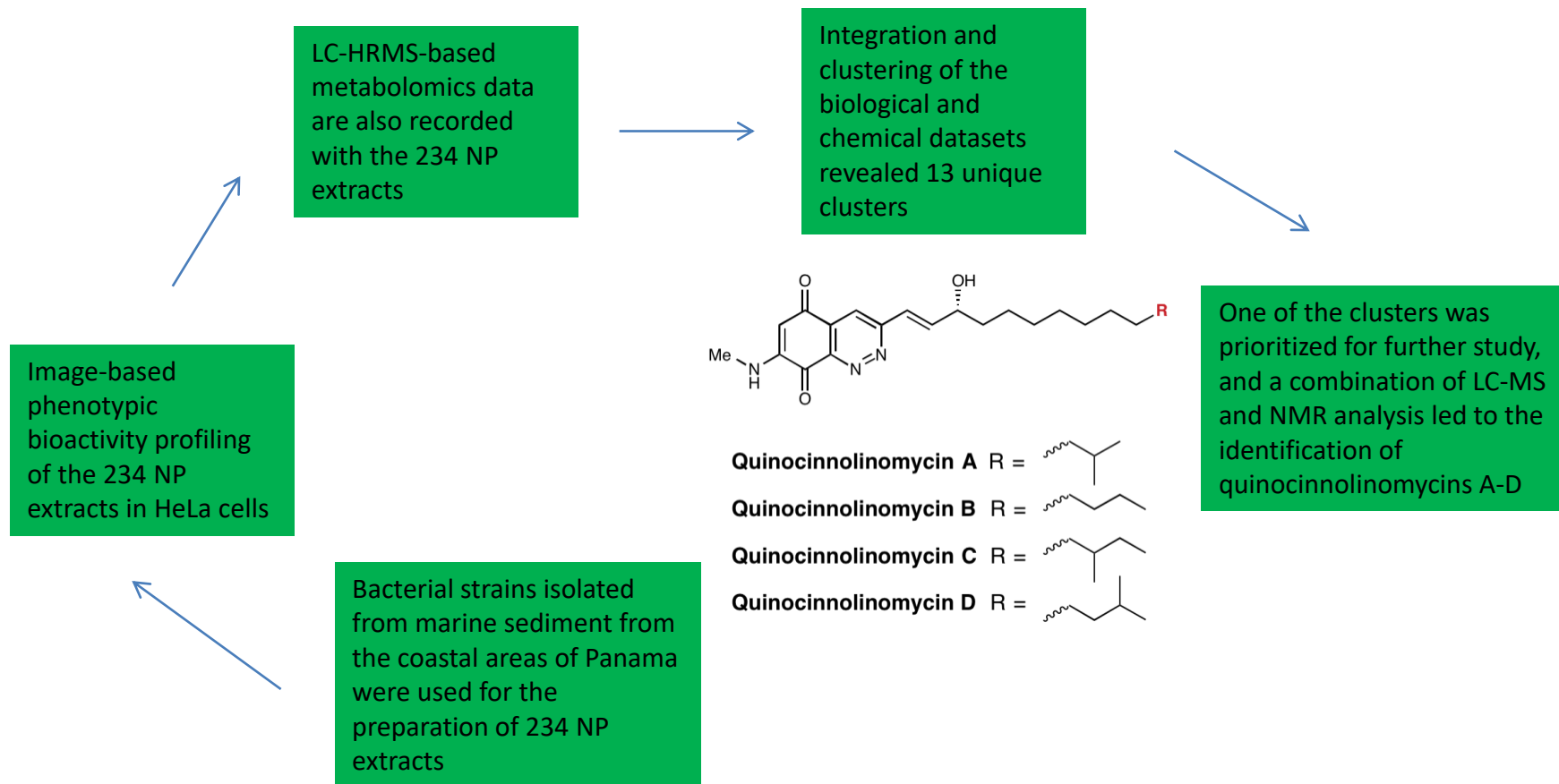
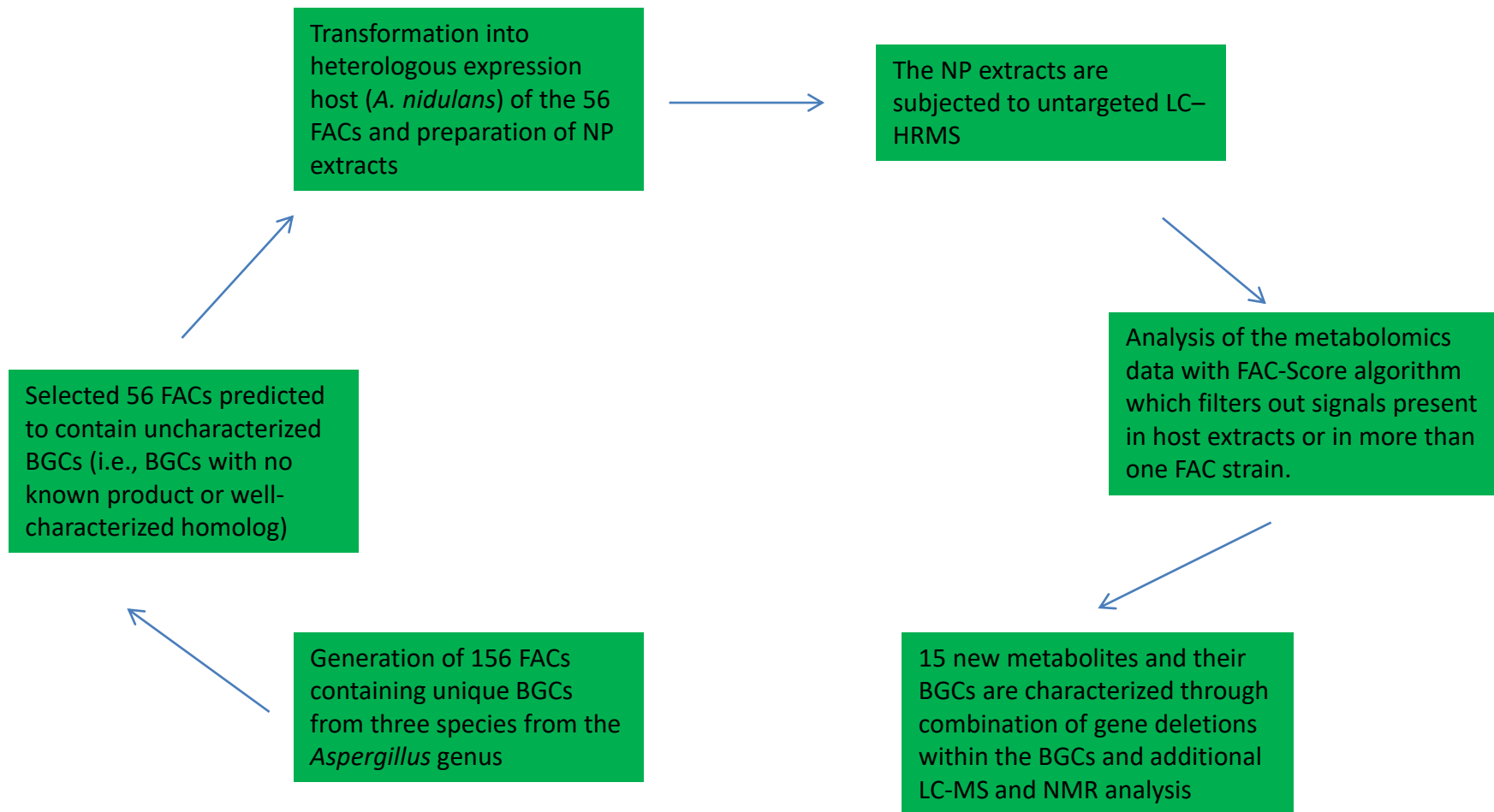


Figure 2A



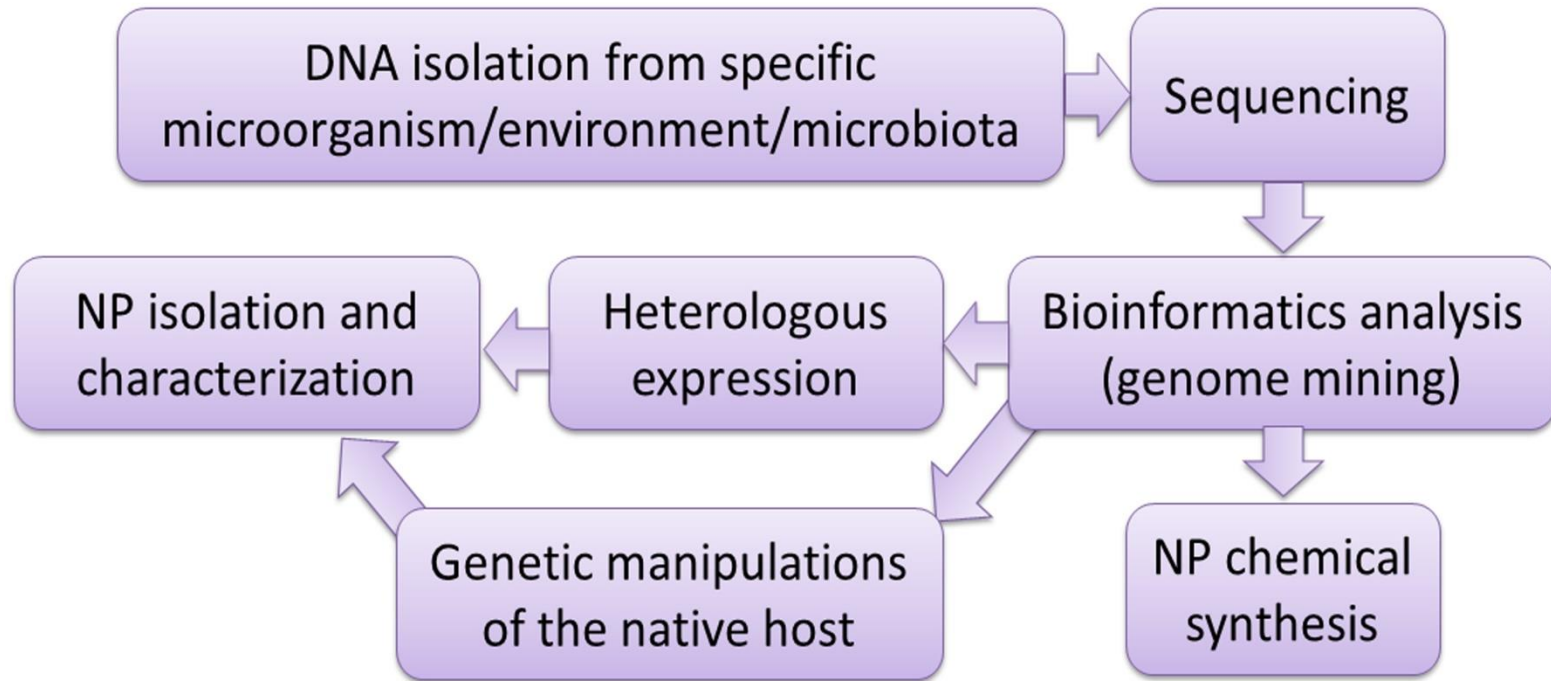
Kurita, K. L., Glassey, E. & Linington, R. G. Integration of high-content screening and untargeted metabolomics for comprehensive functional annotation of natural product libraries. *Proc. Natl. Acad. Sci. U. S. A.* 112, 11999–2004 (2015).

