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-5975
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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