

1 **Supplementary Data**

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3 **Supplementary Figure 1. Representative ultra-performance liquid chromatography-**
4 **quadrupole-time-of-flight mass spectrometry (UPLC-Q-TPF MS) profiles of *Morus* root**
5 **cultivars:** 1, mulberrofuran G; 2, mulberrofuran I; 3, kuwanon G; 4, kuwanon D/F/T; 5,
6 kuwanon H; 6, kuwanon D/F/T; 7, luteolin-methyl ester-glycoside fragment; 8, kuwanon A/B;
7 9, kuwanon D/F/T, and 10, morusin analyzed by ESI-positive mode.

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9 **Supplementary Figure 2. Partial least-squares discriminant analysis (PLS-DA) scores,**
10 **their quality parameters, and heat map for ESI-positive mode (A), (B) and (C) from**
11 ***Morus* roots with different cultivars, respectively.** The quality of the PLS-DA scores plots
12 was evaluated by R2X, R2Y, Q2, and p-values (A) and validated by permutation tests (B). The
13 heat map was drawn by R with ggplot2 and the green-red color represents the z-score
14 transformed raw data of *Morus* root metabolites with significant difference among sample
15 groups. Red and green colors indicate a decrease and an increase of metabolite level,
16 respectively.

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18 **Supplementary Figure 3. Box plot graphs of *Morus* roots with different cultivars for ESI-**
19 **positive mode.** Box plot graphs show minimum, first quartile, median, third quartile, maximum
20 and outliers.

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22 **Supplementary Figure 4. Effect of *Morus* roots with different cultivars on the PSA**
23 **expression in LNCaP.** LNCaP cells were treated with MR cultivars *Igsu* (1, 5, or 10 $\mu\text{g/ml}$)
24 with or without DHT (10 nM), and the mRNA expression of VEGF and MMP-2 was detected
25 by RT-PCR. Data are representative of three independent experiments. Data are expressed as

26 the means±S.E.M. ** $p < 0.01$ compared with the only DHT treated group.

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28 **Supplementary Figure 5. Effect of *Morus* roots according to different cultivars (*Simheung*,**
29 ***Daesim*, *Cheong-il*, *Sangchon*, *Daeseong*, *Suhong*, *Suwon*, and *Igsu*) on cell viability in**
30 **LNCaP cells. Cell viability was measured by the MTT assay.** LNCaP cells were treated with
31 MR cultivars (10 µg/ml) for 24 h prior to the MTT assay.

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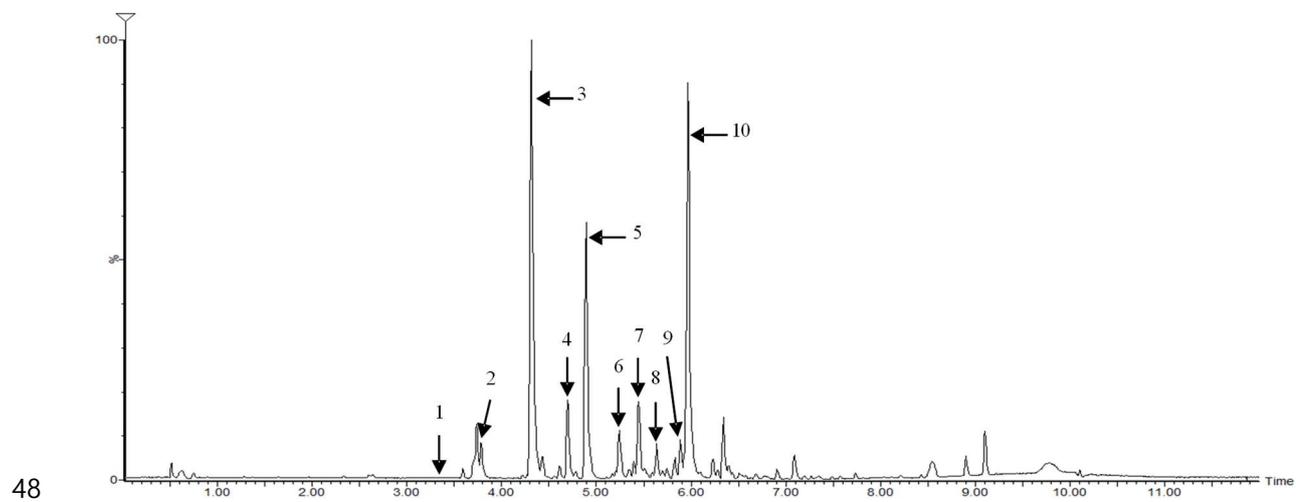
33 **Supplementary Figure 6. Western blot analysis of PSA in LNCaP Cells.** LNCaP cells were
34 incubated in medium containing DHT (10 nM) or *Igsu* (1, 5 or 10 µg/ml) for 24 h, and then
35 cell lysates (30 µg) were assayed for expression level of PSA by Western blotting. Data are
36 representative of three independent experiments. Data are expressed as the means±S.E.M. ##
37 $p < 0.01$ compared with the DHT non treated group; ** $p < 0.01$ compared with the only DHT
38 treated group.