Figure 2B



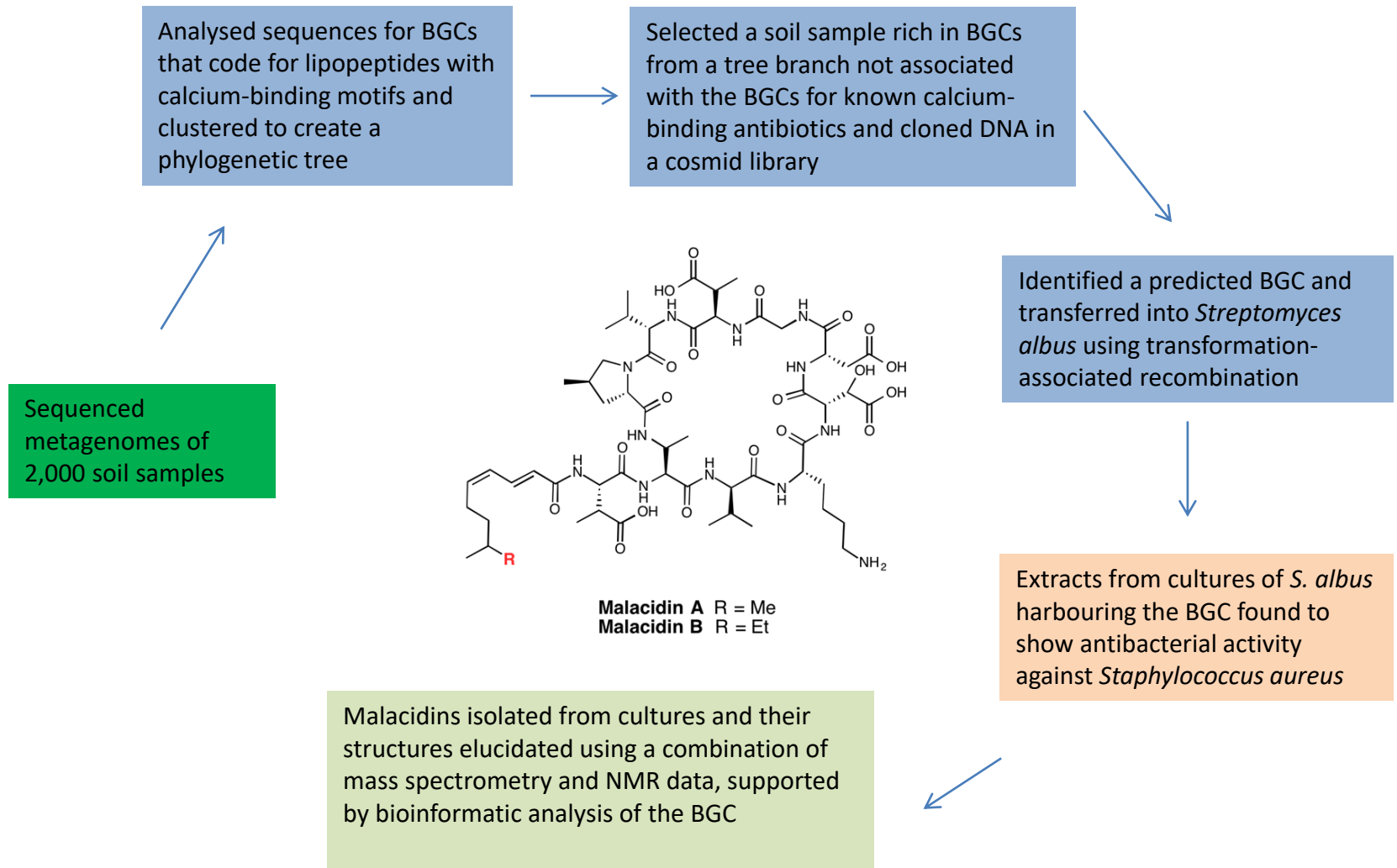
Clevenger, K. D. et al. A scalable platform to identify fungal secondary metabolites and their gene clusters. *Nat. Chem. Biol.* 13, 895–901 (2017).

