A bovine viral diarrhea virus (BVDV), GS5, of the BVDV-1a subgenotype was isolated from dairy cattle in Gansu Province, northwest China. Its near-full-length genome was determined to be closely related to an early Belgian BVDV-1a strain, WAX-N, but the relatedness to domestic strains is relatively low, indicating that different genetic evolution occurred between the viral strains in cattle in China.

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