Draft Genome Sequence of *Thermodesulfovibrio aggregans* TGE-P1\textsuperscript{T}, an Obligately Anaerobic, Thermophilic, Sulfate-Reducing Bacterium in the Phylum *Nitrospirae*

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We report a high-quality draft genome sequence of the type strain (TGE-P1\textsuperscript{T}) of *Thermodesulfovibrio aggregans*, an obligately anaerobic, thermophilic, sulfate-reducing bacterium in the phylum *Nitrospirae*. The genome comprises 2.00 Mb in 16 contigs (3 scaffolds), has a G+C content of 34.5%, and contains 1,998 predicted protein-encoding genes.
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