Genomic comparison of the N-glycan processing locus 1 in *C. acnes* using the comparative genomics platform Sybil (http://sybil.sourceforge.net). The *C. acnes* 266 strain was used as reference genome (region 1545000-1564000). A subset of genomes was selected to represent the different *C. acnes* phylogenetic groups (IA1, IA2, IB, II and III). The GH genes of each locus are highlighted. Predicted function of other genes is given in Figure 2.