

Table 2. Changes of gene expression in *hos9-1*, genes with lower expression in *hos9-1* by 2-fold

| AGI ID | Gene Title | <i>hos9-1</i> vs_C24_0C-24H_Signal Log Ratio | C24-0C-24H_Signal | C24-0C-24H_Detection | <i>hos9-1</i> _0C-24H_Signal | <i>hos9-1</i> _0C-24H_Detection |
|-----------|--|--|-------------------|----------------------|------------------------------|---------------------------------|
| At1g75830 | plant defensin protein, putative (PDF1.1) | -2.3 | 120 | P | 29.8 | A |
| At3g28220 | expressed protein | -2.3 | 1423.8 | P | 320.5 | P |
| At5g44120 | 12S seed storage protein (CRA1) | -1.9 | 436.1 | P | 122.1 | P |
| At3g55970 | oxidoreductase, 2OG-Fe(II) oxygenase family | -1.8 | 290.7 | P | 92.6 | A |
| At1g60590 | polygalacturonase, putative | -1.7 | 523.1 | P | 146.5 | P |
| At5g44420 | plant defensin protein, putative (PDF1.2a) | -1.7 | 5058.4 | P | 1292.8 | P |
| At5g40420 | oleosin | -1.6 | 195.9 | P | 57.8 | A |
| At2g41260 | glycine-rich, late embryogenesis abundant M17 protein | -1.5 | 102.5 | P | 39.3 | A |
| At2g21210 | auxin-induced (indole-3-acetic acid induced) protein, putative | -1.5 | 475.9 | P | 173.6 | A |
| At4g34760 | auxin-induced (indole-3-acetic acid induced) protein family | -1.5 | 201.6 | P | 75.1 | A |
| At5g57180 | CIA2 (CIA2) | -1.5 | 900.1 | P | 297.2 | P |
| At1g19670 | coronatine-induced protein 1 (COR1) | -1.3 | 722 | P | 318.9 | P |
| At4g38860 | auxin-induced (indole-3-acetic acid induced) protein, putative | -1.3 | 228.5 | P | 82.8 | P |
| At5g03350 | expressed protein | -1.3 | 229.1 | P | 77 | A |
| At2g26020 | plant defensin protein, putative (PDF1.2b) | -1.3 | 14437 | P | 5323.6 | P |
| At2g39030 | GCN5-related N-acetyltransferase (GNAT) family | -1.2 | 1850.3 | P | 921.2 | P |
| At2g28200 | zinc-finger protein -related | -1.2 | 274.9 | P | 107 | P |
| At1g78995 | expressed protein | -1.2 | 452.9 | P | 164.7 | A |
| At1g19960 | expressed protein | -1.2 | 883 | P | 362.4 | P |
| At4g26150 | GATA zinc finger protein | -1.2 | 1173.2 | P | 504.4 | P |
| At1g70700 | expressed protein | -1.1 | 1102.4 | P | 676.9 | P |
| At4g25990 | CIL protein | -1.1 | 255.8 | P | 114.3 | P |
| At4g28520 | 12S seed storage protein (cruciferin), putative | -1.1 | 445.1 | P | 132.5 | P |
| At3g49620 | oxidoreductase (din11), putative | -1.1 | 1752.3 | P | 854 | P |
| At5g63530 | copper chaperone (CCH)-related | -1.1 | 355.3 | P | 151.9 | P |
| At2g38240 | oxidoreductase, 2OG-Fe(II) oxygenase family | -1 | 602.8 | P | 293.8 | M |
| At2g28410 | predicted GPI-anchored protein | -1 | 109.4 | P | 83.9 | A |
| At1g10640 | polygalacturonase, putative | -1 | 183.8 | P | 98.3 | A |
| At1g06160 | ethylene response factor, putative | -1 | 141.2 | P | 78.8 | A |
| At1g52400 | glycosyl hydrolase family 1, beta-glucosidase (BG1) | -1 | 3821.8 | P | 1705.9 | P |
| At3g03190 | glutathione transferase, putative | -1 | 239 | P | 139.3 | A |
| At3g15095 | hypothetical protein | -1 | 341.2 | P | 158 | P |
| At1g51780 | auxin conjugate hydrolase/IAA-amino acid hydrolase (ILL5) | -1 | 315.6 | P | 180.3 | P |
| At4g12970 | expressed protein | -1 | 249.6 | P | 119.5 | A |
| At5g05600 | oxidoreductase, 2OG-Fe(II) oxygenase family | -1 | 1022.4 | P | 526.7 | P |