# Invertebrate methylomes provide insight into mechanisms of environmental tolerance and reveal methodological biases

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<sup>\*</sup> joint first authors/equal contribution

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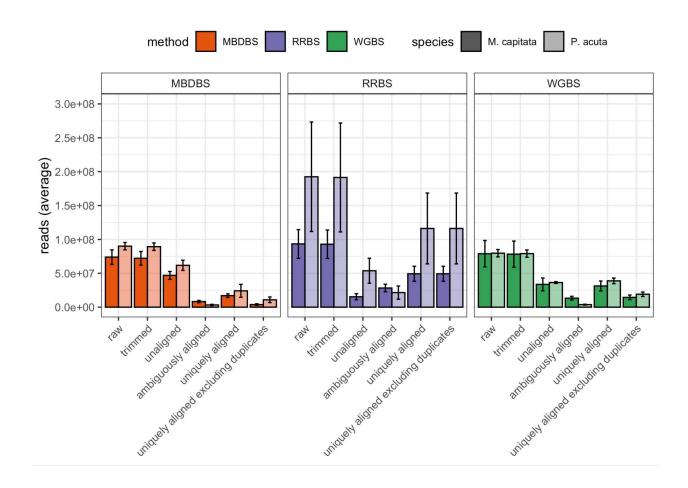
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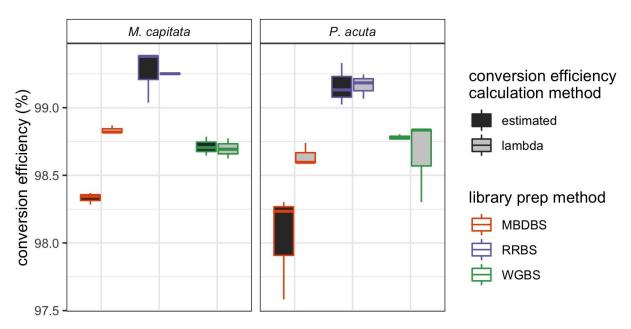
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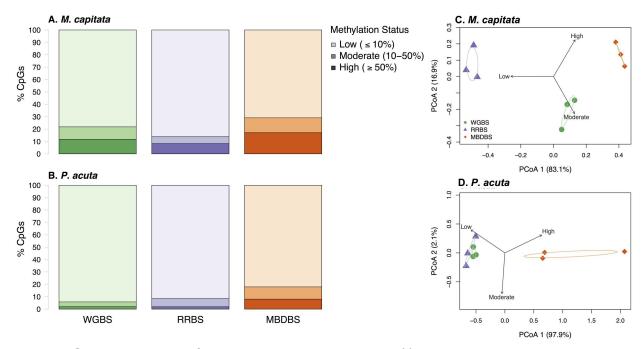
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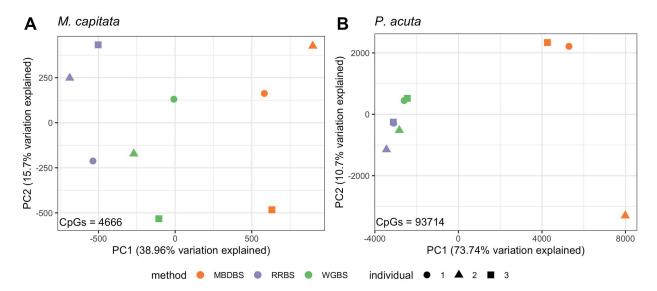
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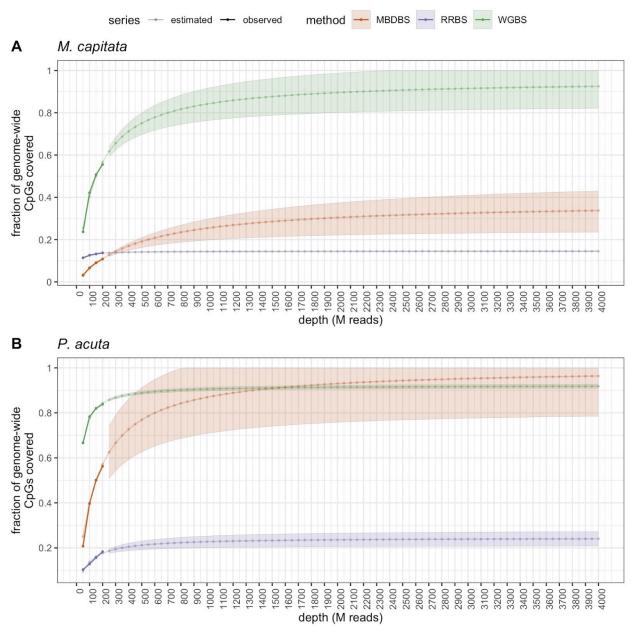
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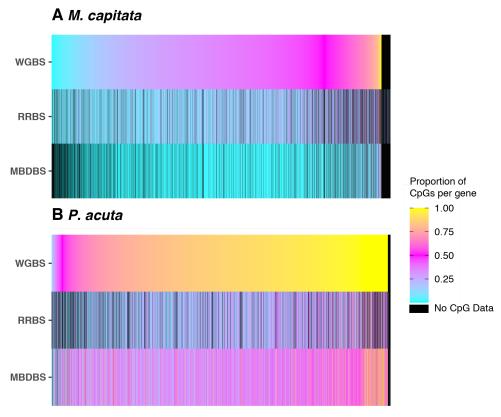
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**Figure SF6.** Mean proportion (n=3 samples per method) of CpGs per gene that have at least 5x coverage in all of the one-to-one orthologous genes, as identified by OrthoFinder (Putnam et al., 2020) for **A**) *M. capitata* and **B**) *P. acuta*.