

Supplementary Table 7. Altered blood coagulation factors according to miRNA processing gene genotypes

Genotype	PLT		PT (sec)		aPTT (sec)		Fibrinogen		Antithrombin	
	Mean±SD (n)	CV (%)	Mean±SD (n)	CV (%)	Mean±SD (n)	CV (%)	Mean±SD (n)	CV (%)	Mean±SD (n)	CV (%)
<i>DICER1</i> rs13078 A>T										
AA	246.77±81.74 (879)	33.1	11.79±0.94 (765)	8.0	31.52±11.89 (765)	37.7	418.21±129.43 (601)	30.9	93.46±18.55 (605)	19.8
AT	247.18±63.58 (97)	25.7	11.67±0.77 (85)	6.6	31.1±6.63 (85)	21.3	439.63±120.52 (65)	27.4	93.85±18.42 (65)	19.6
TT	214.5±14.85 (2)	6.9	11.3±0.57 (2)	5.0	27.35±3.32 (2)	12.1	406.50±78.49 (2)	19.3	78.00±9.9 (2)	12.7
P	0.849		0.385		0.835		0.433		0.492	
<i>DICER1</i> rs3742330 A>G										
AA	246.99±92.07 (315)	37.3	11.85±1.12 (274)	9.5	31.13±6.16 (276)	19.8	410.30±120.95 (210)	29.5	93.08±15.17 (212)	16.3
AG	247.58±69.72 (454)	28.2	11.76±0.79 (396)	6.7	31.06±6.99 (395)	22.5	427.96±133.83 (309)	31.3	94.07±19.11 (310)	20.3
GG	244.58±81.77 (209)	33.4	11.73±0.86 (182)	7.3	32.87±21.3 (181)	64.8	418.45±127.35 (149)	30.4	92.67±21.41 (150)	23.1
P	0.903		0.308		0.178		0.299		0.708	
<i>DROSHA</i> rs6877842 C>G										
CC	246.18±79.98 (909)	32.5	11.78±0.92 (796)	7.8	31.39±11.21 (796)	35.7	419.72±130.14 (623)	31.0	93.49±18.42 (628)	19.7
CG	253.02±79.67 (67)	31.5	11.78±0.93 (55)	7.9	32.55±14.76 (55)	45.3	429.26±102.89 (44)	24.0	92.67±20.19 (43)	21.8
GG	295.5±144.96 (2)	49.1	11.5±0 (1)	0.0	35.4±0 (1)	0.0	406 (1)		105 (1)	
P	0.549		0.954		0.722		0.894		0.790	
<i>DROSHA</i> rs10719 T>C										
TT	246.71±75.2 (528)	30.5	11.75±0.79 (454)	6.7	31.07±7.06 (456)	22.7	419.98±160.20 (358)	38.1	94.67±17.64 (367)	18.6
TC	249.64±86.03 (387)	34.5	11.79±1.06 (341)	9.0	31.2±7.3 (339)	23.4	417.08±125.99 (262)	30.2	91.02±17.50 (257)	19.2
CC	229.29±80.02 (63)	34.9	11.94±1 (57)	8.4	36.23±35.28 (57)	97.4	439.99±160.20 (48)	36.4	97.32±27.29 (48)	28.0
P	0.173		0.352		0.005		0.525		0.017	
<i>RAN</i> rs14035 C>T										
CC	246.41±83.11 (603)	33.7	11.76±0.8 (531)	6.8	31.02±7.31 (531)	23.6	420.57±126.71 (419)	30.1	93.72±19.24 (421)	20.5
CT	246.1±75.26 (338)	30.6	11.78±1.07 (287)	9.1	32.49±16.96 (287)	52.2	419.74±134.66 (221)	32.1	92.03±15.98 (223)	17.4
TT	258.03±71.59 (37)	27.7	12.04±1.37 (34)	11.4	29.75±4.54 (34)	15.3	420.44±107.37 (28)	25.5	100.62±24.28 (28)	24.1
P	0.682		0.252		0.146		0.997		0.061	
<i>XPO5</i> rs11077 A>C										
AA	247.43±82.24 (810)	33.2	11.79±0.95 (708)	8.1	31.65±12.4 (708)	39.2	417.55±127.20 (561)	30.5	93.17±17.48 (562)	18.8
AC	244.47±68.65 (162)	28.1	11.7±0.76 (138)	6.5	30.42±4.51 (138)	14.8	433.61±132.90 (103)	30.6	95.03±23.41 (106)	24.6
CC	216.5±61.75 (6)	28.5	12.42±0.84 (6)	6.8	33.68±5.49 (6)	16.3	453.99±201.62 (4)	44.4	90.36±19.84 (4)	22.0
P	0.593		0.145		0.461		0.437		0.609	

P-value calculated by one-way analysis of variance test.
 PLT, platelet; PT, prothrombin time; aPTT, activated partial thromboplastin time; CV, coefficient of variation; RAN, Ran GTPase; XPO5, exportin 5.