

Supplementary table 1. Combined genotypes of *VEGFA* and *KDR* single nucleotide polymorphisms in high grade glioma cases and controls.

Genotypes	Cases (No, %)	Controls (No, %)	p value	OR (95% CI)
<i>VEGFA -2578C/A + KDR -604T/C</i>				
CC + TT	23 (21.7)	21 (18.6)	0.52	1.24 (0.63 – 2.46)
CA+AA + TC+CC	83 (78.3)	92 (81.4)		Reference
CC+CA + TT+TC	133 (96.4)	119 (93.7)	0.47	1.53 (0.47 – 4.99)
AA + CC	5 (3.6)	8 (6.3)		Reference
<i>VEGFA -2578C/A + KDR -271G/A</i>				
CC + AA	28 (25.5)	20 (17.7)	0.25	1.47 (0.75 – 2.90)
CA+AA + GG+GA	82 (74.5)	93 (82.3)		Reference
CC+CA + GA+AA	139 (97.2)	118 (92.2)	0.13	2.52 (0.75 – 8.41)
AA + GG	4 (2.8)	10 (7.8)		Reference
<i>VEGFA -2578C/A + KDR +1192G/A</i>				
CC + GG	75 (72.8)	53 (58.9)	0.09	1.69 (0.91 – 3.14)
CA+AA + GA+AA	28 (27.2)	37 (41.1)		Reference
CC+CA + GG+GA	186 (100.0)	161 (98.8)	NA	NA
AA + AA	0 (0.0)	2 (1.2)		Reference
<i>VEGFA -2578C/A + KDR +1719A/T</i>				
CC + TT	68 (63.0)	51 (56.7)	0.41	1.27 (0.71 – 2.28)
CA+AA + AA+AT	40 (37.0)	39 (43.3)		Reference
CC+CA + AT+TT	183 (100.0)	159 (98.1)	NA	NA

AA + AA	0 (0.0)	3 (1.9)		Reference
VEGFA -2489C/T + KDR -604T/C				
TT + TT	4 (2.7)	10 (7.8)	0.12	0.38 (0.11 – 1.29)
CC+CT + TC+CC	144 (97.3)	118 (92.2)		Reference
CT+TT + TT+TC	79 (71.8)	94 (81.7)	0.08	0.56 (0.29 – 1.08)
CC + CC	31 (28.2)	21 (18.3)		Reference
VEGFA -2489C/T + KDR -271G/A				
TT + AA	3 (2.1)	9 (7.0)	0.14	0.36 (0.09 – 1.40)
CC+CT + GG+GA	137 (97.9)	119 (93.0)		Reference
CT+TT + GA+AA	80 (76.2)	94 (79.7)	0.52	0.80 (0.42 – 1.55)
CC + GG	25 (23.8)	24 (20.3)		Reference
VEGFA -2489C/T + KDR +1192G/A				
TT + GG	13 (22.8)	27 (35.1)	0.06	0.40 (0.17 – 0.90)
CC+CT + GA+AA	44 (77.2)	50 (64.9)		Reference
CT+TT + GG+GA	108 (98.2)	125 (96.9)	0.59	1.63 (0.27 – 9.70)
CC + AA	2 (1.8)	4 (3.1)		Reference
VEGFA -2489C/T + KDR +1719A/T				
TT + TT	18 (23.1)	30 (40.0)	0.05	0.36 (0.16 – 0.80)
CT+TT + AA+AT	60 (76.9)	50 (60.0)		Reference
CT+TT + AT+TT	104 (99.0)	122 (96.8)	0.18	4.49 (0.48 – 41.87)
CC + AA	1 (1.0)	4 (3.2)		Reference
VEGFA -1154G/A + KDR -604T/C				

