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| **Table 1. Studies included in this analysis.** |
| **Characteristic** | **UK Biobank** | **GOCHA** | **ISGC ICH Study** | **GERFHS** |
| **Analytical stage** | AssociationCholesterol level ~ PRS | AssociationICH Risk ~ PRS | AssociationICH Risk ~ PRS | AssociationICH Risk ~ PRS |
| **Study design** | Cohort | Case / Control | Case / Control | Case / Control |
| **Study participants** | 316,428 | 277 / 248 | 563 / 523 | 446 / 490 |
| **Age, mean (SD)** | 68 (8) | 73 (10) / 72 (8) | 71 (14) / 66 (16) | 70 (14) / 68 (13) |
| **Female sex n, %** | 170,871 (54) | 130 (47) / 123 (50) | 252 (45) / 255 (49) | 211 (47) / 235 (48) |
| **Genotyping platform** | Affymetrix UK Biobank array | Illumina HumanHap550 | Illumina HumanHap550 | Affymetrix 6.0 |
| **Genotyped SNPs** | 820,967 | 527,508 | 527,508 | 580,491 |
| **Imputed SNPs** | 73,355,667 | 7,965,700 | 7,965,700 | 7,967,430 |

**Acronyms:** UK: United Kingdom; GOCHA: Genetics of Cerebral Hemorrhage with Anticoagulation; ISGC: International Stroke Genetics Consortium; GERFHS: Genetic and Environmental Risk Factors for Hemorrhagic Stroke; ICH: intracerebral hemorrhage; PRS: polygenic risk score; SNPs (single nucleotide polymorphisms); SD: standard deviation.