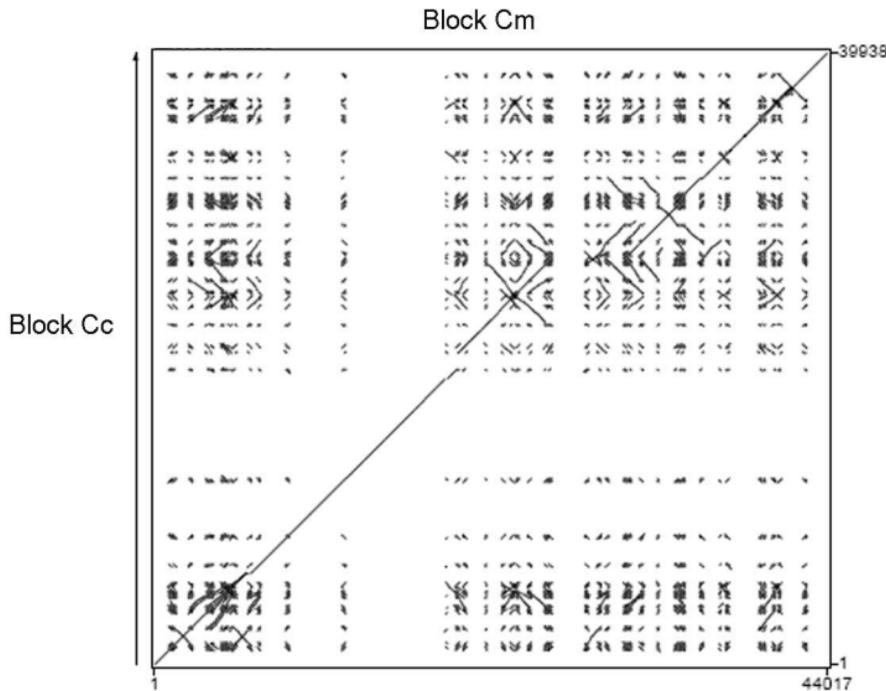


Supplementary Figure 1



Results of Pipmaker analysis. The homologous region (approx. 44Kb) between Blocks Cc and Cm was analysed. The Block Cc sequence has been plotted against a Block Cm sequence. Note the much higher homology of the first part of the blocks (approx. 17Kb; 98.8%) compared with the last 22kb (95.2%).

PIPMAKER analysis was performed at the web site <http://nog.cse.psu.edu/cgi-bin/pipmaker?advanced>, according to the authors' instructions .