GG + TT	30 (32.3)	26 (26.0)	0.32	1.37 (0.73 – 2.58)
GA+AA + TC+CC	63 (67.7)	74 (74.0)		Reference
GG+GA + TT+TC	0 (0.0)	0 (0.0)	NA	NA
AA + CC	136 (100.0)	132 (100.0)		Reference
VEGFA -1154G/A + KDR -271G/A				
GG + AA	35 (36.1)	22 (23.4)	0.13	1.65 (0.85 – 3.18)
GA+AA + GG+GA	62 (63.9)	72 (76.6)		Reference
GG+GA + GA+AA	140 (97.9)	128 (97.0)	0.79	1.23 (0.26 – 5.78)
AA + GG	3 (2.1)	4 (3.0)		Reference
VEGFA -1154G/A + KDR +1192G/A				
GG + GG	92 (83.6)	70 (69.3)	0.05	1.97 (0.99 – 3.90)
GA+AA + GA+AA	18 (16.4)	31 (30.7)		Reference
GG+GA + GG+GA	188 (100.0)	178 (98.3)	NA	NA
AA + AA	0 (0.0)	3 (1.7)		Reference
VEGFA -1154G/A + KDR +1719A/T				
GG + TT	83 (74.8)	69 (67.0)	0.25	1.42 (0.77 – 2.62)
GA+AA + AA+AT	28 (25.2)	34 (33.0)		Reference
GG+GA + AT+TT	185 (100.0)	172 (100.0)	NA	NA
AA + AA	0 (0.0)	0 (0.0)		Reference
VEGFA -634G/C + KDR -604T/C				
CC + TT	7 (5.2)	5 (3.7)	0.54	1.46 (0.42 – 5.00)
GC+CC + TC+CC	128 (94.8)	129 (96.3)		Reference

CC+GC + TT+TC	97 (84.3)	88 (76.5)	0.14	1.66 (0.83 – 3.29)
GG + CC	18 (15.7)	27 (23.5)		Reference

VEGFA -634G/C + KDR -271G/A

CC + AA	6 (4.7)	5 (3.7)	0.73	1.25 (0.35 – 4.46)
GG+GC + GG+GA	121 (95.3)	131 (96.3)		Reference
GC+CC + GA+AA	101 (87.1)	82 (77.4)	0.13	1.74 (0.84 – 3.60)
GG + GG	15 (12.9)	24 (22.6)		Reference

VEGFA -634G/C + KDR +1192G/A

CC + GG	30 (40.5)	20 (26.0)	0.14	1.69 (0.82 – 3.46)
GG+ GC + GA+AA	44 (59.5)	57 (74.0)		Reference
GC+CC + GG+GA	138 (99.3)	113 (96.6)	0.19	4.45 (0.46 – 42.80)
GG + AA	1 (0.7)	4 (3.4)		Reference

VEGFA -634G/C + KDR +1719A/T

CC + TT	25 (30.1)	17 (22.7)	0.38	1.38 (0.66 – 2.88)
GG+GC + AA+AT	58 (69.9)	58 (77.3)		Reference
GC+CC + AT+TT	136 (98.6)	110 (96.5)	0.40	2.09 (0.37 – 11.86)
GG + AA	2 (1.4)	4 (3.5)		Reference

VEGFA -460C/T + KDR -604T/C

TT + TT	19 (17.0)	20 (17.2)	0.97	0.99 (0.49 – 2.00)
CC+CT + TC+CC	93 (83.0)	96 (82.8)		Reference
CT+TT + TT+TC	124 (94.7)	116 (90.6)	0.21	1.87 (0.69 – 5.08)

CC + CC	7 (5.3)	12 (9.4)		Reference
VEGFA -460C/T + KDR -271G/A				
TT + AA	21 (19.1)	17 (15.2)	0.63	1.19 (0.57 – 2.49)
CC+CT + GG+GA	89 (80.9)	95 (84.8)		Reference
CT+TT + GA+AA	132 (94.3)	112 (91.1)	0.49	1.40 (0.53 – 3.68)
CC + GG	8 (5.7)	11 (8.9)		Reference
VEGFA -460C/T + KDR +1192G/A				
TT + GG	64 (67.4)	50 (56.2)	0.18	1.50 (0.81 – 2.77)
CC+CT + GA+AA	31 (32.6)	39 (43.8)		Reference
CT+TT + GG+GA	175 (100.0)	154 (98.7)	NA	NA
CC + AA	0 (0.0)	2 (1.3)		Reference
VEGFA -460C/T + KDR +1719A/T				
TT + TT	57 (57.0)	48 (53.9)	0.72	1.11 (0.62 – 1.99)
CC+CT + AA+AT	43 (43.0)	41 (46.1)		Reference
CT+TT + AT+TT	173 (99.4)	152 (98.1)	0.37	2.84 (0.28 – 28.14)
CC + AA	1 (0.6)	3 (1.9)		Reference
KDR -604T/C + -271G/A				
TT + AA	0 (0.0)	3 (3.0)	NA	NA
TC+CC + GG+GA	103 (100.0)	96 (97.0)		Reference
TT+TC + GA+AA	91 (98.9)	92 (97.9)	0.47	2.46 (0.20 – 30.13)
CC + GG	1 (1.1)	2 (2.1)		Reference

