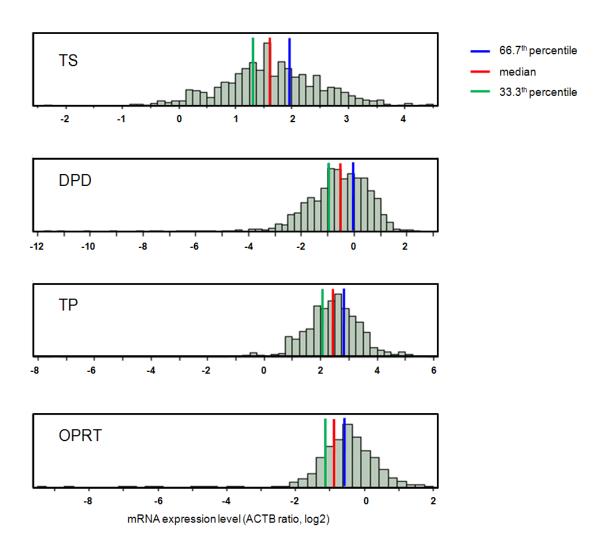
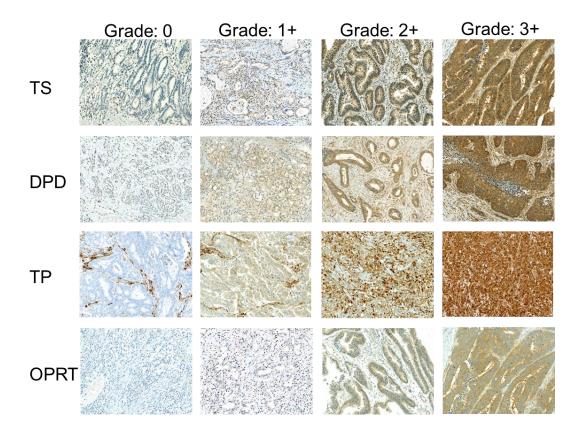
Supplementary Figure S1.



Supplementary Figure S1.

Histograms showing the distribution of TS, DPD, TP and OPRT gene expression measured by RT-PCR (x axis, expressed as log2). Each gene expression level at the 33.3rd, 50th, and 66.7th percentile was as follows: 2.47, 3.03, and 3.87 for TS; 0.50, 0.69, and 0.97 for DPD; 4.19, 5.44, and 7.09 for TP; and 0.45, 0.54, and 0.67 for OPRT, respectively.

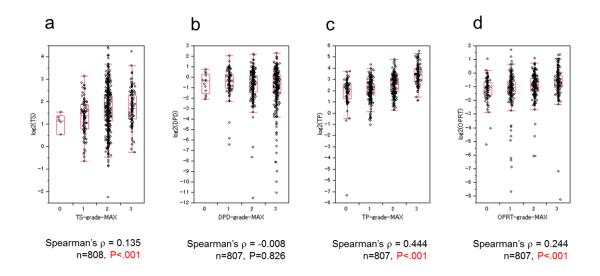
Supplementary Figure S2.



Supplementary Figure S2.

Representative examples of differently staged gastric carcinomas immunochemically stained for TS, DPD, TP and OPRT (magnification $\times 100$).

Supplementary Figure S3.

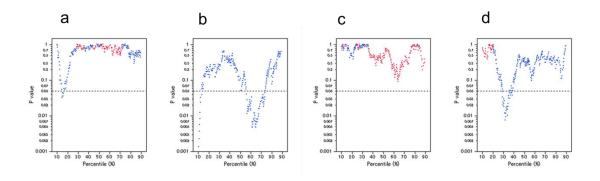


Supplementary Figure S3.

within 1.5 IQR of the upper quartile.

Relationship between TS, DPD, TP and OPRT expression levels in gastric tumors measured by RT-PCR (y axis, expressed as log2) and immunohistochemistry (IHC; x axis). a) Median TS mRNA levels for 4 IHC scores: 0, 1.15 (N=6); 1+, 1.38 (N=106); 2+, 1.61 (N=574); and 3+, 1.70 (N=122). b) Median DPD mRNA levels for 4 IHC scores: 0, -0.33 (N=13); 1+, -0.39 (N=102); 2+, -0.60 (N=258); and 3+, -0.53 (N=434). c) Median TP mRNA levels for 4 IHC scores: 0, 2.15 (N=91); 1+, 2.15 (N=290); 2+, 2.55 (N=295); and 3+, 3.43 (N=131). d) Median OPRT mRNA levels for 4 IHC scores: 0, -1.05 (N=86); 1+, -1.05 (N=218); 2+, -0.84 (N=271); and 3+, -0.68 (N=232). The Spearman's test was used to assess correlations between groups. Box plots show the 25th (bottom) and 75th (top) percentiles, and the median (middle); the lowest datum was within 1.5 IQR of the lower quartile, and the highest datum was

Supplementary Figure S4.



Supplementary Figure S4.

Correlations between cutoff values used for stratification and P values from log-rank tests for TS and DPD gene expression. a) P values for TS in the surgery-only group; b) P values for TS in the S-1-treated group; c) P values for DPD in the surgery-only group; d) P values for DPD in the S-1-treated group. Results below the dotted line (at P<0.05) are considered statistically significant.