

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: GO enrichment of genes in different clusters of Figure 1c. P-values are calculated based on the accumulative hypergeometric distribution.

File Name: Supplementary Data 2

Description: Species specific or conserved orthologous genes between rhesus cluster 1 and human cluster 1.

File Name: Supplementary Data 3

Description: Species specific or conserved orthologous genes between rhesus cluster 2 and human cluster 2.