



Supplementary Fig. S2. Cartoon of maximum-likelihood topology of *rbcl* and *nrSSU* (nuclear-encoded small subunit ribosomal RNA) concatenated dataset of "Pyropia" species with 'Bangia 3' used as an outgroup, following Sutherland et al. (2011). Values on generic and inter-generic branches are displayed on branches or in triangles: non-parametric bootstrap % / Bayesian posterior probabilities / approximate likelihood-ratio test % / Ultrafast bootstrap %.