

Molecular diagnosis of pediatric patients with citrin deficiency in China: *SLC25A13* mutation spectrum and the geographic distribution

Wei-Xia Lin, Han-Shi Zeng, Zhan-Hui Zhang, Man Mao, Qi-Qi Zheng, Shu-Tao Zhao, Ying Cheng, Feng-Ping Chen,

Wang-Rong Wen & Yuan-Zong Song

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Homo_sapiens	FLKYASIEKNGEFFMSPNDFVTRYLNIFGESQPNPKTVELLSGVVDQTKDGLISFQEFVA
C. elegans	FDRFATKEIKGKKLMTPEDFIRGYLGLYTEENYNKETVRLLASAADTTKDGLISFEFCA
Opossum	---YASIEKDGEFFMCPEDFLCRFLNIFGGCQPSPELQLLGGVVDQTKDGLINFQEFIA
X. tropicalis	FYKYASVEKNGDYFMSPLDFVRRYLNLSGDGEPNPKTLQLLAGVVDQTKDGLISFQEFVA
Chicken	FLKYASVEKNGEFFMSPNDFVTRYLKIIGDGLPNANTVQLLAGVVDQTKDGLISFQEFVA
Pig	FKKYASIEKNGEFFMSPNDFVTRYLNIFGESQPNPKTVELLSGVVDQTKDGLISFQEFVA
Cow	FKQYASIEKNGEFFMSPNDFVTRYLNIFGESQPNPKTVELLSGVVDQTKDGLISFQEFVA
Mouse	FLKYASIEKNGEFFMSPHDFVTRYLNIFGESQPNPKTVELLSGVVDQTKDGLISFQEFVA
Rat	FLKYASIEKNGEFFMSPHDFVTRYLNIFGESQPNPKTVELLSGVVDQTKDGLISFQEFVA
Chimpanzee	FLKYASIEKNGEFFMSPNDFVTRYLNIFGESQPNPKTVELLSGVVDQTKDGLISFQEFVA
Dog	FLKYASIEKNGEFFMSPNDFVTRYLNIFGESQPNPKTVELLSGVVDQTKDGLISFQEFVA

Supplemental Information 1. Alignment analysis of the homologous proteins. The mutation p.M35V affected an amino acid highly conserved in all 11 homologous proteins from 11 eukaryotic species, including human (*Homo sapiens*), chimpanzee, dog, pig, cow, rat, mouse, chicken, *Xenopus tropicalis* (*X. tropicalis*), opossum and *Caenorhabditis elegans* (*C. elegans*).