KDR -604T/C + +1192G/A

TT + GG	43 (48.9)	42 (47.7)	0.85	1.05 (0.58 – 1.93)
TC+CC + GA+AA	45 (51.1)	46 (52.3)		Reference
TT+TC + GG+GA	143 (98.6)	145 (97.3)	0.85	1.18 (0.19 – 7.14)
CC + AA	2 (1.4)	4 (2.7)		Reference

KDR -604T/C + +1719A/T

TT + TT	26 (35.6)	38 (45.2)	0.36	0.73 (0.37 – 1.42)
TC+CC + AA+AT	47 (64.4)	46 (54.8)		Reference
TT+TC + AT+TT	139 (99.3)	143 (96.6)	0.16	4.65 (0.52 – 41.45)
CC + AA	1 (0.7)	5 (3.4)		Reference

KDR -271G/A + +1192G/A

AA + GG	35 (53.0)	36 (46.2)	0.70	1.14 (0.57 – 2.26)
GG+GA + GA+AA	31 (47.0)	42 (53.8)		Reference
GA+AA + GG+GA	149 (99.3)	140 (98.6)	0.42	2.71 (0.23 – 30.83)
GG + AA	1 (0.7)	2 (1.4)		Reference

KDR -271G/A + +1719A/T

AA + TT	39 (41.9)	37 (44.0)	0.72	0.89 (0.48 – 1.64)
GG+GA + AA +AT	54 (58.1)	47 (56.0)		Reference
GA+AA + AT+TT	146 (99.3)	136 (99.3)	0.79	1.48 (0.07 – 27.86)

GG + AA	1 (0.7)	1 (0.7)		Reference
KDR +1192G/A + 1719A/T				
GG + TT	114 (80.3)	110 (78.0)	0.75	1.10 (0.60 – 1.98)
GA+AA + AA+AT	28 (19.7)	31 (22.0)		Reference
GG+GA + AT+TT	197 (99.5)	186 (100.0)	NA	NA
AA + AA	1 (0.5)	0 (0.0)		Reference

No, number, %: percent, OR: odds ratio, CI: confidence interval. Genotypes were analyzed according to the functional role of the alleles, and combined genotypes associated with lower VEGF or KDR production or lower binding efficiency of KDR for VEGFA as references. In cases of VEGFRA -634G/C and -460C/T, GG and CC genotypes were taken as references in analyses, respectively, since GC+CC and CT+TT genotypes were associated with increased risks of high grade gliomas in the present study.

Supplementary table 2. Genotypes of VEGFA and KDR single nucleotide polymorphisms in clinicopathological features of high grade glioma cases.

Haplotypes*	Age		p value	Gender		p value	Histological grade*		p value
	≤ 54 (No, %)	> 54 (No, %)		Male (No, %)	Female (No, %)		III (No, %)	IV (No, %)	
VEGFA -2578C/A									
CC	49 (48.5)	48 (46.2)		66 (48.9)	31 (44.3)		17 (47.2)	80 (47.6)	
CA+AA	52 (51.5)	56 (53.8)	0.735	69 (51.1)	39 (55.7)	0.531	19 (52.8)	88 (52.4)	0.965
CC+CA	95 (94.1)	94 (90.4)		124 (91.9)	65 (92.9)		35 (97.2)	153 (91.1)	
AA	6 (5.9)	10 (9.6)	0.437	11 (8.1)	5 (7.1)	1.000	1 (2.8)	15 (8.9)	0.315
VEGFA -2489C/T									
CC	48 (47.5)	48 (46.2)		65 (48.1)	31 (44.3)		17 (47.2)	79 (47.0)	

