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IGEMS: The Consortium on Interplay of Genes and Environment across Multiple Studies-- An Update

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IGEMS consortium**Abstract**

The Interplay of Genes and Environment across Multiple Studies (IGEMS) is a consortium of 18 twin studies from 5 different countries (Sweden, Denmark, Finland, United States, and Australia) established to explore the nature of gene-environment interplay in functioning across the adult lifespan. Fifteen of the studies are longitudinal, with follow-up as long as 59 years after baseline. The combined data from over 76,000 participants aged 14–103 at intake (including over 10,000 monozygotic and over 17,000 dizygotic twin pairs) support two primary research emphases: (1) investigation of models of GE interplay of early life adversity, and social factors at micro and macro environmental levels and with diverse outcomes including mortality, physical functioning, and psychological functioning, and (2) improved understanding of risk and protective factors for dementia by incorporating unmeasured and measured genetic factors with a wide range of exposures measured in young adulthood, midlife and later life.

Although the association between social context and late-life health and functioning is well established (Cacioppo et al., 2002; Cohen, 2004), the mechanisms for these associations or how social context relates to the biological and genetic factors known to contribute to later life functioning has yet to be fully understood. The advantages of twin studies are the strengthening of causal inference through co-twin control methods (McGue, Osler, & Christensen, 2010), the use of biometric models to quantify genetic and environmental variance (Rijsdijk & Sham, 2002), studying sex effects by leveraging data from opposite sex pairs, determining the extent to which associations between risk and outcome are driven by the same genetic or the same environmental influences, and testing whether familial factors (genetic and rearing effects) on the outcome may change as a function of the exposure (van der Sluis, Posthuma, & Dolan, 2012). As an international consortium with harmonized

measures of risk, pathway, and contextual factors, assessed longitudinally on a large number of twins, IGEMS is particularly well suited for investigating the contribution of GE interplay to functioning in multiple domains across adulthood.

The IGEMS consortium includes more than 76,000 twins from 18 studies representing 5 countries (Sweden, Denmark, Finland, United States, and Australia). The sample spans a wide age range (15 to 103 years at intake) and has sufficient power to address issues that typically elude most studies. IGEMS also includes a set of well-characterized longitudinal phenotypes, including measures of physical health, cognitive health, and emotional health, and measures of multiple facets of adult SES (e.g., occupation, education, financial strain), as well as rearing SES that are harmonized over time and across studies. The twin structure of the dataset permits using established twin methods to test key hypotheses on the nature of GE interplay, while the dense genotyping of a large subset of IGEMS participants allows us to confirm and extend these twin analyses through analyses of polygenic risk scores (PRS) for health outcomes and for education. Importantly, IGEMS cohorts span multiple countries and historical periods, allowing us to determine whether models of GE interplay established at the micro (i.e., individual) level also apply at the macro (i.e., country and historical period) level.

There are two substantive emphases in the IGEMS consortium: First, socio-economic conditions are a major social determinant of health (Freedman, Martin, Schoeni, & Cornman, 2008; Mensah, Mokdad, Ford, Greenlund, & Croft, 2005; Mirowsky & Ross, 2003; Sattler, Toro, Schonknecht, & Schroder, 2012; Sharp & Gatz, 2011). The oft-cited “gradient” for socio-economic status (SES) represents the association between health and SES as continuous and monotonic and not fully explained by poorer health among those who are impoverished (Adler et al., 1994). Most research focuses on individual-level socio-economic status—social status that accrues to occupational classification, education, and income as well as access to social, human- and income- capital. Others consider the macro-economic environment, such as the extent of social inequality in a country (Kawachi, Levine, Miller, Lasch, & Amick, 1994; Lynch et al., 2004; Lynch, Smith, Kaplan, & S., 2000). IGEMS is unusual in integrating individual- and country-level contributors to health gradients. Further, although both genetic and environmental factors are known to contribute to the SES-health gradient (Lahey, D’Onofrio, & Waldman, 2009; McGue et al., 2010; Rutter, 2009), the mechanisms by which these factors combine to influence health outcomes (Gene x Environment interplay) are poorly understood.

