

**Table 1** | Summary statistics for exome sequencing - mapping and coverage

<b>ID</b>	<b>Variants</b>	<b>% Autosome heterozygosity</b>	<b>% X heterozygosity</b>	<b>Pipeline gender</b>
<b>II(1)</b>	24934	61.358	18.49	male
<b>II(2)</b>	25001	61.2082	18.13	male

*Variants – total called variants; % Het – genome wide % of calls heterozygous; % X Het - % calls mapped to X chromosome heterozygous; Pipeline. Gender – apparent gender based upon % X Het.*