CT+TT	53 (52.5)	56 (53.8)	0.844	70 (51.9)	39 (55.7)	0.599	19 (52.8)	89 (53.0)	0.983
CC+CT	95 (94.1)	93 (89.4)		123 (91.1)	65 (92.9)		35 (97.2)	152 (90.5)	
TT	6 (5.9)	11 (10.6)	0.312	12 (8.9)	5 (7.1)	0.793	1 (2.8)	16 (9.5)	0.317
VEGFA -1154G/A									
GG	58 (57.4)	66 (63.5)		83 (61.5)	41 (58.6)		20 (55.6)	104 (61.9)	
GA+AA	43 (42.6)	38 (36.5)	0.377	52 (38.5)	29 (41.4)	0.686	16 (44.4)	64 (38.1)	0.479
GG+GA	94 (93.1)	97 (93.3)		128 (94.8)	63 (90.0)		33 (91.7)	157 (93.5)	
AA	7 (6.9)	7 (6.7)	1.000	7 (5.2)	7 (10.0)	0.244	3 (8.3)	11 (6.5)	0.717
VEGFA -634G/C									
GG	29 (28.7)	36 (34.6)		39 (28.9)	26 (37.1)		9 (25.0)	55 (32.7)	
GC+CC	72 (71.3)	68 (65.4)	0.364	96 (71.1)	44 (62.9)	0.228	27 (75.0)	113 (67.3)	0.432
GG+GC	83 (82.2)	86 (82.7)		111 (82.2)	58 (82.9)		30 (83.3)	138 (82.1)	
CC	18 (17.8)	18 (17.3)	0.923	24 (17.8)	12 (17.1)	0.910	6 (16.7)	30 (17.9)	1.000
VEGFA -460C/T									
CC	11 (10.9)	16 (15.4)	0.342	18 (13.3)	9 (12.9)	1.000	1 (2.8)	25 (14.9)	0.054
CT+TT	90 (89.1)	88 (84.6)		117 (86.7)	61 (87.1)		35 (97.2)	143 (85.1)	
CC+CT	60 (59.4)	62 (59.6)	0.976	81 (60.0)	41 (58.6)	0.843	22 (61.1)	99 (58.9)	0.809
TT	41 (40.6)	42 (40.4)		54 (40.0)	29 (41.4)		14 (38.9)	69 (41.1)	
KDR -604T/C									
TT	27 (26.7)	21 (20.2)	0.323	30 (22.2)	18 (25.7)	0.576	12 (33.3)	35 (20.8)	0.106
TC+CC	74 (73.3)	83 (79.8)		105 (77.8)	52 (74.3)		24 (66.7)	133 (79.2)	
TT+TC	72 (71.3)	72 (69.2)	0.747	99 (73.3)	45 (64.3)	0.179	27 (75.0)	116 (69.0)	0.551

CC	29 (28.7)	32 (30.8)		36 (26.7)	25 (35.7)		9 (25.0)	52 (31.0)
<i>KDR -271G/A</i>								
GG	28 (27.7)	26 (25.0)	0.658	34 (25.2)	20 (28.6)	0.602	13 (36.1)	40 (23.8)
GA+AA	73 (72.3)	78 (75.0)		101 (74.8)	50 (71.4)		23 (63.9)	128 (76.2)
GG+GA	78 (77.2)	73 (70.2)	0.253	101 (74.8)	50 (71.4)	0.602	29 (80.6)	121 (72.0)
AA	23 (22.8)	31 (29.8)		34 (25.2)	20 (28.6)		7 (19.4)	47 (28.0)
<i>KDR +1192G/A</i>								
GG	75 (74.3)	80 (76.9)		106 (78.5)	49 (70.0)		28 (77.8)	126 (75.0)
GA+AA	26 (25.7)	24 (23.1)	0.657	29 (21.5)	21 (30.0)	0.178	8 (22.2)	42 (25.0)
GG+GA	100 (99.0)	102 (98.1)		133 (98.5)	69 (98.6)		35 (97.2)	166 (98.8)
AA	1 (1.0)	2 (1.9)	1.000	2 (1.5)	1 (1.4)	1.000	1 (2.8)	2 (1.2)
<i>KDR +1719A/T</i>								
AA	3 (3.0)	3 (2.9)	1.000	5 (3.7)	1 (1.4)	0.666	2 (5.6)	4 (2.4)
AT+TT	98 (97.0)	101 (97.1)		130 (96.3)	69 (98.6)		34 (94.4)	164 (97.6)
AA+AT	32 (31.7)	37 (35.6)	0.555	40 (29.6)	29 (41.4)	0.090	13 (36.1)	55 (32.7)
TT	69 (68.3)	67 (64.4)		95 (70.4)	41 (58.6)		23 (63.9)	113 (67.3)