Figure 3A



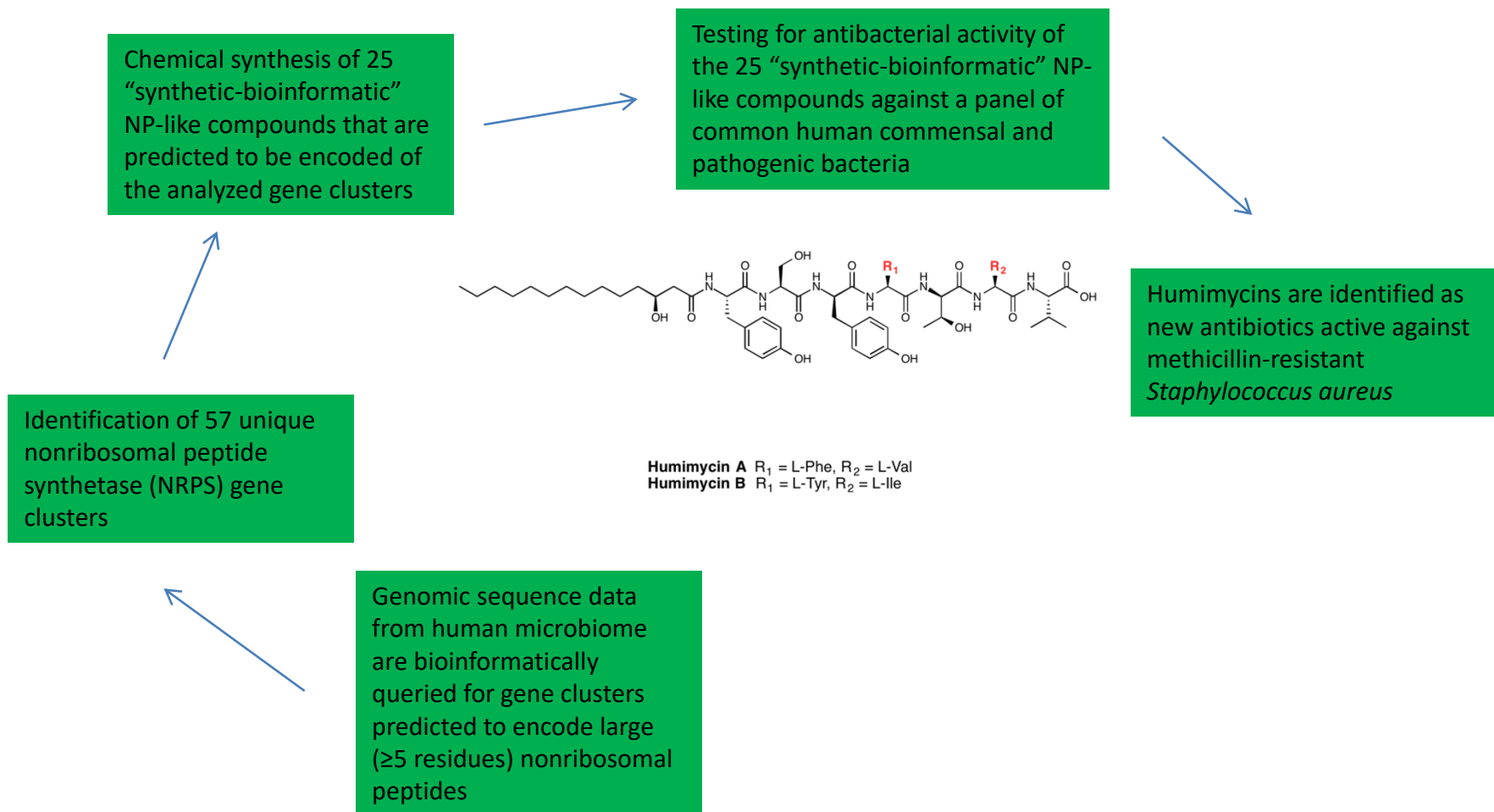
General scheme

Figure 3B



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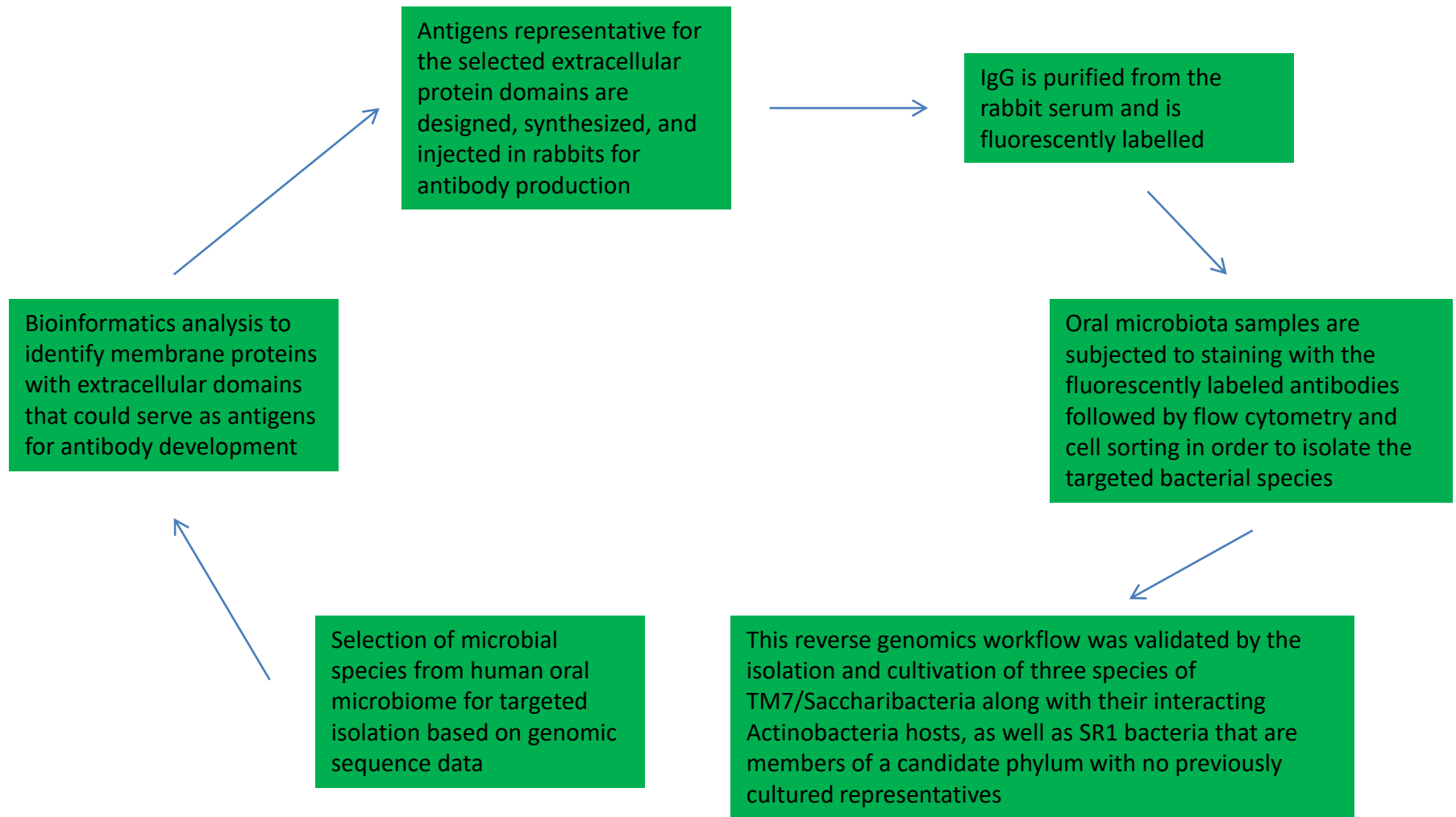
Figure 3C



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Figure 4B



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