

**Sequencing coverage and variant calls in each sample**

<b>Sample ID</b>	<b>Total reads</b>	<b>Mean target coverage</b>	<b>Variants passed QC</b>
<b>4001</b>	62,135,534	50.2	200,298
<b>4002</b>	66,022,486	49.7	215,897
<b>5001</b>	87,865,300	64.3	264,505
<b>5003</b>	70,384,372	58.6	232,068
<b>5004</b>	61,150,770	43.4	205,672
<b>5005</b>	65,183,680	51.5	223,182