|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **Study Name** | **No. of IS cases** | **No. of LVD cases** | **No. of CE cases** | **No. of SVD cases** | **No. of controls** |
| Portugal\* | 395 | 62 | 99 | 37 | 264 |
| Krakow\* | 295 | 50 | 100 | 28 | 297 |
| Lund\* | 346 | 0 | 0 | 0 | 191 |
| Leuven\* | 123 | 25 | 38 | 14 | 0 |
| Germany\* | 3,264 | 2,013 | 669 | 246 | 1,074 |
| USA\* | 714 | 106 | 157 | 128 | 214 |
| CADISP\*\* | 555 | 67 | 211 | 31 | 9,259 |
| NINDS-SiGN\*\* | 7,743 | 1,130 | 2,001 | 1,408 | 17,970 |
| RACE1\*\*\* | 1,218 | 200 | 229 | 192 | 1,158 |
| RACE2\*\*\* | 1,167 | 155 | 193 | 122 | 4,035 |

**Table e-2:** Overview of samples used in the replication analysis. \* indicates de-novo genotyped Caucasian samples, \*\* indicates in-silico lookup Caucasian samples, \*\*\* indicates in-silico lookup South Asian samples. NA indicates that no samples were available for this category.