1	Supplementary Data
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3	Metagenomic profiling of placental tissue suggests DNA virus infection of the placenta is
4	rare
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6	Adam A Witney, Sean Aller and Blair L Strang
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9	Supplementary Figure 1 Number of reads corresponding to phiX174 in the
10	placental metagenomes. The number of reads corresponding to phiX174 in each
11	placental metagenomic sample was determined using Kraken2. The sample accession
12	of each metagenomic sample was noted on each x-axis. Metagenomic samples came
13	from either full-term pregnancies, pre-term pregnancies, full term pregnancies that had
14	reported antenatal infection or pre-term pregnancies that had reported antenata
15	infections (Pt + inf).
16	
17	Supplementary Figure 2 Number of reads corresponding to human herpesvirus
18	genomes in placental metagenomes using a human herpesvirus database. The
19	number of reads corresponding to human herpesvirus genomes in each placental
20	metagenomic sample was determined by comparing reads to a database of human
21	herpesvirus genomes using Kraken2. The sample accession of each metagenomic
22	sample was noted on the x-axis. Metagenomic samples came from either full-term

pregnancies, pre-term pregnancies, full term pregnancies that had reported antenatal

24 infection or pre-term pregnancies that had reported antenatal infections (Pt + inf). HCMV

is shown as human herpesvirus 5.

Supplementary Figure 3 Number of reads corresponding to virus genomes in placental metagenomes using the GATK PathSeq tool. The number of reads corresponding to human viruses in each placental metagenomic sample was determined using GATK PathSeq. The sample accession of each metagenomic sample was noted on the x-axis. Metagenomic samples came from either full-term pregnancies, pre-term pregnancies, full term pregnancies that had reported antenatal infection or pre-term pregnancies that had reported antenatal infections (Pt + inf). HCMV is shown as human betaherpesvirus 5.

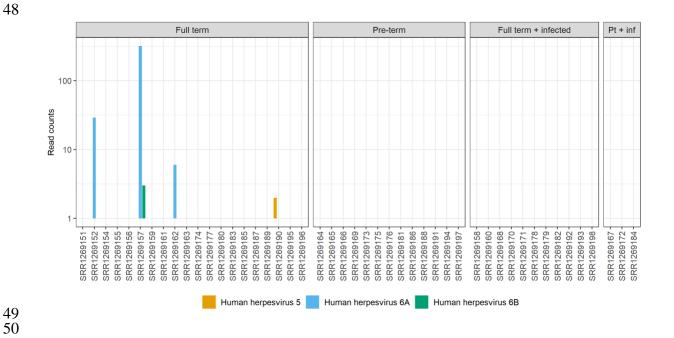
Table S1 Accession numbers for Human Microbiome Project data. See excel sheet

the accompanies this manuscript.

Supplementary figure 1

 SRR1269153
SRR1269163

Supplementary figure 2



Supplementary figure 3

