

Figure S1. Amino acid sequence alignment comparison of human SLURP1 and a few selected proteins. An amino acid sequence alignment was conducted for human SLURP1, human SLURP2, Human Lynx1, Human Lypd6, Mouse SLURP1, Neurotoxin 1, Erabutoxin, and α - Bungarotoxin. The degree of sequence conservation was also demonstrated.

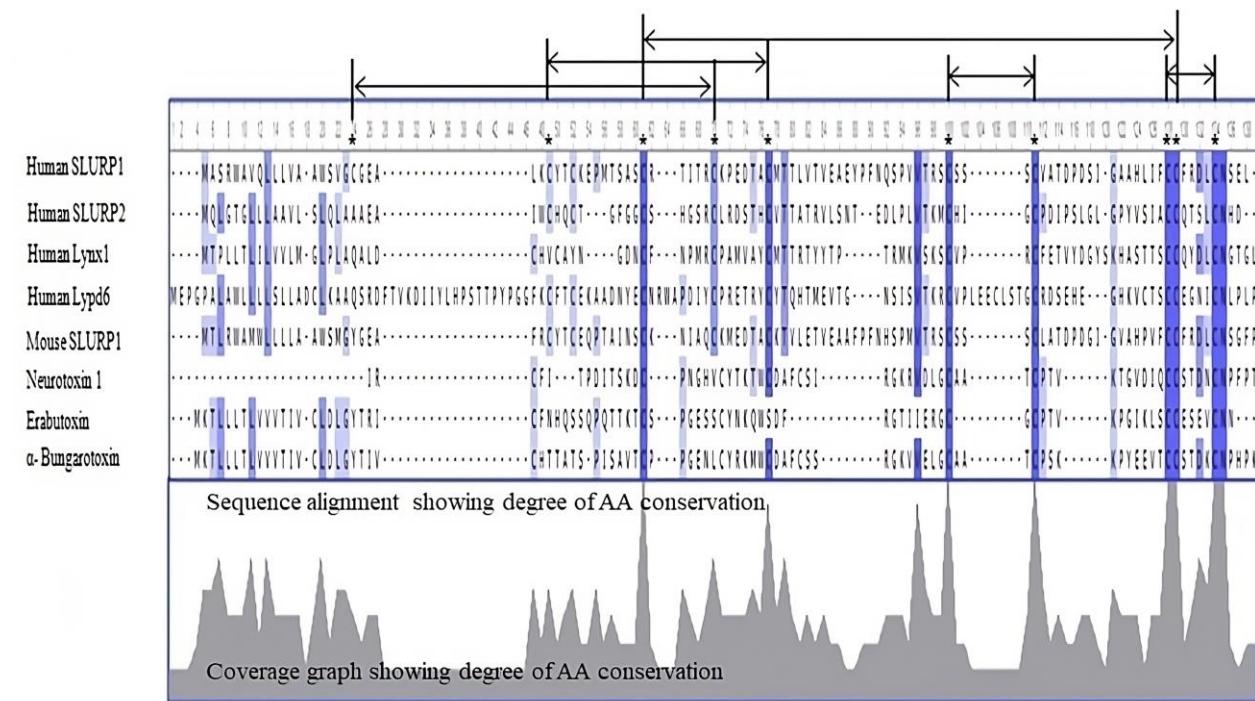


Figure S2. The original uncropped blot image showing all the bands with all molecular weight. Lanes 1 and 6 represent the molecular marker (size range from 10 to 180 kDa). Lanes 2 and 5 denote the purified rSLURP1 protein expressed by *E. coli* (DE3) as a positive control. Lane 3 indicates Vector control (VC) (ST JOL1800 with an empty vector), and lane 4 represents the therapeutic strain, JOL2238.

