Supplementary Data 1

Exploring the cause of diarrhoea and poor growth in 8-11 weeks old pigs from an Australian pig herd using metagenomics sequencing

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>PCV2 Conjugated 40 PC46 BC26
ACGTCATATCTGAAAACGAAAGAAGTGCGCTGTAAGTATTACCAGCGCACTTCGGCAGCGGC
AGCACCTCGGCAGCACCTCAGCAGCAACATGCCCAGCAAGAAGAATGGAAGAAGCGGACCCC
AACCACATAAAAGGGGGGTGTTCACTCTGAATAATCCTTCCGAAGACGAGCGCAAGAAAATA
CGGGAGCTCCCAATCTCCCTGTTTGATTATTTTATTGTTGGCGAGGAGGGTAATGAGGAAGG
ACGAACACCTCACCTCCAGGGGTTCGCTAATTTTGTGAAGAAGCAAACTTTTAATAAAGTGA
AGTGGTATTTGGGTGCCCGCTGCCACATCGAGAAAGCCAAAGGAACTGATCAGCAGAATAAA
GAATATTGCAGTAAAGAAGGCAACTTACTTATTGAATGTGGAGCTCCTCGATCTCAAGGACA
ACGGAGTGACCTGTCTACTGCTGTGAGTACCTTGTTGGAGAGCGGGAGTCTGGTGACCGTTG
CAGAGCAGCACCCTGTAACGTTTGTCAGAAATTTCCGCGGGCTGGCTGAACTTTTGAAAGTG
AGCGGGAAAATGCAGAAGCGTGATTGGAAGACCAATGTACACGTCATTGTGGGGCCACCTGG
GTGTGGTAAAAGCAAATGGGCTGCTAATTTTGCAGACCCGGAAACCACATACTGGAAACCAC
CTAGAAACAAGTGGTGGGATGGTTACCATGGTGAAGAAGTGGTTGTTATTGATGACTTTTAT
GGCTGGCTGCCGTGGGATGATCTACTGAGACTGTGTGATCGATATCCATTGACTGTAGAGAC
TAAAGGTGGAACTGTACCTTTTTTGGCCCGCAGTATTCTGATTACCAGCAATCAGACCCCGT
TGGAATGGTACTCCTCAACTGCTGTCCCAGCTGTAGAAGCTCTCTATCGGAGGATTACTTCC
TTGGTATTTTGGAAGAATGCTACAGAACAATCCACGGAGGAAGGGGGCCAGTTCGTCACCCT
TTCCCCCCCCTGCCCTGAATTTCCATATGAAATAAATTACTGAGTCTTTTTTTATCACTTCG
TAATGGTTTTTATTATTCATTAAGGGTTAAGTGGGGGGTCTTTAAGATTAAATTCTCTGAAT
TGTACATACATGGTTACACGGATATTGTATTCCTGGTCGTATATACTGTTTTCGAACGCAGT
GCCGAGGCCTACGTGGTCTACATTTCCAGCAGTTTGTAGTCTCAGCCACAGCTGGTTTCTTT
TGTTGTTTGGTTGGAAGTAATCAATAGTGGAATCTAGGACAGGTTTGGGGGTGAAGTAGCGG
GAGTGGTAGGAGAAGGGCTGGGTTATGGTATGGCGGGAGGAGTAGTTTACATAGGGGTCATA
GGTGAGGGCTGTGGCCTTTGTTACAAAGTTATCATCTAGAATAACAGCACTGGAGCCCACTC
CCCTGTCACCCTGGGTGATCGGGGAGCAGGGCCAGAATTCAACCTTAACCTTTCTTATTCTG
TAGTATTCAAAGGGCACAGAGCGGGGGTTTGAGCCCCCTCCTGGGGGAAGAAAGTCATTAAT
ATTGAATCTCATCATGTCCACCGCCCAGGAGGGCGTTCTGACTGTGGTTCGCTTGATAGTAT
ATCTGAAGGGGCGGGGGAGGCGGGTATTGAAGATGCCGTTTTTCCTTCTCCAGCGGTAACGG
TGGCGGGGGTGGACGAGCCAGGGGCGGCGGCGGAGGATCTGGCCAAGATGGCTGCGGGGGCG
GTGTCTTCTTCTCCGGTAACGCCTCCTTGGATACGTCATATCTGAAAACGAAAGAAGTGCGC
TGTAAGTATTACCAGCGCACTTCGGCAGCGGCAGCACCTCGGCAGCACCT