**Table S2: Massively parallel sequencing results.**

|  |  |  |  |
| --- | --- | --- | --- |
| **Mouse** | **Coverage (%)** | **Depth (fold)** | **Number of SNVs** |
| Deaf18.D002 | 89.7 | 95 | 10 |
| Deaf18.D012 | 89.7 | 84 | 14 |