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Multicore Homology.

We design and implement a framework for parallel computation of homology of cellular spaces over field coefficients, by decomposing the space. Theoretically, we show that optimal decomposition into local pieces is NP-Hard. In practice, we achieve roughly an $8\times$ speedup of homology computation on a 3-dimensional complex with about 10 million simplices using 11 cores. (Received August 15, 2012)