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or 5-[4-N-(aralkylsulfonyl)anil ino]pyridine having an excellent antiinflammatory, antiarthritic and analgesic activity, a process for preparing it and an antiarthritic or analgesic pharmaceutical composition containing the pyridine as an effective ingredient. 2. Prior Art We have already found that 5-(4-N-(arylsulfonyl)anili no)pyridine derivatives having a carbamoyl or carbamoyloxy group at the 4-position thereof have an

excellent antiinflammatory, antiarthritic and analgesic activity (see U.S. Pat. Nos. 4,172,124 and 4,206,212). The compounds of the present invention have an excellent activity and are different from the compounds disclosed in the foregoing patents. A security strategy for genome wide association studies. Genome wide association studies (GWAS) have contributed to the identification of hundreds of common single nucleotide

polymorphisms (SNPs) that are associated with a variety of disease traits, and have come to dominate the field of genetics today. However, the statistical power of GWAS to identify diseaseassociated loci is still low, even after accounting for multiple hypothesis testing. This is because GWAS used have been criticized as an epidemiological reverse image mapping technique, and the underlying assumptions of which are

often not fulfilled. As such, it is suggested that a newer approach of using prior knowledge and known genetic links between genotypes and phenotypes is needed to increase the power of GWAS. In this review, we show that GWAS has been successful at identifying disease genes that have been previously validated by knock out experiments and prioritize these gene candidates for future functional research.

However, most disease SNPs identified by GWAS cannot be validated by existing knock out or similar technology. We propose to integrate GWAS with gene expression and proteinprotein interaction data to prioritize genes that are important for mediating any phenotypic effects that are associated with 0cc13bf012

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