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40 **Supplementary Figure 7. RT-PCR analysis of mRNA expression VEGF and MMP-2 in**
41 **LNCaP cells.** LNCaP cells were treated with *Igsu* (1, 5, or 10 µg/ml) with or without DHT (10
42 nM), and the mRNA expression of VEGF and MMP-2 was detected by RT-PCR. Data are
43 representative of three independent experiments. Data are expressed as the means±S.E.M. ##
44 $p < 0.01$ compared with the DHT non treated group; ** $p < 0.01$ compared with the only DHT
45 treated group.

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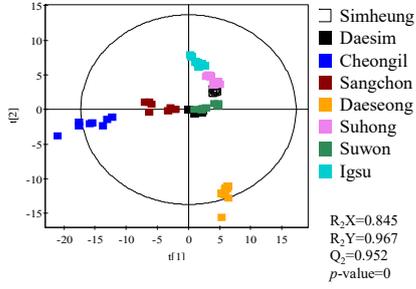


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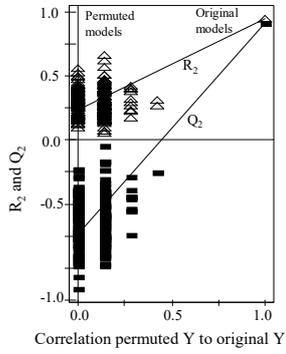
49 **Supplementary Figure 1. Choi et al**

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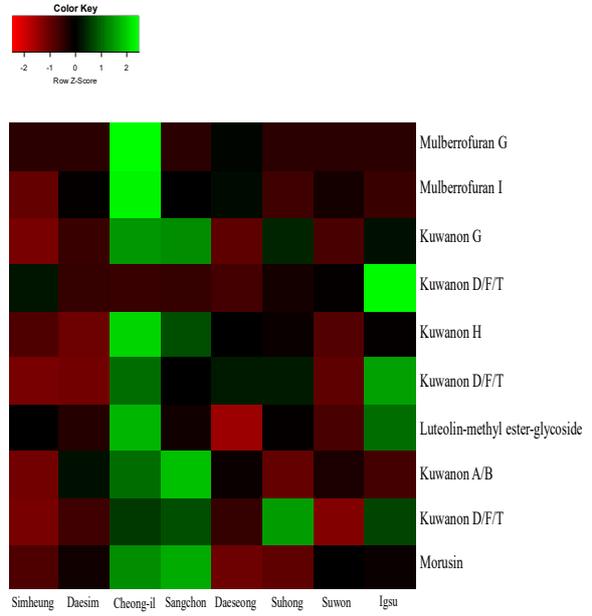
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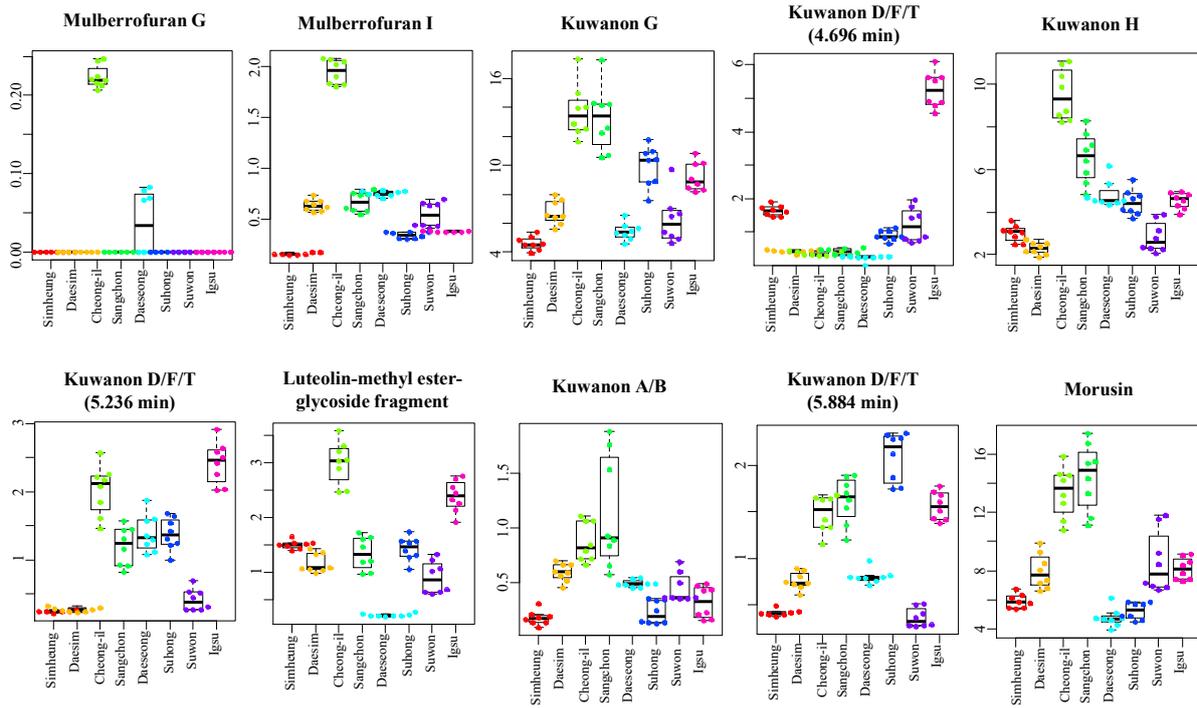