No: number; %, percent. *: histological grade classified according to World Health Organization criteria, where grade III included anaplastic astrocytoma and oligoastrocytoma and grade IV included glioblastoma multiforme; the histological grade was not identified in one patient.

Supplementary table 3. Combined genotypes of *VEGFA* and *KDR* single nucleotide polymorphisms in clinicopathological features of high grade gliomas cases.

CC + TT	41 (44.1)	42 (42.9)		54 (43.9)	29 (42.6)		14 (42.4)	69 (43.9)	
CA+AA + CC+CT	52 (55.9)	56 (57.1)	0.88	69 (56.1)	39 (57.4)	0.88	19 (57.6)	88 (56.1)	1.00
CC+CA + CT+TT	90 (93.8)	88 (89.8)		117 (91.4)	61 (92.4)		35 (97.2)	143 (90.5)	
AA + CC	6 (6.3)	10 (10.2)	0.43	11 (8.6)	5 (7.6)	1.00	1 (2.8)	15 (9.5)	0.31

VEGFA -2489C/T + -1154G/A

TT + GG	1 (2.6)	1 (3.4)		2 (4.5)	0 (0)		0 (0)	2 (3.8)	
CC+CT + GA+AA	38 (97.4)	28 (96.6)	1.00	42 (95.5)	24 (100)	0.53	15 (100)	50 (96.2)	1.00
CT+TT + GG+GA	48 (96.0)	52 (94.5)		64 (98.5)	36 (90.0)		17 (94.4)	82 (95.3)	
CC + AA	2 (4.0)	3 (5.5)	1.00	1 (1.5)	4 (10.0)	0.06	1 (5.6)	4 (4.7)	1.00

VEGFA -2489C/T + -634G/C

TT + CC	0 (0)	0 (0)		0 (0)	0 (0)		0 (0)	0 (0)	
CC+CT + GG+GC	77 (100)	75 (100)	1.00	99 (100)	53 (100)	1.00	29 (100)	122 (100)	1.00
TT+CT + GC+CC	28 (87.5)	23 (88.5)		35 (89.7)	16 (84.2)		11 (91.7)	40 (87.0)	
CC + GG	4 (12.5)	3 (11.5)	1.00	4 (10.3)	3 (15.8)	0.67	1 (8.3)	6 (13.0)	1.00

VEGFA -2489C/T + -460C/T

TT + TT	0 (0)	0 (0)		0 (0)	0 (0)		0 (0)	0 (0)	
CC+CT + CC+CT	54 (100)	51 (100)	1.00	69 (100)	36 (100)	1.00	21 (100)	83 (100)	1.00
CT+TT + CT+CT	42 (100)	40 (100)		52 (100)	30 (100)		18 (100)	64 (100)	
CC + CC	0 (0)	0 (0)	1.00	0 (0)	0 (0)	1.00	0 (0)	0 (0)	1.00

