

Supplementary File1

Possible Prognostic Impact of PKC α Genetic Variants in Prostate Cancer.

Amna Hafeez¹, Maria Shabbir¹, Khushbukhat Khan¹, Janeen H. Trembley^{2,3,4}, Yasmin Badshah¹, Sameen Zafar¹, Kanza Shahid¹, Hania Shah¹, Naeem Mahmood Ashraf⁵, Arslan Hamid⁶, Tayyaba Afsar⁷, Ali Almajwal⁷, Suhail Razak⁷

Table 1: PRKCI primer sequences for Tetra-ARMS PCR

PRKCI primers for Tetra-ARM's PCR		
G34W (rs1199520604)	Forward inner	5'-GTGAAAGCCTACTACCACG-3'
	Reverse inner	5'-TCCCAGGACACTCATCA-3'
	Outer forward	5'-AGGTGGGCAGGTAGGT-3'
	Outer reverse	5'-CACCCCTATCACTTCGTC-3'.
F66Y (rs1197750201)	Forward inner	5' -GTGTTCTTTTGACAACGAACAGCTGT-3'
	Reverse inner	5'-CTTCCTCATCTATCCATTTTCATGGAGA-3'
	Forward outer	5'- AAGTTCATCAAATTGTCAAGCATTCAGT-3'
	Reverse outer	5'-AAGAAAAAACAAGAATGGCATCAAATTT-3'

Table 2: HADDOCK results for the docking interactions

Interaction	Wild-Type PKCI- Par-6	PKCI-F66Y-Par-6	PKCI-G34W-Par-6
HADDOCK score	56.7 +/- 13.0	90.5 +/- 16.3	15.6 +/- 26.0
RMSD from the overall lowest-energy structure	21.7 +/- 0.7	22.0 +/- 0.3	0.8 +/- 0.5
Van der Waals energy	-81.7 +/- 7.4	-96.3 +/- 7.0	-79.6 +/- 8.3
Electrostatic energy	-167.6 +/- 46.1	-184.8 +/- 7.3	-182.0 +/- 34.6
Desolvation energy	-14.5 +/- 6.3	-30.0 +/- 2.8	-36.4 +/- 3.5
Buried Surface Area	2803.2 +/- 269.5	3635.1 +/- 100.9	2970.3 +/- 38.1
Z-Score	-2.0	-1.7	-1.8