Primer	VWF Fragment	SEQUENCING / AMPLIFICATION	Description
1	cDNA-1	AACTTGGAGCTATTGCAGGC	5'UTR (1-F)
2	cDNA-1	GGCAGTAGCAGCCTTCAAGA	Amplifies fragment with EcoRI (1-R)
3	cDNA-1	aaaaagcggccgcAACTTGGAGCTATTGCAGGCAGAGGAATGGGC	5'UTR with NotI (1-Fnest)
4	cDNA-1	TTCATCCGGGAGGGAGAG	Amplifies fragment with EcoRI (1-Rnest)
5	cDNA-2	CAGGTGTACCTGCAGTGTGG	Amplifies fragment with EcoRI (2-F)
6	cDNA-2	TAGCGGACCAAATTCCTAGC	Amplifies fragment with NheI (2-R)
7	cDNA-2	CTGGCTAGCAGTCAGGAGCA	Amplifies fragment with NheI (2-Rnest)
8	cDNA-3	TGGCCTCTACCAGTGAGGTT	Amplifies fragment with NheI (3-F)
9	cDNA-3	GCAAAATTGACCGCCCTGAAGC	Amplifies fragment with NheI (3-Fnest)
10	cDNA-3	ATCCACTGCCATCTCCATGT	Amplifies fragment with SphI (3-R)
11	cDNA-3	GAGACGCCAGCATGCTTAAT	Amplifies fragment with SphI (3-Rnest)
12	cDNA-4	AAGCCTGCATGAAGTCCATT	Amplifies fragment with SphI (4-F)
13	cDNA-4	ATTAAGCATGCTGGCGTCTC	Amplifies fragment with SphI (4-Fnest)
14	cDNA-4	tctagaCTTGCTGCAGTTCCGGGGAGAACAC	VWF 3' with XbaI (4-R)
15	5'RACE	CCTGTCACGGGGCTTTTCTGTG	Outer 5' RACE (Exon 2-R)
16	5'RACE	TCTTGCCCATACAAACAGGGGCTGTA	Inner 5' RACE (Exon 1-R)
17	3'RACE	ACTGCACCAATGGCTCTGTC	Exon 52-F
18	Exon 11	TTGTCATGGGGAAAAGGGTA	PolyPhred, intron 10 (F)
19	Exon 11	CCACTTAGCCCTCCATTTCA	PolyPhred, intron 11 (R)
20	Exon 11	CCTATGGATCTGCAGCAATG	PolyPhred, exon 10 (F)
21	Exon 11	TGCTCCCCCGTGCTTCAGTTT	PolyPhred, exon 12 (R)
22	+774	GGGAGCAATGCCAGCTACT	FE, exon 7 (F)
23	+774	GGCACTGTGGTCAGTCCAG	FE, exon 7 (R)
24	+774	CAGACAGCATCAGGGTCATC	FE, exon 12 (R)
25	+774	CATGCGCACTCTGGCCCCGT	FE, exon 7 SNP, 5'Cy-5 labeled (R)
26	+7970	TATCGGGAAGAGAAGAGCCA	FE, exon 48 (F)
27	+7970	CTTGCTTCAGGGTCATGATC	FE, exon 48 (R)
28	+7970	AGTTCCGGGGAGAACACCT	FE, exon 52 (R)
29	+7970	TATTCAGCTAAGAGGAGGAC	FE, exon 48 SNP, 5'Cy-5 labeled (F)
30	"A" Upper Strand	CTAGAGCGGCCGCTGCTGAATTCTGCTGCTAGCTGCTGCATGCTGCTTCTAGATGATAG	
31	"A" Lower Strand	GCCCTATCATCTAGAAGCAGCATGCAGCAGCTAGCAGCAGAATTCAGCAGCGGCCGCT	
32	"+" Upper Strand	CTAGAGAACAAAAACTCATCTCAGAAGAGGATCTGTGAG	
33	"+" Lower Strand	GGCCTCACAGATCCTCTTCTGAGATGAGTTTTTGTTCT	
34	AA2657	TAAGAGGAGGACgGATCATGACCCT	A/J Strand
35	AA122	GACTCTACCTAGAACtTGAGGCTGGGTACTATAAG	A/J Upper Strand
36	AA122	CTTATAGTACCCAGCCTCAaGTTCTAGGTAGAGTC	A/J Lower Strand
37	AA122	GACTCTACCTAGAACgCGAGGCTGGGTACTATAAG	CASA/Rk Upper Strand
38	AA122	CTTATAGTACCCAGCCTCGcGTTCTAGGTAGAGTC	CASA/Rk Lower Strand