VEGFA -1154G/A + -634G/C

GG + CC	16 (28.1)	15 (30.0)		23 (31.1)	8 (24.2)		6 (27.3)	25 (29.8)	
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GA+AA + GG+GC	41 (71.9)	35 (70.0)	0.83	51 (68.9)	25 (75.8)	0.64	16 (72.7)	59 (70.2)	1.00
GG+GA + GC+CC	69 (94.5)	65 (94.2)		95 (94.1)	39 (95.1)		25 (96.2)	109 (94.0)	
AA + GG	4 (5.5)	4 (5.8)	1.00	6 (5.9)	2 (4.9)	1.00	1 (3.8)	7 (6.0)	1.00
VEGFA -1154G/A + -460C/T									
GG + TT	38 (48.7)	39 (52.7)		53 (51.0)	24 (50.0)		13 (46.4)	64 (52.0)	
GA+AA + CC+CT	40 (51.3)	35 (47.3)	0.63	51 (49.0)	24 (50.0)	1.00	15 (53.6)	59 (48.0)	0.67
GG+GA + CT+TT	87 (95.6)	85 (95.5)		116 (95.1)	56 (96.6)		33 (97.1)	139 (95.2)	
AA + CC	4 (4.4)	4 (4.5)	1.00	6 (4.9)	2 (3.4)	1.00	1 (2.9)	7 (4.8)	1.00
VEGFA 634G/C + -460C/T									
CC + TT	18 (23.1)	18 (22.5)	1.00	24 (22.9)	12 (22.6)	1.00	6 (21.4)	30 (23.3)	1.00
GG+GC + CC+CT	60 (76.9)	62 (77.5)		81 (77.1)	41 (77.4)		22 (78.6)	99 (76.7)	
GC+CC + CT+TT	72 (86.7)	66 (82.5)	0.51	94 (85.5)	44 (83.0)	0.81	27 (96.4)	111 (82.8)	0.08
GG + CC	11 (13.3)	14 (17.5)		16 (14.5)	9 (17.0)		1 (3.6)	23 (17.2)	
VEGFA -2578C/A + KDR -604T/C									
CC + TT	14 (26.4)	9 (17.0)		17 (23.3)	6 (18.2)		7 (33.3)	16 (18.8)	
CA+AA + TC+CC	39 (73.6)	44 (83.0)	0.34	56 (76.7)	27 (81.8)	0.62	14 (66.7)	69 (81.2)	0.23
CC+CA + TT+TC	68 (97.1)	65 (95.6)		90 (97.8)	43 (93.5)		26 (100)	106 (95.5)	
AA + CC	2 (2.9)	3 (4.4)	0.67	2 (2.2)	3 (6.5)	0.33	0 (0)	5 (4.5)	0.58
VEGFA -2578C/A + KDR -271G/A									
CC + AA	12 (22.6)	16 (28.1)		20 (26.7)	8 (22.9)		3 (16.7)	25 (27.5)	
CA+AA + GG+GA	41 (77.4)	41 (71.9)	0.66	55 (73.3)	27 (77.1)	0.81	15 (83.3)	66 (72.5)	0.55
CC+CA + GA+AA	68 (98.6)	71 (95.9)		93 (96.9)	46 (97.9)		23 (95.8)	116 (97.5)	

TT + GG	3 (11.5)	8 (27.6)		9 (25.7)	2 (10.0)		1 (11.1)	10 (21.7)	
CC+CT + GA+AA	23 (88.5)	21 (72.4)	0.18	26 (74.3)	18 (90.0)	0.29	8 (88.9)	36 (78.3)	0.66
CT+TT + GG+GA	53 (98.1)	55 (98.2)		69 (98.6)	39 (97.5)		19 (95.0)	88 (98.9)	
CC + AA	1 (1.9)	1 (1.8)		1 (1.4)	1 (2.5)		1 (5.0)	1 (1.1)	0.33
VEGFA -2489C/T + KDR +1719A/T									
TT + TT	4 (11.8)	9 (20.5)		10 (20.8)	3 (10.0)		0 (0)	13 (20.0)	
CT+TT + AA+AT	30 (88.2)	35 (79.5)	0.37	38 (79.2)	27 (90.0)	0.34	12 (100)	52 (80.0)	0.20
CT+TT + AT+TT	50 (100)	54 (98.2)		66 (98.5)	38 (100)		17 (100)	86 (98.9)	
CC + AA	0 (0)	1 (1.8)		1 (1.5)	0 (0)	1.00	0 (0)	1 (1.1)	1.00
VEGFA -1154G/A + KDR -604T/C									
GG + TT	17 (34.0)	13 (30.2)		20 (32.3)	10 (32.3)		8 (40.0)	22 (30.1)	
GA+AA + TC+CC	33 (66.0)	30 (69.8)	0.82	42 (67.7)	21 (67.7)	1.00	12 (60.0)	51 (69.9)	0.42
GG+GA + TT+TC	69 (100)	67 (100)		94 (100)	42 (100)		26 (100)	109 (100)	
AA + CC	0 (0)	0 (0)	1.00	0 (0)	0 (0)	1.00	0 (0)	0 (0)	1.00
VEGFA -1154G/A + KDR -271G/A									
GG + AA	13 (28.3)	22 (43.1)		24 (36.4)	11 (35.5)		3 (20.0)	32 (39.5)	
GA+AA + GG+GA	33 (71.7)	29 (56.9)	0.14	42 (63.6)	20 (64.5)	1.00	12 (80.0)	49 (60.5)	0.24
GG+GA + GA+AA	67 (98.5)	73 (97.3)		96 (98.0)	44 (97.8)		21 (95.5)	119 (98.3)	
AA + GG	1 (1.5)	2 (2.7)	1.00	2 (2.0)	1 (2.2)	1.00	1 (4.5)	2 (1.7)	0.39
VEGFA -1154G/A + KDR +1192G/A									
GG + GG	42 (80.8)	50 (86.2)		65 (85.5)	27 (79.4)		15 (83.3)	77 (83.7)	
GA+AA + GA+AA	10 (19.2)	8 (13.8)	0.45	11 (14.5)	7 (20.6)	0.41	3 (16.7)	15 (16.3)	1.00

