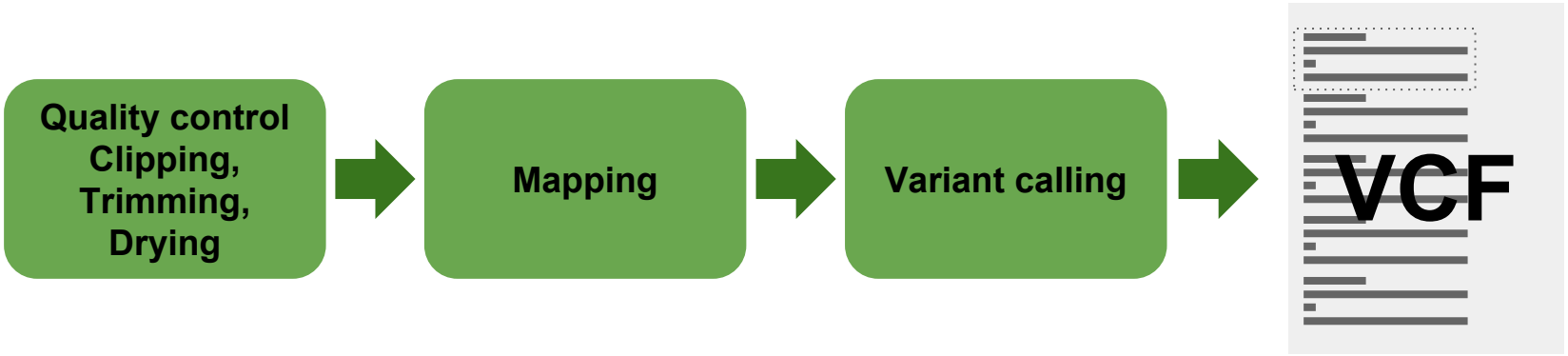
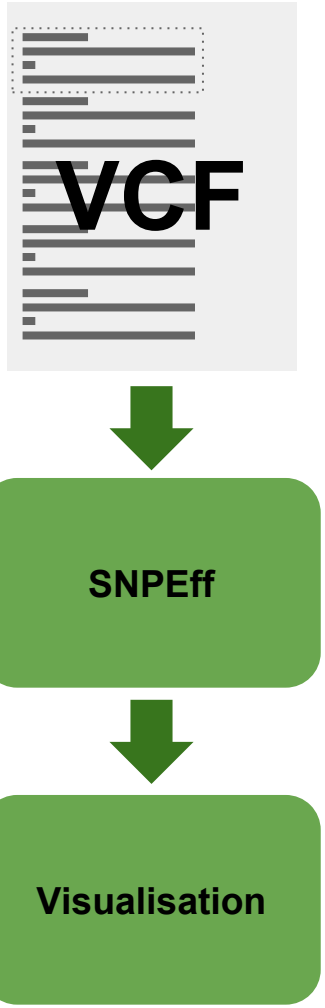
Call me, maybe - Integrating SNP visualisation from CandiSNP into Galaxy

