

***FEN1* -96G>A and 4150G>T polymorphisms and cancer risk in Chinese population**

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Supplement Table S1 Frequency distribution of the *FEN1* -69G>A and 4150G>T haplotypes

First author	Case				Control			
	A <sub>-69</sub> T <sub>4150</sub>	G <sub>-69</sub> G <sub>4150</sub>	A <sub>-69</sub> G <sub>4150</sub>	G <sub>-69</sub> T <sub>4150</sub>	A <sub>-69</sub> T <sub>4150</sub>	G <sub>-69</sub> G <sub>4150</sub>	A <sub>-69</sub> G <sub>4150</sub>	G <sub>-69</sub> T <sub>4150</sub>
Lv, Z.	46	119	616	1419	11	17	1075	1697
Chen, Y. D.	243	286	1	4	478	1110	8	8
Liu, L.	1183	54	1317	1082	1736	43	1536	1126
Yang, M.	1251	2893	50	105	1558	2287	30	41

Supplement Table S2 Main Characteristics of *FEN1* -69G>A polymorphism and *FEN1* mRNA expression

First author	Tissue type	Assay method	Internal reference	Sample number	Genotype	<i>FEN1</i> mRNA expression (mean $\pm$ SE)	SD
Liu, L.	Liver	SYBR-Green real-time quantity PCR	$\beta$ -actin	5	AA	0.217 $\pm$ 0.038	0.0850
				15	AG+GG	0.099 $\pm$ 0.013	0.0503
	Esophagus			4	AA	0.242 $\pm$ 0.015	0.0300
				11	AG+GG	0.108 $\pm$ 0.042	0.1393
	Stomach			3	AA	0.268 $\pm$ 0.060	0.1039
				9	AG+GG	0.120 $\pm$ 0.035	0.1050
	Colon and rectum			3	AA	0.198 $\pm$ 0.034	0.0589
				10	AG+GG	0.091 $\pm$ 0.012	0.0379
Lv, Z.	Breast	SYBR-Green real-time quantity PCR	$\beta$ -actin	6	AA	0.193 $\pm$ 0.027	0.0661
				19	AG+GG	0.084 $\pm$ 0.010	0.0436