**Supplementary Table 2 | Primers for Real-time Quantitative PCR**

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| **Mouse Gene** | **Transcript ID** | **Primer Sequence (from 5’ to 3’)** | **Tm** | **Exon Spanning,****Amplicon Length** |
| *Icam-1* | NM\_010493.3 | FOR: GTGGCGGGAAAGTTCCTG | 60.0 | YES, 170 bp |
| REV: CGTCTGCAGGTCATCTTAGGAG |
| *Icam-2* | NM\_010494.2 | FOR: ATCAACTGCAGCACCAACTG | 60.0 | YES, 215 bp |
| REV: ACTTGAGCTGGAGGCTGGTA |
| *Vcam-1* | NM\_011693.3 | FOR: AGTTGGGGATTCGGTTGTTC | 60.0 | NO, 107 bp |
| REV: CATTCCTTACCACCCCATTG |
| *Sele* / *E-selectin* | NM\_011345.2 | FOR: AGCTACCCATGGAACACGAC | 60.0 | YES, 199 bp |
| REV: ACGCAAGTTCTCCAGCTGTT |
| *Selp* / *P-selectin* | NM\_011347.2 | FOR: GTCCACGGAGAGTTTGGTGT | 60.0 | YES, 241 bp |
| REV: AAGTGGTGTTCGGACCAAAG |
| *Hprt* | NM\_000194.3 | FOR: CTTCCTCCTCAGACCGCTTT | 60.0 | YES, 146 bp |
| REV: TTTTCCAAATCCTCGGCATA |