

Supplementary Table 1. Exome sequencing coverage and mapping in PCD-221 II:1

Read length	76 bp
Total bases	3 Gb
Total reads	40,375,630
Reads mapped	40,132,202 (99.3%)
% Target bases with coverage $\geq 4X$	93.1%
% Target bases with coverage $\geq 10X$	84.9%
% Target bases with coverage $\geq 20X$	68.3%
Duplication rate	5.91%