

Open Peer Review on Qeios

FAM13A wt Allele

National Cancer Institute

Source

National Cancer Institute. <u>FAM13A wt Allele</u>. NCI Thesaurus. Code C90309.

Human FAM13A wild-type allele is located in the vicinity of 4q22.1 and is approximately 550 kb in length. This allele, which encodes protein FAM13A, may play a role in the modulation of GTP metabolism. Polymorphism may be associated with chronic obstructive pulmonary disease and aberrant expression may play a role in leukemia relapse.