



Figure S4. Whole genome alignment of *Hermetia illucens* GCA_009835165.1 and iHerIII assemblies. Chromosome alignment of GCA_009835165.1 scaffolds to the iHerIII assembly both per chromosome (top left to bottom left) and whole genome overview (bottom right). Unique forward alignments = blue, unique reverse alignments = orange and repetitive sequences = green.