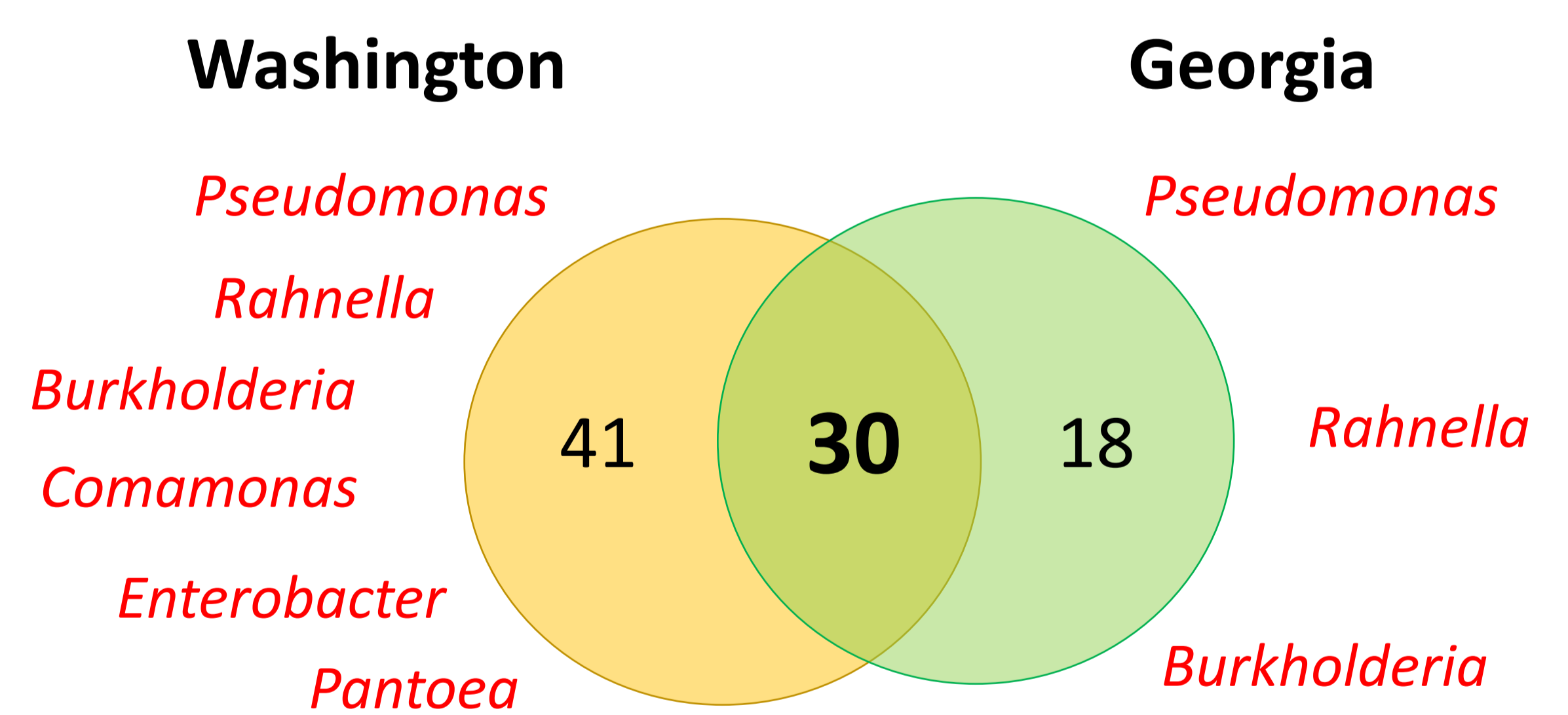
2. Methodology:

- Metagenomes from the endophyte community onion bulb cultivars stored in Washington State and Georgia facilities.
- Metagenomes were extracted from both healthy and diseased bulbs.



4. Results – Distribution of viral populations:

- Distinct populations in bulb cultivars from Georgia and Washington State (only 30 viruses are shared between cultivars).
- The viral population in bulbs from Washington cultivars are more diverse (41 unique viruses) and includes a broader range of hosts (indicated in red).



6. Conclusions

- High diversity of endophytic bacterial viruses in onion bulbs with bacterial rot.
- Results suggest that endophytic viral populations might be regionally distinct, reflecting the different bacterial hosts observed in cultivars from these regions.
- Endophytic viruses show marker genes that might help in bacterial host fitness and population control (biocontrol agents).

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References

- Asselin JAE, Bonasera JM, Beer SV. Center Rot of Onion (*Allium cepa*) Caused by *Pantoea ananatis* Requires pepM, a Predicted Phosphonate-Related Gene. *Mol Plant Microbe Interact*. 2018 Dec;31(12):1291-1300. doi: 10.1094/MPMI-04-18-0077-R. Epub 2018 Oct 24. PMID: 29953334.
- Gourion B, Berrabah F, Ratet P, Stacey G. Rhizobium-legume symbioses: the crucial role of plant immunity. *Trends Plant Sci*. 2015 Mar;20(3):186-94. doi: 10.1016/j.tplants.2014.11.008. Epub 2014 Dec 24. PMID: 25543258.
- Ortiz-Castro, R., Contreras-Cornejo, H. A., Macías-Rodríguez, L., & López-Bucio, J. (2009). The role of microbial signals in plant growth and development. *Plant signaling & behavior*, 4(8), 701–712. <https://doi.org/10.4161/psb.4.8.9047>
- Rohwer, F., Prangishvili, D. and Lindell, D. (2009). Roles of viruses in the environment. *Environmental Microbiology*, 11: 2771-2774. <https://doi.org/10.1111/j.1462-2920.2009.02101.x>

