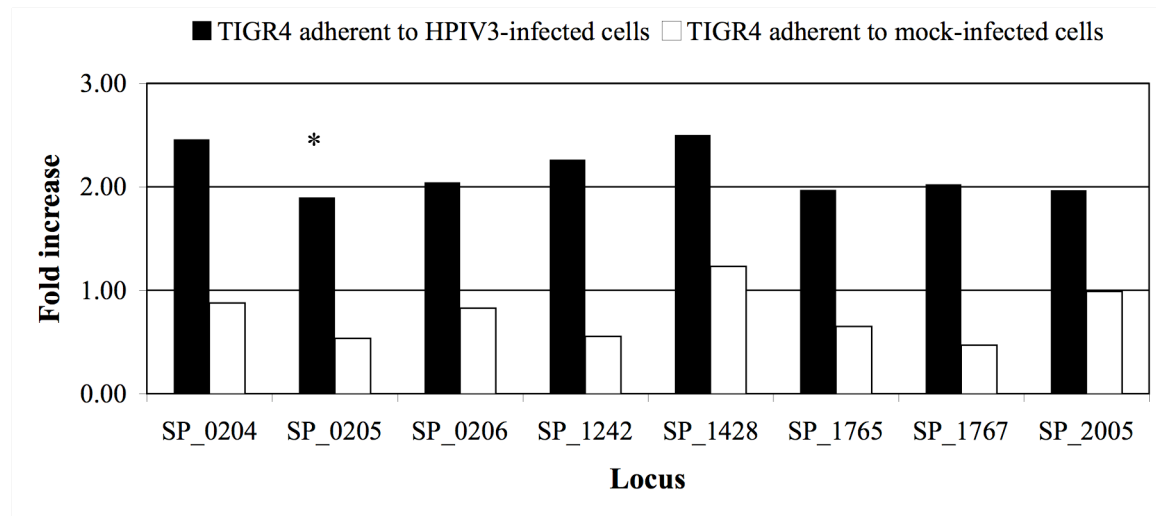


Figure 2.



SP_0204 – SP_0206: ribonucleotide reductase (*nrdDG* operon); SP_1242: *qInQ*; SP_1428 & SP_2005: hypothetical proteins; SP_1765 – SP_1767: glycosyl transferase genes

* SP_0205 was co-opted into the final analysis since it appeared to be co-regulated as part of an operon