Recent research has identified alternative models of GE interplay important to understanding health and disease (Boardman, Daw, & Freese, 2013; Reiss, Leve, & Neiderhiser, 2013; Shanahan & Boardman, 2009; Shanahan & Hofer, 2005). Although these models recognize that individuals inherit differential sensitivity to the environment, they differ in their environmental focus (disease-triggering effects of toxic environments vs. health-promoting benefits of favorable environments) and the expected genetic contribution to disease (maximized in adverse environments, in favorable environments, or at both extremes). The differences between models of GE interplay have implications beyond resolving an academic dispute. Environmental improvements would be expected to reduce or eliminate genetically based health disparities under some models (e.g., diathesis-stress) but expand

them under others (e.g., social distinction) or have a mixed impact (e.g., differential susceptibility (Boardman et al., 2013; Reiss et al., 2013; Shanahan & Boardman, 2009; Shanahan & Hofer, 2005). Understanding whether socially enriched environments compensate for genetic vulnerability or whether they preferentially promote good health among genetically selected individuals, for example, is essential for both translating research into effective prevention strategies and for anticipating consequences of social policies.

A second substantive emphasis of the IGEMS consortium is cognitive functioning in adulthood. In particular, Alzheimer's disease and related dementias (ADRD), along with mild cognitive impairment (MCI), present a major public health challenge due to the large numbers of people affected and lack of a clear path to prevention or cure (Katzman, 2004; Livingston, Sommerlad, & Orgeta, 2017; Wang, MacDonald, Dekhtyar, & Fratiglioni, 2017). While it is generally recognized that Alzheimer's disease (except for rare dominantly inherited forms) is caused by multiple genetic and environmental factors, it remains unclear how these factors contribute to the disease, whether they function independently or through interactions with each other (Gatz, Mortimer, & Fratiglioni, 2007), the ages at which these factors have the greatest impact (Wang et al., 2017), and if these factors affect men and women equally (Mielke, Vemuri, & Rocca, 2014). Moreover, it remains to be determined whether many of these factors represent modifiable targets appropriate for intervention, or actually reflect pre-existing genetic vulnerability to ADRD and its risk factors.

IGEMS STUDIES

From an original consortium of 8 twin studies (Pedersen et al., 2013), IGEMS has expanded to include 18 studies from 5 countries, representing the strongest available longitudinal twin studies of adulthood and aging in the world. The total sample size is now 76,233 including both members of 10,266 monozygotic (MZ) pairs and 17,288 dizygotic (DZ) pairs, of which 5,063 pairs are opposite sex DZ (OSDZ). The summary below outlines the sampling principles for each study. Numbers of pairs and age ranges at intake are provided in Table 1, as well as the number of waves and length of follow-up, where appropriate. Total Ns refer to individuals and include members of incomplete pairs. Note that updates for several of the studies are included in this issue.

Sweden.

Swedish studies are drawn from the population-based Swedish Twin Registry. The Swedish Adoption/Twin Study of Aging (SATSA) began in 1984 (Finkel & Pedersen, 2004). The base population comprises all pairs of twins from the registry who indicated that they had been separated before the age of 11 and reared apart, and a sample of twins reared together matched on the basis of gender, date and county of birth. The OCTO-Twin Study (Origins of Variance in the Old-Old) included twin pairs who were over the age of 80 at baseline in 1991 (McClearn et al., 1997). Aging in Women and Men: A Longitudinal Study of Gender Differences in Health Behaviour and Health among Elderly (GENDER) is a study of opposite-sex twin pairs born between 1906 and 1925 (Gold, Malmberg, McClearn, Pedersen, & Berg, 2002). The Twin and Offspring Study in Sweden (TOSS) includes pairs of same-sex twins and their offspring (Neiderhiser & Lichtenstein, 2008). The Study of

Dementia in Swedish Twins (HARMONY) was conducted between 1998 and 2004. HARMONY screened all twins age 65 and over in the Screening Across the Lifespan of Twins (SALT) effort (Lichtenstein et al., 2006) and clinically assessed those who screened positive or whose co-twin screened positive for cognitive impairment (Gatz, Fratiglioni, Johansson, & Berg, 2005)

Denmark.

The Longitudinal Study of Aging Danish Twins (LSADT) began in 1995 with the assessment of members of like-sex twin pairs born in Denmark prior to 1920; twins were recruited regardless of whether their cotwin was alive (Christensen, Holm, McGue, Corder, & Vaupel, 1999). The study of Middle-Aged Danish Twins (MADT) includes twins ranging in age from 46 to 68 years at the original assessment (Osler, McGue, Lund, & Christensen, 2008). The Mid-aged Danish Twin (MIDT) study includes twins representing members of the Danish Twin Registry for the birth years 1931 through 1969 not already participating in MADT.

