GAPS allowable uses and register linkage

If study subgroups or cases and controls have different information, please use one form for each.

Study name: _____

Study subgroup (if applicable): _____

ICD codes and phenotypes allowed for study by GAPS: _____

_____

Can these samples be used to calculate and report aggregated Sweden allele frequencies? _____

Can these samples be used for UNICORN (used in aggregate as controls in association studies)? _____

Can these samples be uploaded to Sanger HRC for imputation? _____

Is it permitted for the Honest Broker to link the genetic data to the national registers? _____

Is re-contact possible?_____ By genotype, phenotype or both? _____

Register linkage

Register	Linkage allowed		Linkage done	
Patient register	yes	no	yes	no
Cause of death register	yes	no	yes	no
Prescribed drug register	yes	no	yes	no
Multi-generation register	yes	no	yes	no
Conscription	yes	no	yes	no
Grades, testing	yes	no	yes	no
Medical birth register	yes	no	yes	no
SES (family status, income, education)	yes	no	yes	no
Population register (birthplace, migration)	yes	no	yes	no
Other _____	yes	no	yes	no
Other _____	yes	no	yes	no

What month/year was the last register linkage done? _____

Additional comments _____

_____

Signature _____