|  |  |  |  |
| --- | --- | --- | --- |
| **Sample** | **Total SNPs identified** | **SNPs verified/SNPs tested** | **Methods** |
| Culture I | 26 | 24/24 | Previously reported 7 |
| Culture II | 42 | 6/6 | Direct PCR & sanger sequencing |
| 30/30 | Previously reported 7 |
| Vesicle II | 197 | 41/41 | long PCR and 454 sequencing |
| CSF I | 35 | 23/23 | long PCR and 454 sequencing |

**Supplementary Table S2 | Confirmation of fixed SNPs identified in assembled consensus sequences**