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Introduction ===== Gastrointestinal stromal tumors (GISTs) are the most common mesenchymal tumors of the gastrointestinal tract. These tumors are characterized by the expression of KIT and/or PDGFRA \[@B1],[@B2]\]. They originate in the stomach (25%), small bowel (28%), colorectum (32%) and esophagus (2%). Most of the patients with GISTs are diagnosed at the early stage of the disease, but some patients, especially when the tumor is located in the small bowel, have been diagnosed with the advanced disease, because these tumors are usually difficult to diagnose by conventional endoscopy and radiologic examinations. Recently, molecular diagnosis of GIST has increased diagnostic accuracy, but because of the tumor heterogeneity, it is still difficult to obtain a tumor-specific diagnostic marker for GIST. In order to develop new tumor-specific molecular markers, it is useful to investigate the difference in gene expression profiles of GISTs compared with other gastrointestinal tumors and normal tissues. In this study, we isolated cDNA samples from GISTs and explored the gene expression profiles of GISTs compared with other gastrointestinal tumors and normal tissues using DNA microarray. We successfully identified KIT as a differentially expressed gene in GISTs. As KIT was previously reported as a diagnostic and therapeutic target for GIST, the identification of this gene was a significant step in the development of new molecular diagnostic markers for GIST.

Materials and methods ===== From 2006 to 2008, seven GISTs from the GI tract of Japanese patients were collected. Tissue samples were removed from surgical specimens, immediately frozen in

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