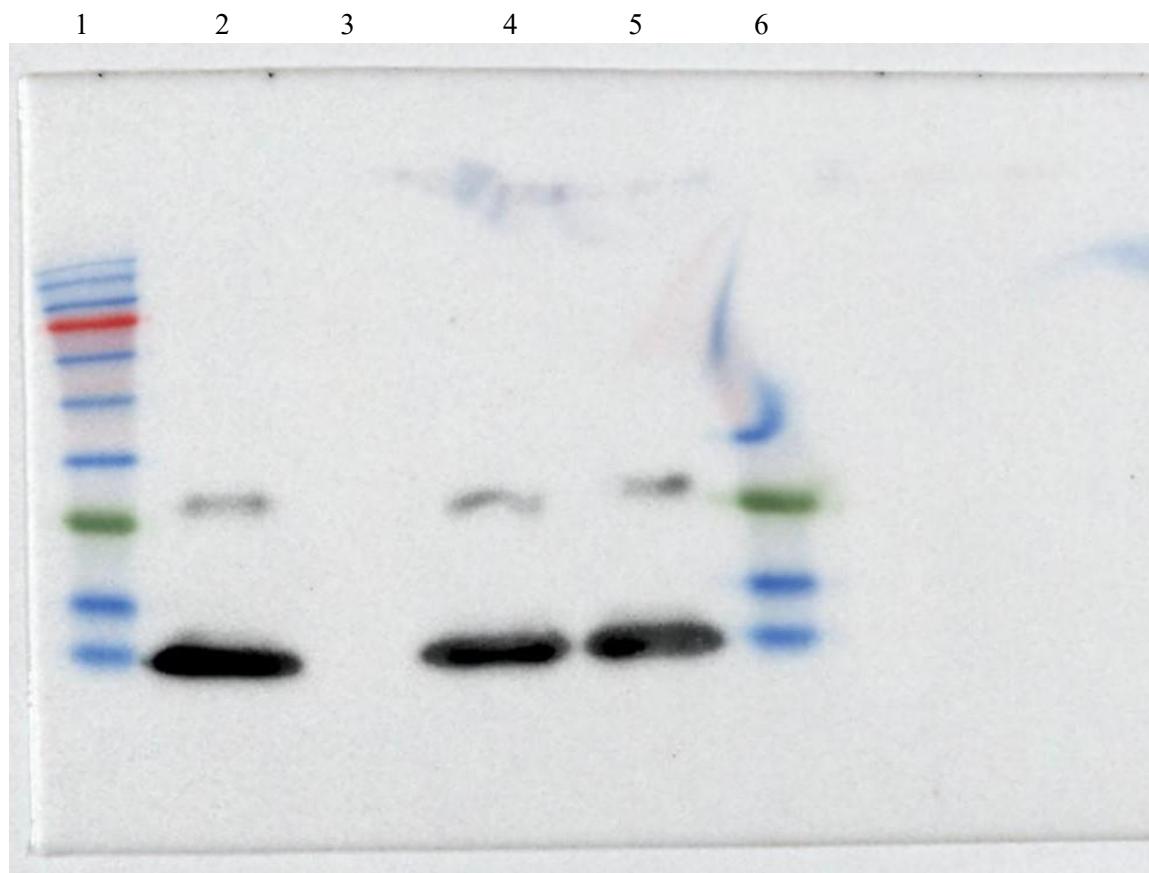


Table S1. Primer sequences used in qRT-PCR.

Gene		Forward	Reverse
Cell cycle-related	Cyclin E	GTCCTGGCTGAATGTATAACATGC	CCCTATTGTTAGACAAACATGGC
	Cyclin B1	GGCTTCTCTGATGTAATTCTTGC	GTATTTGGTCTGACTGCTTGC
	Cyclin B2	CCTCCCTTTCAAGTCCGC	CTCCTGTGTCATATTCTCAAATC
	Cyclin D1	CATCTACACCGACAACCTCCATC	TCTGGCATTGGAGAGGAAG
	Cyclin D2	GCTCTGTGCGCTACCGACTT	CACGCTTCCAGTTGCAATCA
	Cyclin D3	TGATTGCGCACGACTTCCT	CAAAGCCTGCCGGTCACT
	Cyclin A2	CTGCATTTGGCTGTGAAC TAC	ACAAACTCTGCTACTTCTGGG
(Human)	Endogenous receptor	CTGCATTTGGCTGTGAAC TAC	CACAAACTGAGACTAAGGCAG
	SLURP1	TTCTGAGCACGGAGCAATG	TTGCAGGTGTAGCACTTGAG
	NF- κ B	CTCCACAAGGCAGCAAATAGA	ACTGGTCAGAGACTCGGTAAA
	TNF- α	CTTGTTCCTCAGCCTCTCTC	TCAGCTTGAGGGTTGCTAC
	IFN- γ	GTGGAGACCATCAAGGAAGAC	ACCTCGAAACAGCATCTGAC
	IL-1 β	CTGCGTGTGAAAGATGATAAG	CCACATTTCAGCACAGGACTC
	RelA	GGTGGGCACTGGAGTTATT	GAGGATGGATGGTCCAAC TTAC
(Mice)	STAT1	CTGGAGGAGTTGGAACAGAAA	CTGAATGAGCTGCTGGAAGA
	TBK1	AGTACCTGCTGCTCTATCA	GTGGACGGTCTCGTTGTAATC
	IRF3	GTCTTAAGGAGCTGTTAGAGATGG	TGGTCAGAGGTAAAGGGAGATAG
	GAPDH	AACAGCAACTCCC ACTCTTC	CCTGTTGCTGTAGCCGTATT
	NF- κ B	TGGGACCAGCAAAGGTTATT	GATCCCATCCTCACAGTGT TTT
	VEGF	GCGAAGCTACTGCCGTCC	TCTGCATTGTGATGTTGCTC
	Bax	AGGATGCGTCCACCAAGAAGCT	TCCGTGTCCACGT CAGCAATCA
	Bcl2	CCTGTGGATGACTGAGTACCTG	AGCCAGGAGAAATCAAACAGAGG