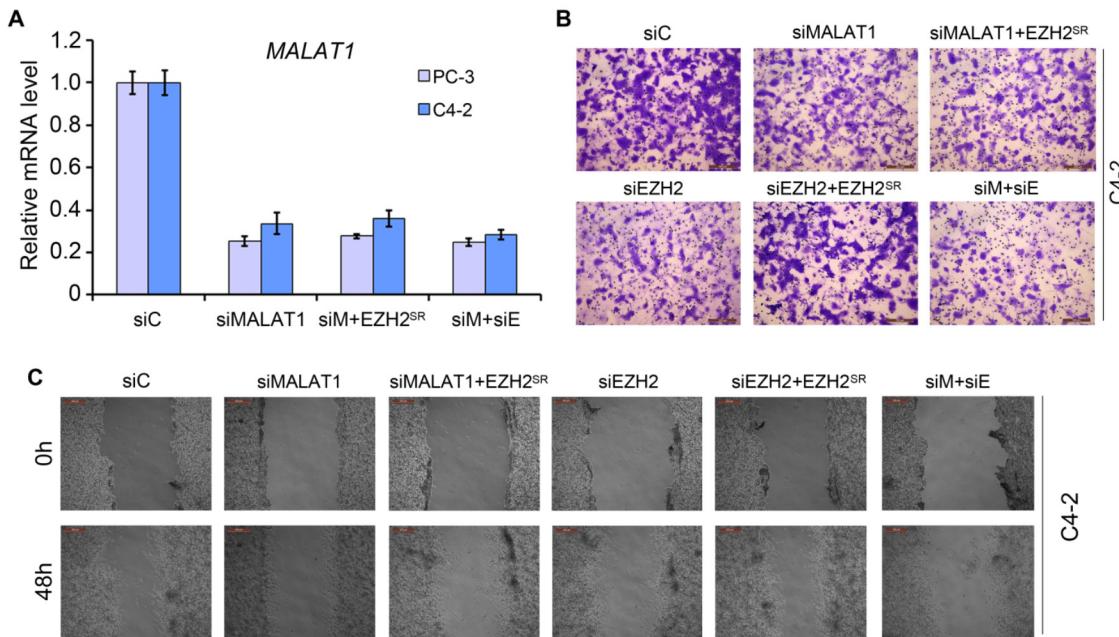


## SUPPLEMENTARY FIGURE AND TABLE



**Supplementary Figure S1: MALAT1 facilitates EZH2-mediated PCa cell migration and invasion.** PC-3 and C4–2 cells were transfected with siRNAs and/or siRNA-resistant EZH2 expression vector (Myc-EZH2<sup>SR</sup>) as indicated in Figure 5A. **A.** At 48 h after transfection, RT-qPCR analysis of *MALAT1* expression in PC-3 and C4–2 cells transfected with control siRNA (siC) and *MALAT1*-specific siRNA (siMALAT1 or siM). Data are means ± S.D. from experiments with three replicates. **B.** C4–2 cells were transfected as indicated in Figure 5A. At 48 h after transfection, cells were used for matrigel invasion assays. Representative images of invasion assay are shown. **C.** At 24 h after transfection, artificial wounds were created on C4–2 cells grown in confluence. Images were taken at 0, 48 h after wound. Scale bar, 200 μm.

Supplementary Table S1: Information of the primers, sequences of siRNAs/shRNAs used

## MALAT1 Cloning Primers

|      |                                     |
|------|-------------------------------------|
| M1-F | CCCAAGCTTGGGTAAAGGACTGGGGCCCCGCAACT |
| M1-R | TATGAAGACTTAGAACAGAGTACCGCTCGAGCGG  |
| M2-F | CCCAAGCTTGGGCATGAGGAAGGAAAAGATA     |
| M2-R | ACCACTCGCTTCCTGTCCGCTCGAGCGG        |
| M3-F | CCCAAGCTTGGTGGTAAAAATCCGTGAGGTC     |
| M3-R | CACTACCATAATCAAACAACCCGCTCGAGCGG    |
| M4-F | CCCAAGCTTGGGTGTTCTCTTTGGAATT        |
| M4-R | TGGTCCATTAAAGAGTGTCCCGCTCGAGCGG     |
| M5-F | CCCAAGCTTGGGATCAGGATTGAGCGGAAG      |
| M5-R | TTGTTGTCTCTGCCACACCGCTCGAGCGG       |
| M6-F | CCCAAGCTTGGGAGCGCTATTATCCTAAGGTC    |
| M6-R | TTTAGAGCTTCTCCATTCCGCTCGAGCGG       |

(Continued)

**MALAT1 Cloning Primers**

| RT-qPCR primers          |                         |
|--------------------------|-------------------------|
| DAB2IP-F                 | ACACGCCATGGAGCCCCGACT   |
| DAB2IP-R                 | GAAGCCCGTGACCCGGAACG    |
| BARCHURY-F               | AGGTGGGGAAGTTCCCTTCT    |
| BARCHURY-R               | GCAAATGAGGTCCCTTGGT     |
| M1-RT-F                  | TTGAGGCCTTCCAAGAGT      |
| TMEM48-RT-F              | AGGTCGCGGGACATACTGT     |
| TMEM48-RT-R              | TGCAGATGGTAGAAATAGCACT  |
| CKS2-RT-F                | TTCGACGAACACTACGAGTACC  |
| CKS2-RT-R                | GGACACCAAGTCTCCTCCAC    |
| KIAA0101-RT-F            | ATGGTGCAGACTAAAGCAGAC   |
| KIAA0101-RT-R            | CCTCGATGAAACTGATGTCGAAT |
| M1-RT-R                  | CGGTTGAGAAGTGGCAAAAT    |
| M2-RT-F                  | GACGGAGGTTGAGATGAAGC    |
| M2-RT-R                  | ATTCGGGGCTCTGTAGTCCT    |
| M3-RT-F                  | CCCACCCCTTAATCAGACT     |
| M3-RT-R                  | CAACAGCACAGCGGTACACT    |
| M4-RT-F                  | GTGTGCCAATGTTCGTTG      |
| M4-RT-R                  | AGGAGAAAGTGCATGGTTG     |
| M5-RT-F                  | AAAGCAAGGTCTCCCCACAAG   |
| M5-RT-R                  | GGTCTGTGCTAGATCAAAGGCA  |
| M6-RT-F                  | CTCCCCACAAGCAACTTCTC    |
| M6-RT-R                  | TTCAACCCACCAAGACCTC     |
| FOXO1-RT-F               | CGTGCTTACAGCCTCTA       |
| FOXO1-RT-R               | ACCTCCATCGTGACAAAA      |
| ChIP-qPCR primers        |                         |
| DAB2IP-F                 | CCTGCTTCTGTTCCCTCTCCTG  |
| DAB2IP-R                 | TTGAACCACCTCCTCCCTCTC   |
| BARCHURY-F               | AGGTGGGGAAGTTCCCTTCT    |
| BARCHURY-R               | GCAAATGAGGTCCCTTGGT     |
| siRNA sequences          |                         |
| non-specific siRNA/shRNA | UAGCGACUAAACACAUCAA     |
| siEZH2 (SMART pool)      | CAAAGAAUCUAGCAUCAUA     |
|                          | GAGGACGGCUUCCCAUUA      |
|                          | GCUGAAGCCUAAUGUUUA      |
|                          | GAAUGGAAACAGCGAAGGA     |
| MALAT1-siRNA (siM1)      | GACCUUGAAAUCCAUGACGUU   |
| MALAT1-siRNA (siM2)      | GAUCCAUAAUCGGUUCAAGGUU  |