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| **Supplementary File 3. Primers for RT-qPCR** | | | | |
| **Gene** | **Bases** | **Forward Primer (5'-3')** | **Bases** | **Reverse Primer (3'-5')** |
| *Dusp1 (Dusp1)* | 466-484 | AGTGCAGAATCCGGATGCA | 523-508 | CTGGGCCCCCCTGATC |
| *Dusp2 (Pac1)* | 395-412 | TGCGAGGCGGTTTCAAAA | 460-443 | CCTGGGCAGGAGCTTCAG |
| *Dusp4 (Mkp2)* | 761-779 | ACCACAAGGCCGACATCAG | 827-805 | CAGTCCTTTACTGCGTCGATGTA |
| *Dusp5 (Hvh3)* | 709-733 | GACATTAGCTCCCACTTTCAAGAAG | 783-764 | GACCAGGACCTTGCCTCCTT |
| *Dusp6* | 345-362 | CGGGCTGCTGCTCAAGAA | 409-389 | TGAAGCCACCTTCCAGGTAGA |
| *Dusp7* | 521-543 | TCCAAGGTGGTTTCAACAAGTTC | 588-567 | CGAGCTGTCCACGTTAGTCTCA |
| *Dusp8* | 218-237 | CTACACGGAGCCAGGTGGAT | 284-262 | TCTCGTGTGCTCTGGTCATACAC |
| *Dusp0* | 1126-1146 | GCCACAGACAGCAACAAACAG | 1200-1180 | CTGGTGAGCTTCCTCGATGAA |
| *Dusp16* | 505-522 | GGCTGCCAGCGAGATGTC | 569-546 | TTTAACACATAGCCAATCCCATTC |