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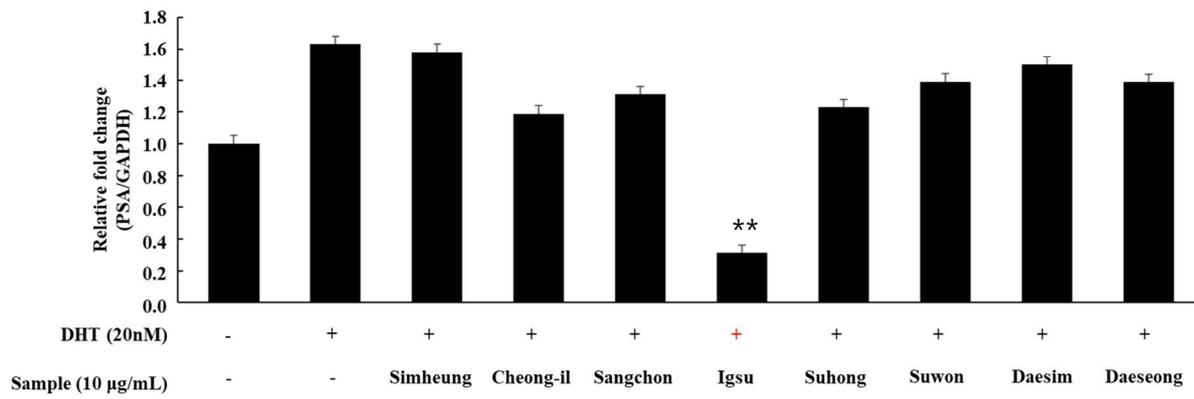
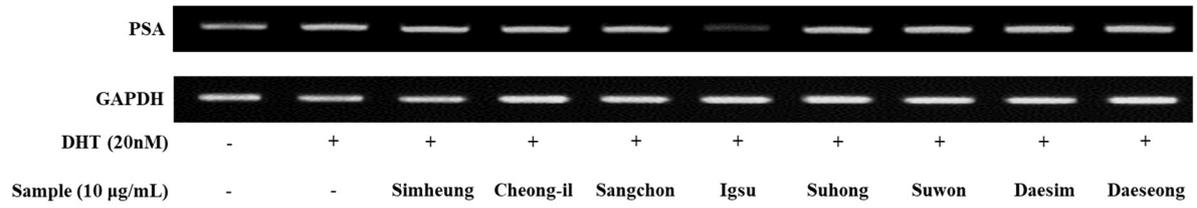


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56 **Supplementary Figure 3. Choi et al**

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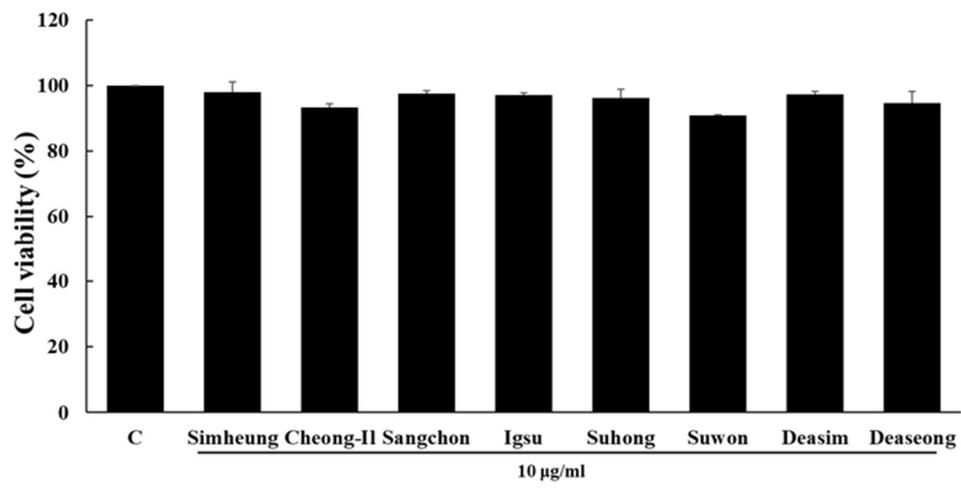