GG+GA + GG+GA	93 (100)	95 (100)		126 (100)	62 (100)		32 (100)	155 (100)
AA + AA	0 (0)	0 (0)	1.00	0 (0)	0 (0)	1.00	0 (0)	0 (0) 1.00
VEGFA -1154G/A + KDR +1719A/T								
GG + TT	41 (73.2)	42 (76.4)		60 (77.9)	23 (67.6)		14 (66.7)	69 (77.5)
GA+AA + AA+AT	15 (26.8)	13 (23.6)	0.82	17 (22.1)	11 (32.4)	0.34	7 (33.3)	20 (22.5) 0.39
GG+GA + AT+TT	91 (100)	94 (100)		123 (100)	62 (100)		31 (100)	153 (100)
AA + AA	0 (0)	0 (0)	1.00	0 (0)	0 (0)	1.00	0 (0)	0 (0) 1.00
VEGFA -634G/C +KDR -604T/C								
CC + TT	4 (6.2)	3 (4.2)		6 (6.5)	1 (2.4)		1 (5.0)	6 (5.2)
GC+GG + TC+CC	60 (93.8)	68 (95.8)	0.70	87 (93.5)	41 (97.6)	0.43	19 (95.0)	109 (94.8) 1.00
CC+GC + TT+TC	49 (89.1)	48 (80.0)		69 (88.5)	28 (75.7)		19 (95.0)	78 (82.1)
GG + CC	6 (10.9)	12 (20.0)	0.20	9 (11.5)	9 (24.3)	0.10	1 (5.0)	17 (17.9) 0.19
VEGFA -634G/C + KDR -271G/A								
CC + AA	2 (3.1)	4 (6.3)		4 (4.7)	2 (4.8)		1 (4.0)	5 (5.0)
GG+GC + GG+GA	62 (96.9)	59 (93.7)	0.44	81 (95.3)	40 (95.2)	1.00	24 (96.0)	96 (95.0) 1.00
GC+CC + GA+AA	52 (86.7)	49 (87.5)		70 (89.7)	31 (81.6)		16 (88.9)	85 (87.6)
GG + GG	8 (13.3)	7 (12.5)	1.00	8 (10.3)	7 (18.4)	0.24	2 (11.1)	12 (12.4) 1.00
VEGFA -634G/C + KDR +1192G/A								
CC + GG	14 (38.9)	16 (42.1)		22 (44.9)	8 (32.0)		6 (42.9)	24 (40.0)
GG+ GC + GA+AA	22 (61.1)	22 (57.9)	0.81	27 (55.1)	17 (68.0)	0.32	8 (57.1)	36 (60.0) 1.00
GC+CC + GG+GA	71 (100)	67 (98.5)		95 (99.0)	43 (100)		26 (100)	112 (99.1)

TT + TT	29 (59.2)	28 (54.9)		39 (60.9)	18 (50.0)		11 (52.4)	46 (59.0)
CC+CT + AA+AT	20 (40.8)	23 (45.1)	0.69	25 (39.1)	18 (50.0)	0.30	10 (47.6)	32 (41.0)
CT+TT + AT+TT	87 (100)	86 (98.9)		112 (100)	61 (98.4)		33 (100)	140 (99.3)
CC + AA	0 (0)	1 (1.1)	1.00	0 (0)	1 (1.6)	0.35	0 (0)	1 (0.7)

