

Supplementary Table 2. Number of variants by different genome regions in KC cases and controls

gene	KC	control
<i>ATP6</i>	37	40
<i>ATP8</i>	8	7
<i>COX1</i>	37	49
<i>COX2</i>	21	26
<i>COX3</i>	28	29
<i>CYTB</i>	54	57
<i>ND1</i>	39	35
<i>ND2</i>	36	39
<i>ND3</i>	14	22
<i>ND4</i>	41	46
<i>ND4L</i>	10	10
<i>ND5</i>	68	68
<i>ND6</i>	25	18
<i>RNR1</i>	23	35
<i>RNR2</i>	27	19
tRNA	32	35
noncoding	189	190