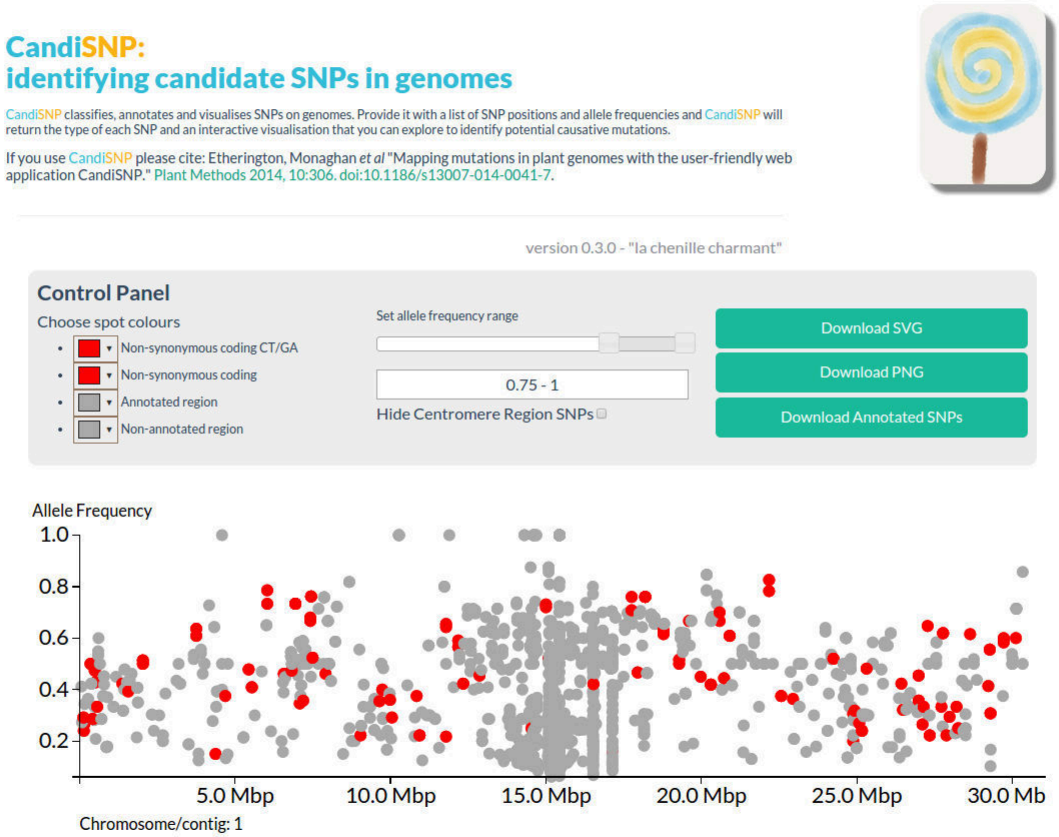
Christian Schudoma, Martin Page, Dan MacLean
Bioinformatics Team, The Sainsbury Laboratory

I got a VCF… and now?



Variant annotation and visualisation with SNPEff and CandiSNP

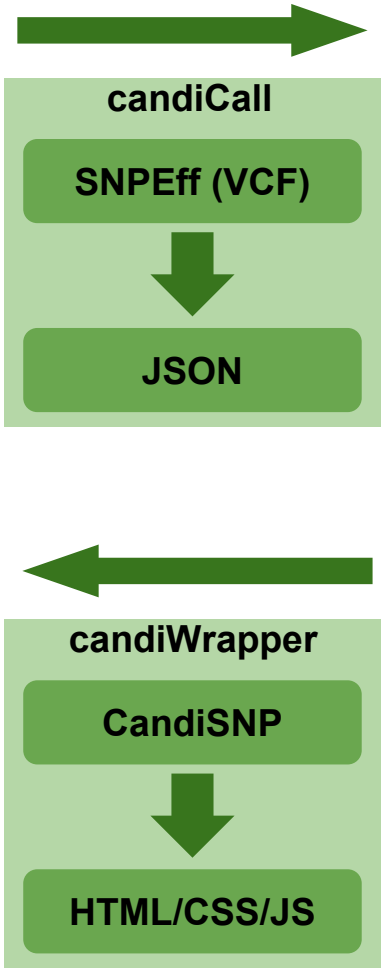




Etherington, Monaghan et al., Plant Methods (2014)

How to get Candi to go out with Galaxy?