Finland.

The older Finnish Twin Cohort (FTC) study spans 4 decades; it was initiated in 1975 by contacting all same-sex Finnish twin pairs born before 1958 with both co-twins alive in 1975 (Kaprio & Koskenvuo, 2002). FinnTwin16 (FT16) is a cohort of younger twins born between 1975 and 1979. Waves 4 and 5, when the study participants were in their 20s and 30s, are included in IGEMS (Kaprio, Pulkkinen, & Rose, 2002).

United States.

Each US study consists of an independent sample. The Minnesota Twin Study of Adult Development and Aging (MTSADA) is a population-based sample drawn from state birth records (Finkel & McGue, 1993; McGue, Hirsch, & Lykken, 1993). The Vietnam Era Twin Study of Aging (VETSA) is a community dwelling sample of male–male twin pairs, all of whom served in some branch of US military service sometime between 1965 and 1975 (Kremen, Franz, & Lyons, 2013). Midlife in the United States (MIDUS) is a national telephone/mail survey originally carried out in 1995–1996 that included specific recruitment methods for twins (South & Krueger, 2012). The Carolina African-American Twin Study of Aging (CAATSA) used public records to identify all living African-American twins in the State of North Carolina born between 1920 and 1970 (Whitfield, 2013). The Project Talent Twin and Sibling Study (PTTS) includes 4481 twins and triplets plus 522 of their siblings, drawn from Project Talent, a longitudinal study begun in 1960 with a nationally representative sample of U.S. high school students born 1942–1946 (Flanagan, 1962). Follow-up surveys were conducted in young adulthood (ages 19, 23 and 29) and then in 2014 (ages 68–72) and 2019 (aged 73–77). 1, 5, and 11 years following the year of expected high school graduation. The Project Talent Twin and Sibling Study (PTTS) tracked 96.4% of the original PT twins (Prescott et al., 2013). The National Academy of Sciences-National Research Council (NAS-NRC) Twin Registry consists of white male twin pairs born in the years 1917 to 1927, both of whom served in the armed forces, mostly during World War II (Page, 2006).

Australia.

The Australian Over 50's study (A50) is based on a questionnaire mailed between 1993 and 1995 to Australian twins age 50-95 (Hopper, 2002; Hopper, Foley, White, & Pollaers, 2013). The Older Australian Twins Study (OATS) incorporates in-person assessments every two years of twins age 65 and older in the three Eastern states of Australia: New South Wales, Victoria and Queensland (Sachdev et al., 2009).

The range in study years and intake ages across the 18 IGEMS studies results in unique coverage of cohorts and historical periods. As shown in Table 2, the IGEMS sample permits sequential comparisons of sex and SES effects across six cohorts.

IGEMS MEASURES

Measures used in IGEMS analyses include aging-relevant outcomes in three broad domains: physical health and functional ability (e.g., self-reported diseases, subjective health, body mass index, grip strength, motor function, activities of daily living), psychological well-being (e.g., depressive symptoms, anxiety symptoms, subjective well-being, loneliness), and cognitive health (i.e., scores on cognitive tests; dementia). Predictors and covariates include health behaviors (e.g., smoking, alcohol, physical activity, cognitively-engaging leisure activity), social resources, and indicators of socioeconomic status. Table 3 presents a list of some of the primary phenotypes assessed and the number of IGEMS studies that include each variable.

Because participating studies differed in how similar constructs were assessed, IGEMS gives emphasis to harmonization of relevant phenotypes and outcomes. Creating scores that are common across studies enables pooling data across samples, in order to increase power. Score harmonization requires overlapping item content across studies as well as across time for longitudinal hypotheses. For some measures, it was straightforward to create a common metric; e.g., BMI, lung function, and blood pressure. For harmonizing education and occupation, we have recoded all studies to the International Standard Classification of Education (ISCED (UNESCO, 1997) and the International Standard Classification of Occupations (ISCO (Ganzeboom, De Graaf, Treiman, & De Leeuw, 1992), as an international standard. Where a common metric was not already available, overlapping item content and response formats were identified and item response theory (IRT) or factor-analytic techniques were implemented to create harmonized scores across studies. Where there were no common items across studies, IGEMS has collected separate samples that were administered the different measures used in different IGEMS studies to measure a given construct, with those results used to establish 'crosswalks' between the different scales (Gatz, Reynolds, et al., 2015).

