## **Supplementary Data**

MicroRNA 138 upregulation is associated with decreasing levels of *CCND1* gene expression and promoting cell death in human prostate cancer cell lines

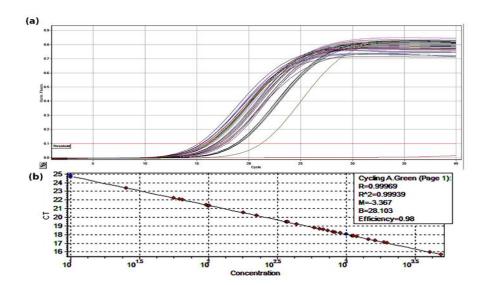
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**Figure S1**. (a) Difference plot of the same sample. Serial dilutions of 10<sup>1</sup>, 10<sup>2</sup> and 10<sup>3</sup> of a pooled cDNA sample were tested with of GAPDH primers. (b) qRT-PCR efficiency of GAPDH gene (data of other set of primers is not shown).

**Table S1.** The expression of miR-138 evidently diminished in PCa clinical samples in comparison U6 and U47 as normalizers.

| Gene       | Туре | Reaction Efficiency | Expression | Std. Error    | 95% C.I.      | P(H1) | Result |
|------------|------|---------------------|------------|---------------|---------------|-------|--------|
| U47        | REF  | 0.7384              | 1.981      |               |               |       |        |
| U6         | REF  | 0.6575              | 0.505      |               |               |       |        |
| mir 138-5p | TRG  | 0.7397              | 0.266      | 0.051 - 1.402 | 0.012 - 5.326 | 0.004 | DOWN   |

**Table S2.** Results of quantitative RT-PCR on all candidate genes in PC3 cell lines. Substantial overexpression of *CCND1* in comparison with *GAPDH* as a housekeeping gene in PC3 cells. The expression profile of other candidate genes, including *ABL*, *CCND3*, *VIM*, *TWIST1*, *HIF1A*, and *TERT* genes exhibited no considerable change in comparison with *GAPDH* in the PC3 cell line.

| Gene  | Туре | Reaction Efficiency | Expression | Std. Error     | 95% C.I.       | P(H1) Result |
|-------|------|---------------------|------------|----------------|----------------|--------------|
| GAPDH | REF  | 0.98                | 1.000      |                |                |              |
| ABL   | TRG  | 0.99                | 0.952      | 0.090 - 9.223  | 0.012 - 25.820 | 0.931        |
| CCND1 | TRG  | 1.0                 | 3.210      | 0.308 - 23.123 | 0.099 - 69.005 | 0.028 UP     |
| CCND3 | TRG  | 1.0                 | 0.408      | 0.032 - 5.532  | 0.005 - 19.999 | 0.178        |
| HIF1a | TRG  | 0.99                | 2.233      | 0.281 - 32.173 | 0.026 - 82.124 | 0.185        |
| VIM   | TRG  | 0.8                 | 0.674      | 0.113 - 3.477  | 0.023 - 11.487 | 0.386        |

| Gene  | Туре | Reaction Efficiency | Expression | Std. Error     | 95% C.I.          | P(H1) | Result |
|-------|------|---------------------|------------|----------------|-------------------|-------|--------|
| GAPDH | REF  | 0.95                | 1.000      |                |                   |       |        |
| Twst1 | TRG  | 0.9                 | 0.623      | 0.022 - 16.699 | 0.002 - 423.948   | 0.384 |        |
| TERT  | TRG  | 0.95                | 1.021      | 0.009 - 54.980 | 0.000 - 3,515.604 | 0.976 |        |

**Table S3.** Results of quantitative RT-PCR on *CCND1* expression in comparison with *GAPDH* following transfection with miR-138-5p-mimic in (a) PC3 and (b) DU145 cells.

(b)

| Gene  | Туре | Reaction Efficiency | Expression | Std. Error    | 95% C.I.      | P(H1) | Result |
|-------|------|---------------------|------------|---------------|---------------|-------|--------|
| GAPDH | REF  | 1.0                 | 1.000      |               |               |       |        |
| CCND1 | TRG  | 1.0                 | 0.530      | 0.260 - 1.149 | 0.124 - 2.037 | 0.090 |        |