KDR -604T/C + -271G/A

TT + AA	0 (0)	0 (0)		0 (0)	0 (0)		0 (0)	0 (0)
TC+CC + GG+GA	51 (100)	52 (100)	1.00	71 (100)	32 (100)	1.00	17 (100)	86 (100)
TT+TC + GA+AA	44 (100)	47 (97.9)		66 (98.5)	25 (100)		14 (100)	77 (98.7)
CC + GG	0	1 (2.1)	1.00	1 (1.5)	0	1.00	0	1 (1.3)

KDR -604T/C + +1192G/A

TT + GG	25 (51.0)	18 (46.2)		28 (50.9)	15 (45.5)		11 (61.1)	31 (44.9)
TC+CC + GA+AA	24 (49.0)	21 (53.8)	0.67	27 (49.1)	18 (54.5)	0.66	7 (38.9)	38 (55.1)
TT+TC + GG+GA	72 (98.6)	71 (98.6)		98 (99.0)	45 (97.8)		27 (96.4)	115 (99.1)
CC + AA	1 (1.4)	1 (1.4)	1.00	1 (1.0)	1 (2.2)	0.53	1 (3.6)	1 (0.9)

KDR -604T/C + +1719A/T

TT + TT	14 (42.4)	12 (30.0)		18 (39.1)	8 (29.6)		7 (46.7)	19 (32.8)
TC+CC + AA+AT	19 (57.6)	28 (70.0)	0.33	28 (60.9)	19 (70.4)	0.45	8 (53.3)	39 (67.2)
TT+TC + AT+TT	70 (98.6)	69 (100)		95 (99.0)	44 (100)		1 (3.7)	0 (0)
CC + AA	1 (1.4)	0 (0)	1.00	1 (1.0)	0 (0)	1.00	26 (96.3)	112 (100)

KDR -271G/A + +1192G/A

AA + GG	13 (44.8)	22 (59.5)		25 (55.6)	10 (47.6)		4 (44.4)	31 (54.4)
GG+GA + GA+AA	16 (55.2)	15 (40.5)	0.32	20 (44.4)	11 (52.4)	0.60	5 (55.6)	26 (45.6) 0.72
GA+AA + GG+GA	72 (100)	77 (98.7)		100 (99.0)	49 (100)		22 (100)	127 (99.2)
GG + AA	0 (0)	1 (1.3)	1.00	1 (1.0)	0 (0)	1.00	0 (0)	1 (0.8) 1.00

KDR -271G/A + +1719A/T

AA + TT	18 (40.0)	21 (43.8)		26 (44.8)	13 (37.1)		5 (31.3)	34 (44.7)
GG+GA + AA+AT	27 (60.0)	27 (56.2)	0.83	32 (55.2)	22 (62.9)	0.52	11 (68.8)	42 (55.3) 0.40
GA+AA + AT+TT	70 (100)	76 (98.7)		97 (99.0)	49 (100)		21 (100)	125 (99.2)
GG+ AA	0 (0)	1 (1.3)	1.00	1 (1.0)	0 (0)	1.00	0 (0)	1 (0.8) 1.00

KDR +1192G/A + 1719A/T

GG + TT	56 (81.2)	58 (79.5)		81 (84.4)	33 (71.7)		22 (75.9)	92 (81.4)
GA+AA + AA+AT	13 (18.8)	15 (20.5)	0.83	15 (15.6)	13 (28.3)	0.11	7 (24.1)	21 (18.6) 0.60
GG+GA + AT+TT	97 (100)	100 (99.0)		129 (99.2)	68 (100)		33 (100)	163 (99.4)
AA + AA	0 (0)	1 (1.0)		1 (0.8)	0 (0)		0 (0)	1 (0.6) 1.00

No, number; %: percent. *: histological grade classified according to World Health Organization criteria, where grade III included anaplastic astrocytoma and oligoastrocytoma and grade IV included glioblastoma multiforme; the histological grade was not identified in one patient.