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60 **Supplementary Figure 4. Choi et al**

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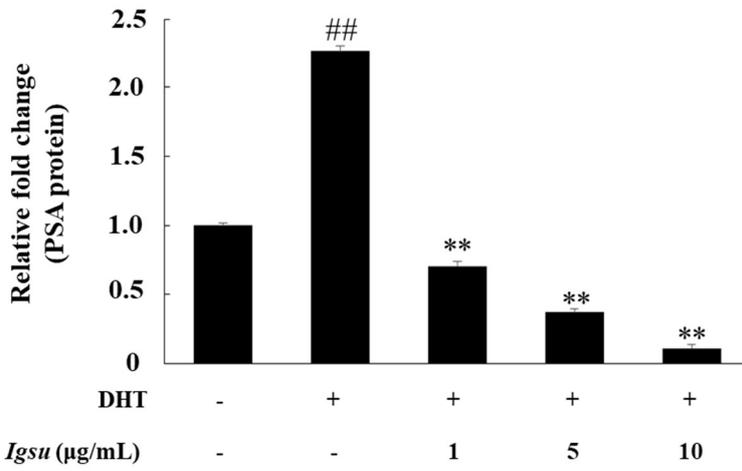
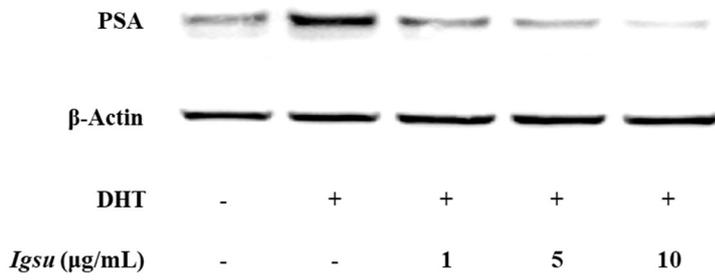
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64 **Supplementary Figure 5. Choi et al**

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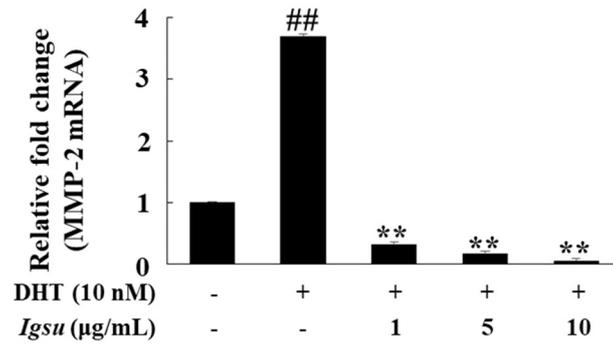
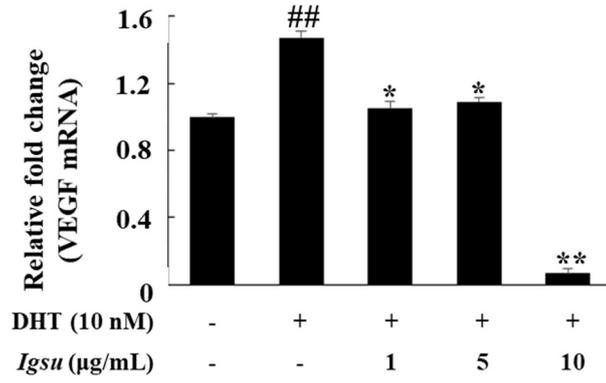
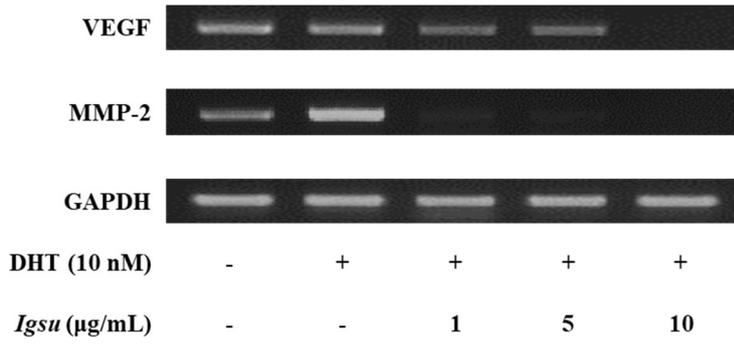


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69 **Supplementary Figure 6. Choi et al**

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73 **Supplementary Figure 7. Choi et al**

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