Across the 18 IGEMS studies, there is genome-wide genotype information available from 22,765 subjects, including MZ cotwins of genotyped individuals (Table 4), which will be available for analysis with appropriate correction for clustering. In Sweden, there are also genotypes available from an additional 14,244 individuals who participated in Screening Across the Lifespan of Twins (SALT) substudies known as TwinGene and SALTY (Scanning Across the Lifespan of Twins – Young (Magnusson et al., 2013), but under age 65

when screened in SALT. Thus, the total number of individuals with genome wide genotyping available to us is 37,009. Polygenic risk scores (PRS) have or will be computed for cardiovascular disease, lipids, Type II diabetes, Alzheimer's disease, neuroticism, major depression and depressive symptoms, smoking and alcohol behaviors, well-being, and educational attainment. We will compute new PRS scores as new GWAS training sets become available.

APPROACHES

IGEMS harnesses the full analytic potential of twin designs to address issues of gene-environment interplay as well as risk and protective factors for aging related outcomes. Methods using MZ within-pair differences allow us to test for the presence of GxE without having a specific measured early environment. With a MZ within-pair approach, we established evidence of GxE for BMI, depressive symptoms, a physical illness index, several cognitive domains (Reynolds et al., 2016), and longitudinal grip strength trajectories (Pedersen et al., 2016). Results also suggested that APOE may act as a 'variability gene' for symptoms of depression and spatial reasoning but not for other cognitive measures or BMI, with greater intrapair differences for non- $\epsilon 4$ carriers. For grip strength trajectories, a buffering effect for $\epsilon 2$ carriers emerged, with lower sensitivity to environments and better-maintained performance.

Co-twin control methods also provide the opportunity to strengthen causal inferences and test whether associations between early life exposures and late life outcomes are due to confounding by common familial (genetic and/or shared rearing) influences. For example, Mosing and colleagues (2016) found that, of a number of birth characteristics, low birth weight was associated with poorer self-rated health in adulthood, when evaluated with a generalized estimating equation adjusting for the twin structure. However, as these associations were attenuated in a co-twin control analyses (first in all pairs, then only in MZ pairs), there is evidence that the association is in part due to familial influences. In subsequent analyses of birth characteristics and cognitive impairment and dementia, we found evidence that low birth weight and small head circumference are risk factors for dementia. Further, head circumference was also significantly associated with age-related cognitive impairment (Mosing, Lundholm, Cnattingius, Gatz, & Pedersen, 2018). Here, within-pair analyses of identical twins suggested that the observed associations between birth characteristics and cognitive decline are likely not due to underlying familial etiology.

To quantify interplay, we have applied biometric moderation models (Purcell, 2002; van der Sluis, Dolan, Neale, & Posthuma, 2008). We have examined GxE interactions in relation to cognitive performance (Pahlen et al., 2018; Zavala et al., 2018), depression (Petkus et al., 2017), subjective health (Franz et al., 2017), and an index of physical illness (Gatz, Petkus, Franz, Kaprio, & Christensen, 2015). For most phenotypes, unique environmental variance was greater at older ages, presumably reflecting the accumulating importance of individual differences in environmental context with age. However, there was a non-uniform pattern for genetic factors over age, in combination with SES or sex moderation. In SES moderation analyses of cognition, for verbal ability and for perceptual speed (Zavala et al., 2018), genetic variance was diminished in those with higher SES, perhaps reflective of a buffering

effect on normative aging processes particularly for speed; whereas for short-term/working memory and spatial performance genetic variance was amplified with higher SES, suggesting stable experiences in enriched (high SES) environments may support genetic variation.

Because IGEMS has genome-wide genetic data, we are able to create PRS scores and incorporate these into our models of GE Interplay. In this case, the PRS scores will be entered as a moderator, together with other indicators of the environment, such as SES. The interaction between PRS and SES in regression models predicting health outcomes of interest will inform whether those at high genetic risk for a health outcome are more or less susceptible to e.g. health-promoting benefits of favorable environments.

