Summary information on Diseases. Homologues. Literature Interaction, Literature, Pathways, GO annotation, 1153 publications firstAuthor title and more Genome-wide association analysis identifi Saxena R trialyceride levels. Meta-analysis of genome-wide association data and large-scale replication Zeggini E identifies additional susceptibility loci for type 2 diabetes. Replication of genome-wide association signals in UK samples reveals risk loci Zeggini E for type 2 diabetes. +-- in Finals detects multiple Scott LJ Disease 5 diseases large-scale identifier name Voiaht BF 604367 LIPODYSTROPHY, FAMILIAL PARTIAL, TYPE 3; FPLD3 125853 DIABETES MELLITUS, NONINSULIN-DEPENDENT; NIDDM 609338 CAROTID INTIMAL MEDIAL THICKNESS 1 137800 GLIOMA SUSCEPTIBILITY 1; GLM1 GLIOMA OF BRAIN, FAMILIAL, INCLUDED

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