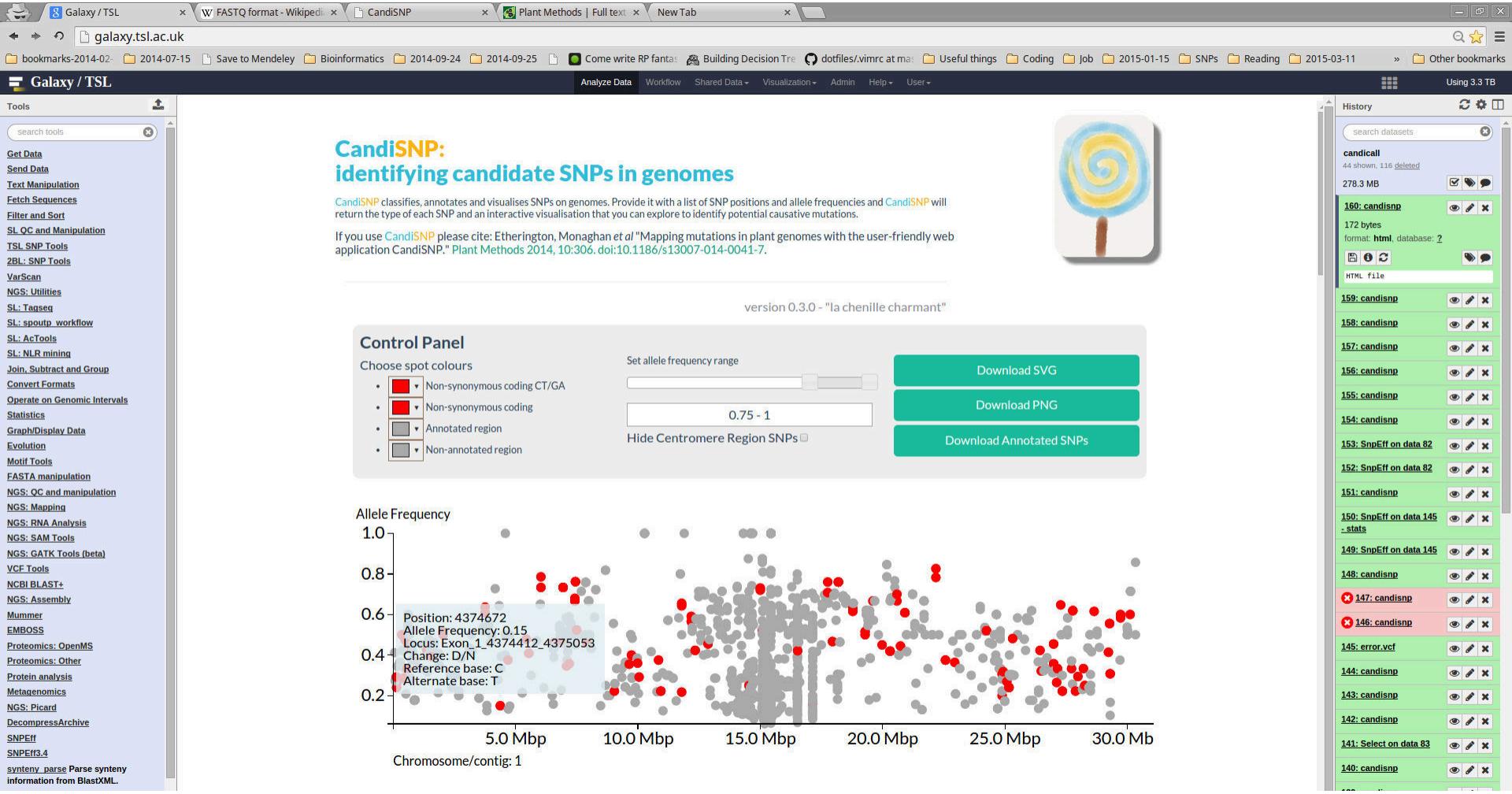
`sanitize_all_html = False`



http://candisnp.tsl.ac.uk

Look for candiSNP in the testtoolshed

Did it work out?



And so they'll live happily ever after..

Planned: full integration using BioJS visualisation in Galaxy

Acknowledgements

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