Country-level SES-indicators are available from various online sources. These data provide historical measures of social and economic conditions from the mid-1800s to the early 2000s for each of the five IGEMS countries. Variables include: average years of education, educational inequality, gross domestic product per capita (GDP), Gini coefficient of income inequality, public social spending, and top 1% income share. As a demonstration of the use of country-level SES indicators, we examined harmonized depressive symptom scores across five countries and a wide range of birth cohorts from 1890 through 1970. We used Top 1% (share of wealth held by top 1% of residents) to index country-level inequality when participants were aged 10 (“World Inequality Database,” 2017). Controlling for age when the depressive symptom measure was completed, gender, and country-level GDP, adult depressive symptom scores were higher among those exposed to greater inequality as youths. Using a modified twin correlation model, we found greater genetic effects on depressive symptoms with exposure to greater inequality (Gatz, Finch, Beam, & Thomas, 2018).

SUMMARY

The IGEMS consortium harnesses a combination of twin designs and multiple studies representing different cohorts and contexts. The accomplishments of the consortium demonstrate the feasibility of this type of collaboration in addressing gene-environment interplay with respect to important age-related outcomes.

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IGEMS data are not publicly available given the variety of data agreements and regulations governing the different studies and countries. However, many of the individual studies participating in IGEMS do have ways to access their data, and many of the datasets may be accessed through National Archive of Computerized Data on Aging (NACDA).

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Table 1.

Number of twins in each study included in IGEMS, by age at intake.

| Study | Age 14 - 40 | | | | | | Age 41 - 65 | | | | | | Age > 65 | | | | | | All Ages | | | Waves | | | |
|---------|-------------|-------|----------|-------|----------|-------|-------------|-------|----------|-------|----------|-------|----------|-------|----------|-------|----------|-------|----------|-------|-------|-------|-------|-------|--|
| | MZ Pairs | | DZ Pairs | | OS Pairs | | MZ Pairs | | DZ Pairs | | OS Pairs | | MZ Pairs | | DZ Pairs | | OS Pairs | | All | DZ | OS | Total | Total | Years | |
| | N | Total | N | Total | N | Total | N | Total | N | Total | N | Total | N | Total | N | Total | N | Total | Pairs | Pairs | Pairs | N | # | Span | |
| SA TSA | 38 | 234 | 61 | 150 | 273 | 1074 | 95 | 184 | 149 | 202 | 647 | 283 | 518 | 2210 | 17 | 30 | | | | | | | | | |
| OCTO | | | | | | | | | | | | | | | | | | | | | | | | | |
| GENDER | | | | | | | | | | | | | | | | | | | | | | | | | |
| TOSS | 75 | 314 | 82 | 307 | 401 | 1420 | | | | | | | | | | | | | | | | | | | |
| HARMONY | | | | | | | | | | | | | | | | | | | | | | | | | |
| LSADT | | | | | | | | | | | | | | | | | | | | | | | | | |
| Danish | | | | | | | | | | | | | | | | | | | | | | | | | |
| MADT | | | | | | | | | | | | | | | | | | | | | | | | | |
| MIDT | | | | | | | | | | | | | | | | | | | | | | | | | |
| FTC | | | | | | | | | | | | | | | | | | | | | | | | | |
| Finn | | | | | | | | | | | | | | | | | | | | | | | | | |
| FT16 | 835 | 1798 | 944 | 5608 | | | | | | | | | | | | | | | | | | | | | |
| MTSADA | 45 | 159 | 27 | 234 | 200 | 956 | 54 | 61 | 244 | 244 | 244 | 333 | 288 | 1359 | 3 | 12 | | | | | | | | | |
| VETSA | | | | | | | | | | | | | | | | | | | | | | | | | |
| MIDUS | 150 | 763 | 213 | 176 | 274 | 1484 | | | | | | | | | | | | | | | | | | | |
| CAATSA | 36 | 198 | 61 | 54 | 102 | 377 | 14 | 15 | 5 | 102 | 104 | 104 | 178 | 677 | 1 | NA | | | | | | | | | |
| PTTS | | | | | | | | | | | | | | | | | | | | | | | | | |
| NAS-NRC | | | | | | | | | | | | | | | | | | | | | | | | | |
| A50 | | | | | | | | | | | | | | | | | | | | | | | | | |
| OATS | | | | | | | | | | | | | | | | | | | | | | | | | |
| Total | 1185 | 2248 | 1064 | 7326 | 9511 | 43744 | 2463 | 5529 | 2069 | 25163 | 10266 | 17288 | 5063 | 76233 | | | | | | | | | | | |

Note: MZ = monozygotic, DZ = dizygotic, OSDZ = opposite sex dizygotic pairs. Total N refers to individuals from both complete and incomplete pairs. Some individuals may have participated in more than one study, e.g., in A50 and OATS. The totals in the bottom row count each pair or individual once.

