

Table SI. Predicted binding sites for ZNF667 in the promoters of 10 angiogenesis-related genes using JASPAR database.

Gene symbol	Number of predicted binding site	Score of predicted binding site	Relative score of predicted binding site	Location of predicted binding site (bp)
<i>VEGFA</i>	3	9.12, 8.91, 8.32	0.81, 0.80, 0.80	-491~-485, -614~-608, -1123~-1117
<i>VASH1</i>	2	9.363, 12.234	0.84, 0.82	-226~-222, -154~-158
<i>SVBP</i>	2	6.84, 7.21	0.80, 0.81	-334~-328, -927~-921
<i>PFKFB3</i>	2	10.0, 8.48	0.85, 0.81	-214~-208, -689~-683
<i>FGF2</i>	1	7.31	0.81	-1799~-1788
<i>KDR</i>	2	7.99, 7.33	0.82, 0.81	-450~-439, -1437~-1426
<i>FLT1</i>	3	5.46, 5.15, 5.51	0.77, 0.77, 0.78	-60~-49, -220~-209, -1190~-1179
<i>ANGPT1</i>	2	7.79, 7.75	0.82, 0.81	-903~-892, -864~-853
<i>ANGPT12</i>	3	7.97, 7.57, 7.32	0.82, 0.81, 0.81	-849~-838, -106~-95, -1120~-1109
<i>TEK</i>	3	7.97, 7.97, 7.89	0.82, 0.82, 0.82	-1503~-1492, -245~-234, -117~-106