Large-scale compositional analysis of invertebrate viromes estimated from public RNA-Seq data
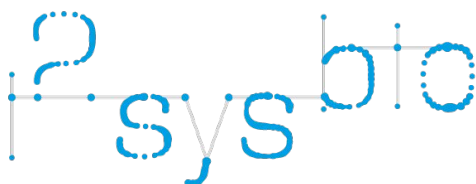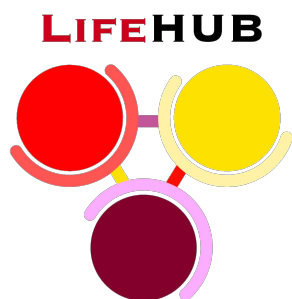

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