Table 2:

Total number of individuals (% female) in each birthyear range (cohort) by age at intake.

| Intake Age | Birthyears | | | | | | | Total | # Cohorts |
|------------|------------|--------------|--------------|--------------|------------|------------|--------------|-------|-----------|
| | <1914 | 1915-1929 | 1930-1944 | 1945-1959 | 1960-1974 | 1975+ | | | |
| <35 | | | | 243 (59%) | 583 (61%) | 5643 (53%) | 6469 (54%) | 3 | |
| 35-49 | | 12,121 (0%) | 341 (50%) | 2842 (59%) | 3420 (60%) | | 18,724 (21%) | 4 | |
| 50-64 | | 2996 (15%) | 5640 (59%) | 14,466 (48%) | 356 (55%) | | 23,428 (47%) | 4 | |
| 65-79 | 524 (65%) | 11,516 (57%) | 9145 (51%) | 1974 (53%) | | | 23,159 (55%) | 4 | |
| 80-94 | 3205 (66%) | 1193 (61%) | | | | | 4398 (64%) | 2 | |
| 95+ | 55 (71%) | | | | | | 55 (71%) | 1 | |
| Total | 3784 (66%) | 27,796 (28%) | 15,126 (54%) | 19,525 (50%) | 4359 (59%) | 5643 (53%) | 76,233 (44%) | | |

Table 3:

Number of the 18 IGEMS studies with key variables

| Variables | Number of Studies |
|---|--------------------------|
| <i>Socioeconomic Status</i> | |
| Occupation: International Standard Classification of Occupations | 18 |
| Education: International Standard Classification of Education | 18 |
| Financial strain | 12 |
| Early life SES/Parental Education | 15 |
| Marital status | 18 |
| <i>Physical Health</i> | |
| Measured Blood Pressure | 7 |
| Measured Grip Strength | 8 |
| Measured Lung Function | 8 |
| Weight, height, Body Mass Index | 18 |
| Self-reported diseases (Cumulative Illness) | 17 |
| Vascular Risk (hypertension and diabetes) | 14 |
| Stroke and cardiovascular disease | 13 |
| Mortality | 16 |
| Measured Motor function and balance | 10 |
| ADL/IADL | 7 |
| Subjective Health | 18 |
| <i>Cognitive Health</i> | |
| Cognitive test scores | 16 |
| Young adult cognitive ability | 12 |
| Mini-Mental State Examination or Telephone Interview for Cognitive Status | 13 |
| Clinical dementia diagnoses | 6 |
| <i>Emotional Health</i> | |
| Anxiety (Symptoms) | 10 |
| Depression (Symptoms) | 16 |
| Subjective Well-Being | 13 |
| Loneliness | 15 |
| <i>Health Behaviors</i> | |
| Physical/Leisure activity | 15 |
| Alcohol, smoking | 18 |

Table 4.

Genotype data available in participating IGEMS twin studies

| Country | N Individuals* | Baseline Age | N full twin pairs | | |
|---|-------------------|-----------------|-------------------|------|----------|
| | | | MZ* | DZ | DZ OS |
| USA | 1329 | 51-60 | 388 | 264 | - |
| Sweden [#] | 4705 | 38-108 | 601 | 633 | 582 |
| additional twins <65 yrs. ^{**} | 14244 | 44-65 | 2209 | 1535 | 1501 |
| Denmark | 1968 | 58-85 | 391 | 353 | 233 |
| Finland | 13087 | 30-95 | 1825 | 3810 | 89 |
| Australia | 1676 | 50-92 | 475 | 123 | 76 |
| TOTAL | 37009 | 30-108 | 5889 | 6718 | 2481 |

* including imputed MZ co-twins

[#] From SATSA, GENDER, HARMONY, TwinGene < 65 yrs.^{**} From TwinGene < 65 yrs. and SALTY

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