**Table S4** Comparison of performance on the independent test set between CDMPred and other methods designed to predict single nucleotide driver variants in cancer using the Delong test

|  |  |
| --- | --- |
| **Method** | **P-value** |
| Cscape-somatic | <0.0001 |
| TransFIC | <0.0001 |
| FATHMM | <0.0001 |
| CanDrA | <0.0001 |
| CHASM\_plus | <0.0001 |
| CHASM | <0.0001 |