A panel of 6 stomach cancer cell lines isolated from both primary and metastatic sites. Each culture contains genomic mutations in one or more of the following genes according to the Sanger COSMIC database: CDKN2A, TP53, SMAD4, CDH1, CTNNB1, KRAS, MLH1 and PIK3CA.

KATOIII, ATCC HTB-103
NCI-N87, ATCC CRL-5822
SNU-16, ATCC CRL-5974
SNU-5, ATCC CRL-5973
AGS, ATCC CRL-1739
SNU-1, ATCC CRL-5971

Appropriate safety procedures should always be used with this material. Laboratory safety is discussed in the current publication of the *Biosafety in Microbiological and Biomedical Laboratories* from the U.S. Department of Health and Human Services Centers for Disease Control and Prevention and National Institutes for Health.

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