



Figure S5. Proteome profiles of control seedlings (A) and seedlings treated with 100 μ M AgNP-PVP (B), 100 μ M AgNP-PVP and 500 μ M cysteine (C), 100 μ M AgNP-CTAB (D), 100 μ M AgNP-CTAB and 500 μ M cysteine (E), 100 μ M AgNO₃ (F) and 100 μ M AgNO₃ and 500 μ M cysteine (G) obtained after 2-DE analysis. Differently abundant proteins (at least 1.5-fold compared to the control) are indicated by circles; upregulated proteins are indicated by black line, while down-regulated proteins are indicated with broken-lined circles. The numbers correspond to the numbers listed in Table 2. M, molecular weight markers