1
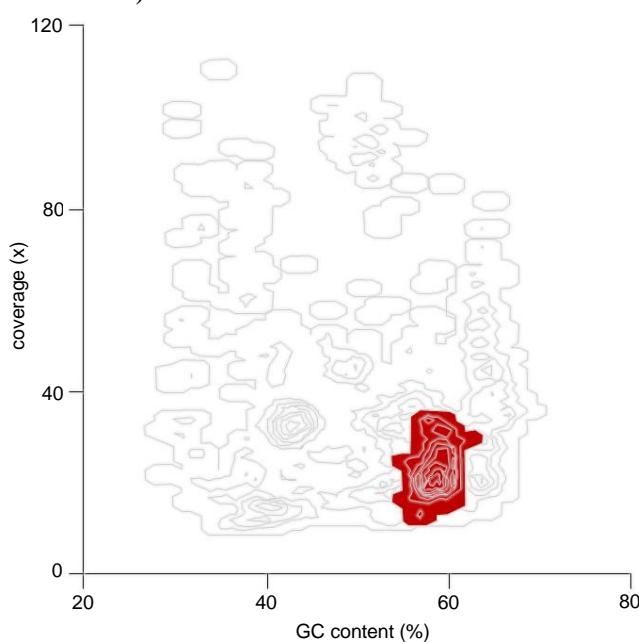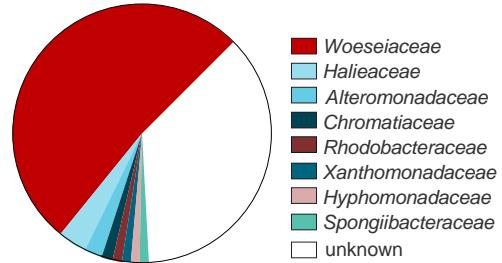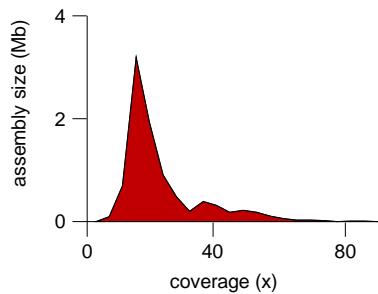

2

3   **Figure   S5.**   GC-content   vs.   sequence   coverage   plot   of   the   Janssand   sediment
4   metagenome   assembly.   A)   Highlighted   in   red   is   the   GC-content   vs.   coverage   area   of
5   bin JSS_woes1 belonging to the *Woeseiaceae*/JTB255. B) Taxonomic distribution of ORFs in bin
6   JSS_woes1. C) Coverage distribution of bin JSS_woes1 based on a 95% minimal identity mapping of
7   paired-end   sequence   reads   from   the   Janssand   metagenome.   Figures   were
8   generated in Metawatt (Strous *et al*., 2012).