Table S4. Percent of variants in coding regions with MAF<1% in various studies. All comparisons involve persons of European ancestry

|  |  |
| --- | --- |
| **Study1** | **%** |
| Present study - observed | 81% |
| Present study - downsampled to match Fu *et al*. | 79% |
| Present study - downsampled to match Tennessen *et al*. | 70% |
| Fu *et al*. | 93% |
| Tennessen *et al*. | 87% |
| Nelson *et al*. downsampled to present study | 95% |

1References: Nelson *et al*. (Science 2012; 337:100-104), Tennessen *et al*. (Science 2012; 337:64-69), Fu *et al*. (Nature 2013; 493:216-220)